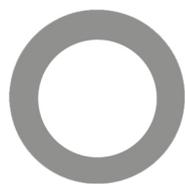


Newsletter

Team
Version

September, 2016



XMU-China Credit

Team Version

Dear all,

Here comes the Team Version for 2016 iGEM Newsletter.

For three years in a row, we have published a dozen Newsletters for iGEM competition. It is of great pleasure to announce that this year 46 teams from 21 countries and regions join us!

This year, we publish only one issue with two different versions: Part Version and Team Version.

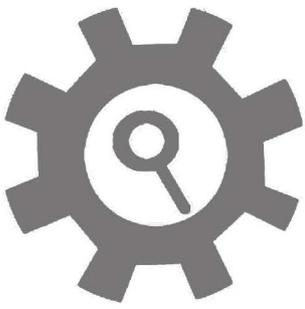
We acknowledge significant help, feedback and suggestions from the following forty-five teams (in alphabetical order):

Aachen, Aalto-Helsinki, AHUT, BGU_ISRAEL, BIT-China, Cardiff_Wales, CGU_Taiwan, Duesseldorf, Edinburgh_UG, EPFL, Evry, Freiburg, Groningen, Hannover, HUST, HokkaidoU_Japan, IIT_Kharagpur, Imperial_College, Jilin_China, Leiden, Manchester, NCTU_Formosa, OUC-China, Oxford, Peking, Peshawar, Pretoria_UP, Queens_Canada, SYSU-CHINA, SYSU-MEDICINE, Tec-Chihuahua, Tel-Hai, Tianjin, Tongji_Shanghai, UCAS, UCC_Ireland, UCL, UNIK_Copenhagen, UPO-Sevilla, UrbanTundra_Edmonton, USTC, Valencia_UPV, Vilnius-Lithuania, Washington and Westminster_UoW.

Many thanks to all of you for your generosity and contributions!
If there are any questions, please reach us at igemxmu@gmail.com.

All the best! Cheer for the event!

XMU-China



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the RWTH University

AACHEN



@iGEMAachen



iGEMAachen



igem@rwth-aachen.de



igem.aachen

We are the iGEM Team 2016 of the RWTH University in Aachen. Our team is composed of 16 students from Biology, Biotechnology, Computer Science and Biomedical Engineering. Like the past two iGEM teams of the RWTH University, we are also a very international team this year, our team members are from Germany, China and India.

Project

Our project deals with the problem caused by Boric Acid in washing detergent. Boric Acid is listed as substance of very high concern according to ECHA. For example there is the suspicion that boric acid causes reproduction toxicity. However, it is still used in liquid washing detergents for stabilization of the proteases. Our idea is to replace boric acid and achieve the same control of activity and stabilization of the protease with another mechanism. The technique we want to use is called photocaging. In our case we want to use a photo-cleavable amino acid, that will be incorporated in the protein in a way that it will inhibit the protease through an extra protection group "Photocage". These groups can be cleaved off by irradiation of light with a specific wavelength and the original amino acid remains functional. The general idea is to connect the protection group with a commonly used amino acid which is added to the active site or other important site for the functionality of the enzyme. For example, we replaced a serine in the original sequence of the protease with a photocaged serine. Thanks to the additional protection group on the unnatural amino acid, the enzyme should remain inactive as long as there's no light shining on. Our intention is making the protease unable to fold correctly because of the additional size of this specific unnatural amino acid and its position in the protease. If we shine light on it, the protease should become active again and fold correctly, since the large protection group is cleaved off and the original amino acid is revealed again.

Photocaging



For the incorporation of our unnatural amino acid we need a special tRNA which recognizes the least used stop codon UAG. UAG is used in exchange for a specific codon in the coding sequence of our protease where we want to put our unnatural amino acid in.

This tRNA needs an specific aminoacyl tRNA synthetase which attaches the appropriate amino acid onto its tRNA. This tRNA and Synthetase need to be an orthogonal pair, which means the Synthetase does not charge other tRNAs or the wrong one.

Collaboration

We are definitely interested in scientific collaborations linked to proteases, proteins or other parts of our project. Moreover we are in general also interested in collaborations regarding the Human and Practice Track.



Human Practice

In this part our team focused on visiting schools and to organize different surveys, which should help us to understand and analyze the opinion of the German public towards synthetic biology.

Aalto University and University of Helsinki

AALTO-HELSINKI



@AaltoHelsinki



Aalto-Helsinki iGEM



team@aaltohelsinki.com



aaltohelsinki_igem



blog.aaltohelsinki.com

Aalto-Helsinki is the first and only iGEM team from Finland, participating for the third time in the synthetic biology competition. Our team is composed of ten students from two different universities, seven study programmes and four nationalities!

Team

Getting our team together

Aalto-Helsinki is first and foremost a student-driven project: the idea to participate in iGEM came from the first team's student team leader; the teams are chosen by the previous years' team members; and we come up with our project on our own. We also gather the funding and sponsors by ourselves. Our team was chosen through an application process where we first sent application letters and were then interviewed. Out of about 50 applicants the ten of us were chosen. Our team got together for the first time in late February and we have been working ever since.

How Aalto-Helsinki 2016 works

Even though we come from two different universities, from different fields of science and even from different countries, we have found a common tune. Our multidisciplinary team has many kinds of expertise and lots of knowledge; we have people from biology and modelling as well as those in between.

Our project has been divided into smaller subprojects with their own teams to make it easier to achieve our goals:

The research team has five members, responsible for the biological side of the project as well as knowing what is

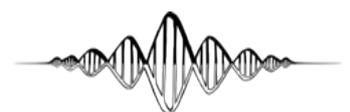
happening in the lab.

The modeling team of two is working their magic in order to provide more understanding of the biology behind our experiments.

The funding team has done an amazing job making sure that our team is able to participate to the Giant Jamboree as well as other iGEM meetings before that.

The design team is there to make sure that the visual side of our project is at least as good as the scientific.

Even though we have people in charge of each team, almost every one of us has done something in every subproject: even our mathematicians got to try out what it is like to work in a lab coat and nitrile gloves by making some liquid cultures. Our team isn't only about hard work and long days at the office or at the lab. One of the most important things is our weekly tradition of choosing a Fun Master and a Cake Master. The Fun Master of the week is responsible for taking the team to do something fun together. So far, we've gone ziplining, hiking and room-escaping, just to name a few. The Cake Master is also vital: our weekly team meetings simply couldn't work if there wasn't any cake!



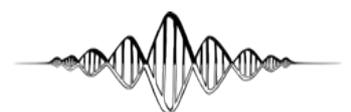
Project

Deciding on the Project Idea

From the beginning we had quite a clear vision that we wanted a project that had local significance in Finland and that lacked a good solution at the moment. Our brainstorming resulted in more than 50 ideas ranging from bacteria in space to a bacterial biocomputer. The final idea seemed promising right from the start: tackling the problem of cyanobacteria. Our idea has evolved as we have spoken to different experts from different fields such as environmental sciences, modelling and synthetic biology. We have had the opportunity to collaborate with researchers and experts, and this has been really productive. Our team's wide range of expertise has also made it possible to tackle our problem from different angles, from the production of proteins to the mathematical modeling of the toxin's effects. This also gives every team member a chance to learn new things outside their own field of study.

Introducing MCYeast

Our project this year is named "MCYeast: Stress-based detection and enzymatic degradation of the cyanobacterial toxin microcystin." Our goal is to build a two-part system to detect and then degrade cyanobacterial toxins known as microcystins (MCs). Our detection system is based on the natural oxidative stress response of the yeast *Saccharomyces cerevisiae*. Exposure to MCs is linked to higher levels of oxidative stress, and we will couple this response to the expression of yellow fluorescent protein. Thus, fluorescence levels will indicate the amount of MCs present in a sample. To understand and validate our MC detection mechanism, we will also create mathematical and molecular models. For degrading the detected toxins, we are expressing and purifying the enzyme microcystinase (MlrA), which is naturally found in some gram-negative bacteria. The enzyme renders the MC harmless by modifying its structure.



Collaboration

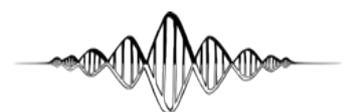
CollabSeeker

One of the judging criteria in iGEM is to do collaboration with other teams. It is a great opportunity to communicate with and learn from different teams all over the world, but how do you find teams with similar visions regarding collaboration? Since in our experience iGEM currently lacks an easy collaboration platform, we at Aalto-Helsinki decided to step up and make one!

We are introducing CollabSeeker, a search engine where you can find all of this year's projects with different keywords. Your team can also login with your team's facebook or twitter account and edit your own team page in order to tell others what kind of collaboration you are looking for, and also to provide the page with additional contact information, something that has till now been quite hard to find.

You can find more detailed information about the CollabSeeker and how to use it from our blog, which has step by step directions on how to use the platform (the blogpost about CollabSeeker: <https://tumblr.co/ZT7snk2AIIIKB>).

You can read more about our team members at www.aaltohelsinki.com. Check also our blog about what is happening in Aalto-Helsinki: <http://blog.aaltohelsinki.com>.



Ben-Gurion University of the Negev

BGU_ISRAEL



@igembgu2016



iGEM BGU



igembgu2016@gmail.com

This year we chose to concentrate our efforts in environmental preservation in an attempt to counteract the ongoing and destructive results of pollution resulting from plastics. Plastic's chemical and physical attributes make it very hard to decompose.

Team

Hello, We are the BGU iGEM Team and our goal is to devise several approaches using synthetic biology tools for efficient plastic biodegradation using bacteria. In addition, we plan to utilize the high energy stored in polyethylene terephthalate (PET) molecules, for electricity production. We have created a game which illustrates the basic idea behind our project, our bacteria secretes LC-cutinase an enzyme found to degrade PET and uses it's biocatalysis products to create energy.



Project

This year we chose to concentrate our efforts in environmental preservation in an attempt to counteract the ongoing and destructive results of pollution resulting from plastics. Plastic's chemical and physical attributes make it very hard to

decompose. Consequently highly toxic waste is accumulating in alarming rates the whole world over. A plastic bottle will decompose in a period of an astonishing 500-700 years, with the slow decomposition rate deriving from the fact that plastic materials are synthetic and considerably new. Organisms have yet to develop an effective way to sustain themselves from those materials as a sole carbon source.

Today there are three main ways of disposing plastic waste:

Recycling - although awareness of recycling has increased considerably over the years, it remains an ineffective solution. Because there are only particular plastic types which are recyclable, plastic waste must be sorted manually, which is expensive and slow in comparison to manufacturing new materials.

Burying plastic waste - a slow degradation process accrues in which greenhouse gasses such as methane are emitted into the atmosphere and contribute to global warming. Furthermore toxic compounds are released into the soil and water sources that can affect public health. The areas in which plastic waste will not cause ecological damage are limited, and plastic disposal in the western world is declining.

Controlled combustion and incineration - If plastic waste is not incinerated and disposed of properly, harmful amounts of toxins can be released and dispersed through the air or waterways.

Our solution – Biological Remediation

For many year scientists have tried to find ways to disintegrate plastic polymers. Some of the most promising research work tries to utilize microorganisms in order to degrade plastic, but they are often challenged by the slow rate of the process.

In our project we are trying to find a solution by means of experimental evolution and “intentional evolution,” referring to genetic engineering and enzyme improvement by mutations. The goal of the research is to create a bacterium that could utilize plastic as a sole carbon source. In addition we hope to show a potential of making electrical energy by creating fuel cells based on the redox reactions taking place as part of the plastic degradation, forming a self-producing system.

In contrast to other solutions available today, our solution does not cause pollution and is relatively inexpensive.

Abit about our project

In order to achieve our goal, different courses of action were chosen:

(i) An Organism Evolution Approach – One of our approaches is to use an organism which has adapted into "finding a solution" to use plastic, and try to improve that "solution" using methods such as experimental evolution and serial passaging. We have chosen to improve a bacteria that was found to degrade polyethylene called *Rhodococcus ruber*.

(ii) A Protein Engineering Approach – Another approach we have adopted, is the engineering of a protein. We chose an enzyme that was found to be one of the most efficient enzymes in breaking down PET polymers into degradable substrates, this enzyme is called LC-cutinase. Based on the enzymes structure that was solved, we have chosen to use a rational mutagenesis approach for our experiment. After the process was completed we have ended up with 4 different variants other than the WT. These mutations are expected to improve the enzymes activity.

(iii) Genetic Engineering of Metabolic Pathways – we will modify two metabolic pathways using engineered enzymatic cascades which lead to two

products, terephthalate and ethyleneglycol resulting from the of PET by LC-cutinase. While the terephthalate will be used to produce succinyl co-A and acetyl co-A, the ethylene glycol will be transformed into malate by using a metabolic pathway that already exists in *E. coli*. This way, our engineered bacteria of choice, *Pseudomonas putida*, will degrade PET and will transform the electrons released from PET degradation into energy by using PET as the carbon source.

(iv) Microbial Fuel Cells - Since PET is a polymer that contains high energy bonds in its carbon-carbon bonds, excess energy released by our engineered microorganisms from carbon-carbon bond degradation will be harnessed and utilized in microbial fuel cells devices, this way plastic biodegradation will be converted into energy.

Thank you for spending your time reading a bit about our project and we hoped you enjoyed our game!

If you think your project could have some sort of collaboration with ours we would love to hear from you!

Human Practice

a link to our game: <http://bgu-plastikiller.netne.net/#> (PC use only)



Beijing Institute of Technology

BIT-CHINA



@iGEM_BIT



igem_bit@outlook.com



iGEM_BIT



iGEM_BIT

iGEM_BIT is an iGEM team that organized by Beijing Institute of Technology, School of Life Science. It is a scientific group that researches the natural of life science. Our team aims to make undergraduates have an all-round development, and we underline the comprehensive quality and ability of scientific research as the core of our team.

Team

iGEM_BIT is an iGEM team that organized by Beijing Institute of Technology, School of Life Science. It is a scientific group that researches the natural of life science. Our team aims to make undergraduates have an all-round development, and we underline the comprehensive quality and ability of scientific research as the core of our team. Team BIT participate in iGEM competition for the first time since 2013, we have been involved three session of the competition, and achieved high grades. After three years of development, Team BIT has gradually formed a feature that we see synthetic biology as the core, then fuse many subjects as supplementary

like: Biological chemistry, Biological medicine, Electronic information science, Computer science, Mathematics and so on.

*OUR MEMBERS

Team BIT is a passionate team which is composed of 19 undergraduate students with different majors. With the shared interest in synthetic biology, we gather together and want to make a difference in this field. The team is supervised by Prof. Deng Yulin, Prof. Li Xiaoqiong, Prof. Lv Xuefei and Dr. Quan Zhenzhen. We are excited to share what we are doing with leading universities and students around the world.

Project

In a short time, and provide patients with reference significance testing data, and the testing costs can be drastically reduced alleviate the contradiction of medical treatment system. In recent years, breast cancer has become one of the highest incidental cancer. Many patients missed the best diagnosis and treatment time and them made patients lost their lives. Studies have shown



that when we have breast cancer, the expression degree of microRNA-21 and microRNA-155 in the human body will be significantly higher than the normal human body. Based on the above background, this project applies the artificial designed biological system to realize the detection of expression levels of microRNA (associated with breast cancer) in the serum environment, the concrete realization method is: Detect microRNA expression level in the serum samples through the engineering bacteria(directional transformed) , the higher microRNA expression degree can produce green fluorescent protein, then using miniaturized signal hardware to detect fluorescence, and through the mathematical modeling of the model to calculate microRNA expression level. Finally deduce the theoretical diagnosis to user and this project is a real-time inspection system for breast cancer detection.

Collaboration

*Searching

In the last year we had collaborated with USTC because we faced the same difficulties in experiment, modeling and hardware. This year our team choose the DIAGNOSTICS as our prime track and we are still looking forward for collaboration with any team who are interested in diagnostics, detection, model, hardware and software. If anyone else would like to collaborate with our team we will be pleased to hear from you. If yours project is similar with us and want to discuss the problems, please do not hesitate to tell us! We are waiting for your connection.



Human Practice

Our project aims to create a real-time device connect with software to detect the specific microRNA in the human body. So besides our experimental work, we reached out of the lab and did some human practice activities. Our human practice consists of online and offline. In the first part we will make a questionnaire which aims to realize practicability of the project, through some social software like Wechat Platform, Facebook and Twitter. In the mean time we can find out how well people know about the breast cancer. In the second part we are going to make a survey in schools, biotechnology companies and tumor hospitals. We will also communicate with technical personnel about our project. By listening to them, we can solve the problems we have meet.

Discussion

Our chassis organism is E. coli (Trans 5 alpha Chemically Competent Cell and Trans1-T1 Phage Resistant Chemically Competent Cell). We have several preventive measures for potential risks: 1) Wearing rubber gloves. 2) Handling bacteria experiments with alcohol burner. 3) Sterilizing bacteria by using 84 disinfectants. 4) Using specific sewer to deal with bacteria waste. As for the product in real world, the bacterium will be made into dry powder by the method of freeze-drying. And the powder will apply to the chamber of micro-fluidic chip. When users need to detect certain disease, users can add their blood sample into the chip and insert chip into the hardware we designed so that results will be simply shown.

The University of Wales

CARDIFF_WALES



@iGEM_Cardiff



golshaiea@cardiff.ac.uk

We will assess the viability of a novel bioluminescence detection system for point-of-care diagnostic testing.

Team

Team members (all 2nd years/going into third year after summer):

Name- Age- Subject

- Christian Donohoe- 20- Chemistry
- Rob Newman- 21- Genetics
- Asal Golshaie- 20- Genetics
- Andrew Brimer- 20- Biomedical Sciences
- Laura Bird- 20- Biology
- David McMaster- 20- Biomedical Sciences
- Nikolas Demetriou- 23- Chemistry

Our supervisors:

Primary PI: Dr Geraint Parry
Secondary PI: Dr Amit Jathoul
Advisor: Jamie Long (Ph.D. student)

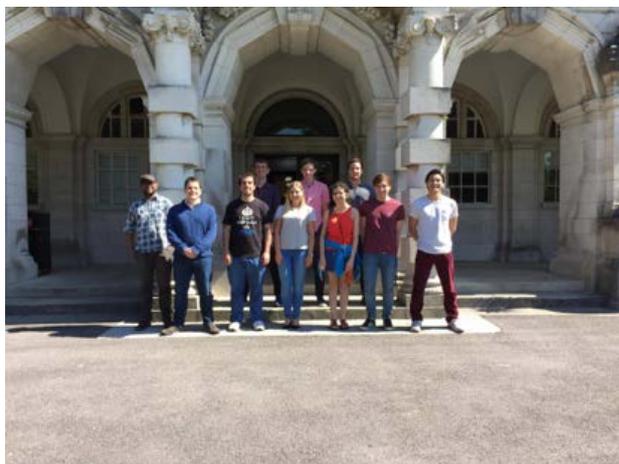


Photo:
Back row, left to right: Geraint, Jamie, David
Front row, left to right: Amit, Andrew, Nikolas, Laura, Asal, Rob, Christian

Project

We will assess the viability of a novel bioluminescence detection system for point-of-care diagnostic testing. In our proposed system, a *Streptococcus pyogenes* dCas9 isoform codon optimised for *Escherichia coli* is fused to the N-or C- terminal fragments of a thermostable pH-tolerant *Pyrophorus plagiopthalmus* luciferase (LUC).

We aim to coexpress guideRNAs to target these chimeric dCas9-LUC proteins to adjacent DNA sequences. This will enable the reconstitution of luciferase activity and subsequent bioluminescence in the presence of luciferin. This light output constitutes a signal for detection of any targeted DNA sequence and is extremely adaptable, dependent on access to the target sequence. We plan to undertake a proof of concept study of this system using gRNAs targeted to the E.coli 16S rRNA locus, aiming to describe both the effective output of this system in vitro, and the optimum distance between gRNA targets. Finally an important aspect of this study will be to investigate the feasibility of this diagnostic system as a clinical test, potentially using cell-free components.

Human Practice

We have been exploring GM regulations, getting expert advice on the design aspects and applications of our diagnostic, and setting up public engagement events for September and October. We'll be doing a talk with Cardiff's Science Cafe in October, taking part in the Live Mars event at Cardiff University with an exciting activity (using 3D printed E.coli and luciferase), and setting up a stall at the Biology Rocks event in the National Museum in October. We are also in continued talks to set up a stall and demonstration at Techniquest. The GM regulations issues we faced with this endeavour inspired us to look deeper into safety, and GM regulations. We have had displays at an open day and the STEM conference at Cardiff University for sixth formers. Two group members represented us at a European iGEM teams meet up in Paris, and two members will also be attending the UK iGEM teams meet up at Westminster this week (17th-18th August).

Chang Gung University

CGU_Taiwan



@iGEMCGU_Taiwan



CGU iGEM Taiwan



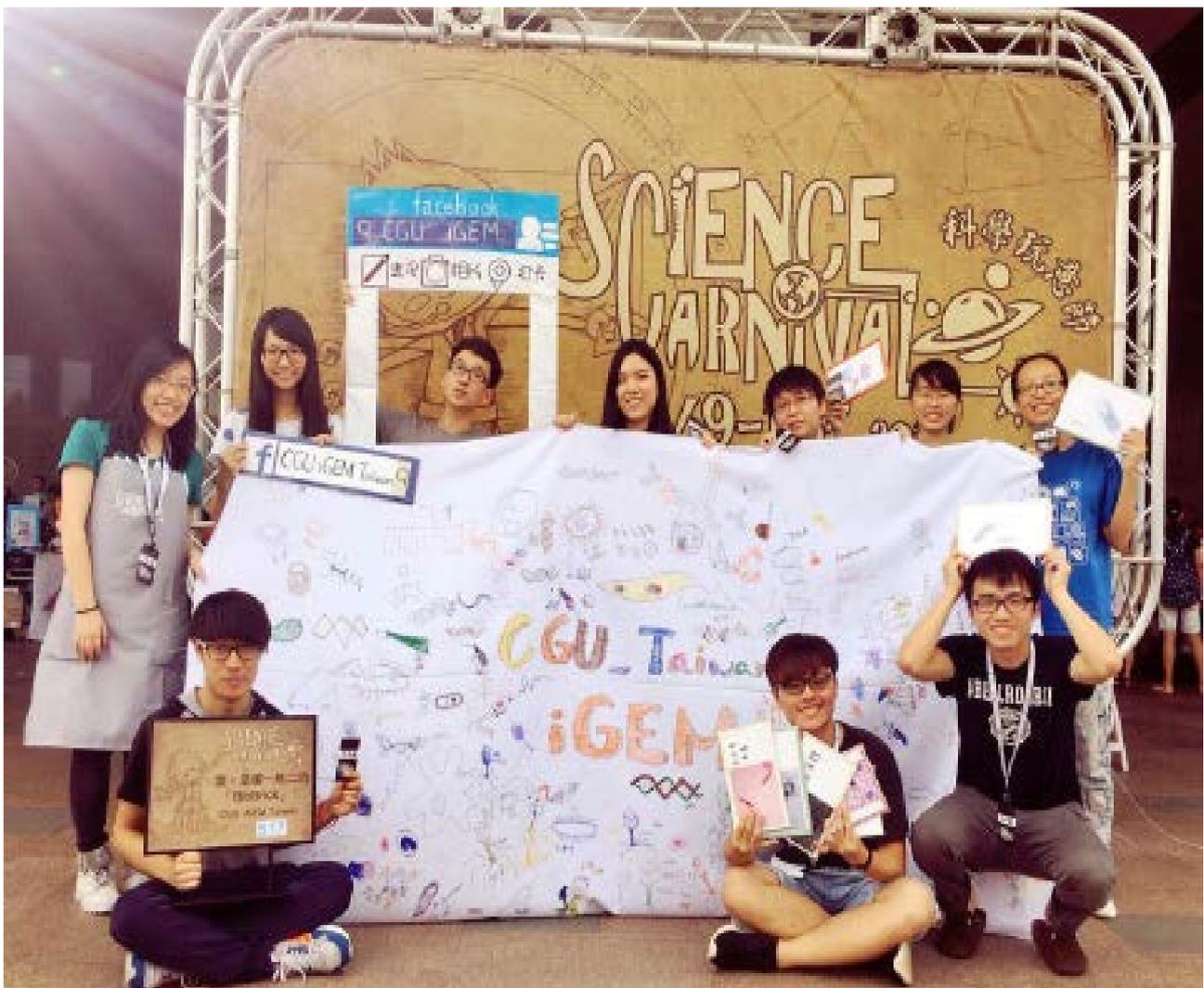
igemcgu@gmail.com

Our team consists of 12 members majoring in biomedical science and electronic engineering to represent CGU participating in iGEM. We are resourceful, thriving on challenge, and eager learners. Everyone brings what he or she has learned into full play.

Team

Our team consists of 12 members majoring in biomedical science and electronic engineering to represent CGU participating in iGEM. We are resourceful, thriving on challenge, and eager learners. Everyone brings what he or she has learned into full play.

After interdisciplinary brainstorm, we hope to contribute to the society and world by developing a new adjuvant delivery way during the iGEM project. We also expect to communicate with other teams and, through science, build up a friendship with lots of iGEMers!



Project

*What is the context of this research?

Leishmania, a trypanosomatid protozoan, are aerobic organisms, relying on oxidative phosphorylation, but are defective in the synthesis of heme. The genetic deficiency of heme biosynthesis in *Leishmania* makes it possible to produce transgenic mutants (DT), which are inducible with delta-aminolevulinate (ALA) for accumulation of uroporphyrin I (URO) as an endogenous photosensitizer. Another photosensitizer, PC, will also be loaded exogenously. After URO and PC are illuminated by specific wavelength of light, double inactivation will kill *Leishmania* thoroughly with proven effectiveness. Professor Kwang Poo Chang established the double photo inactivation system of the *Leishmania* and validated its possibility as a cancer vaccine.

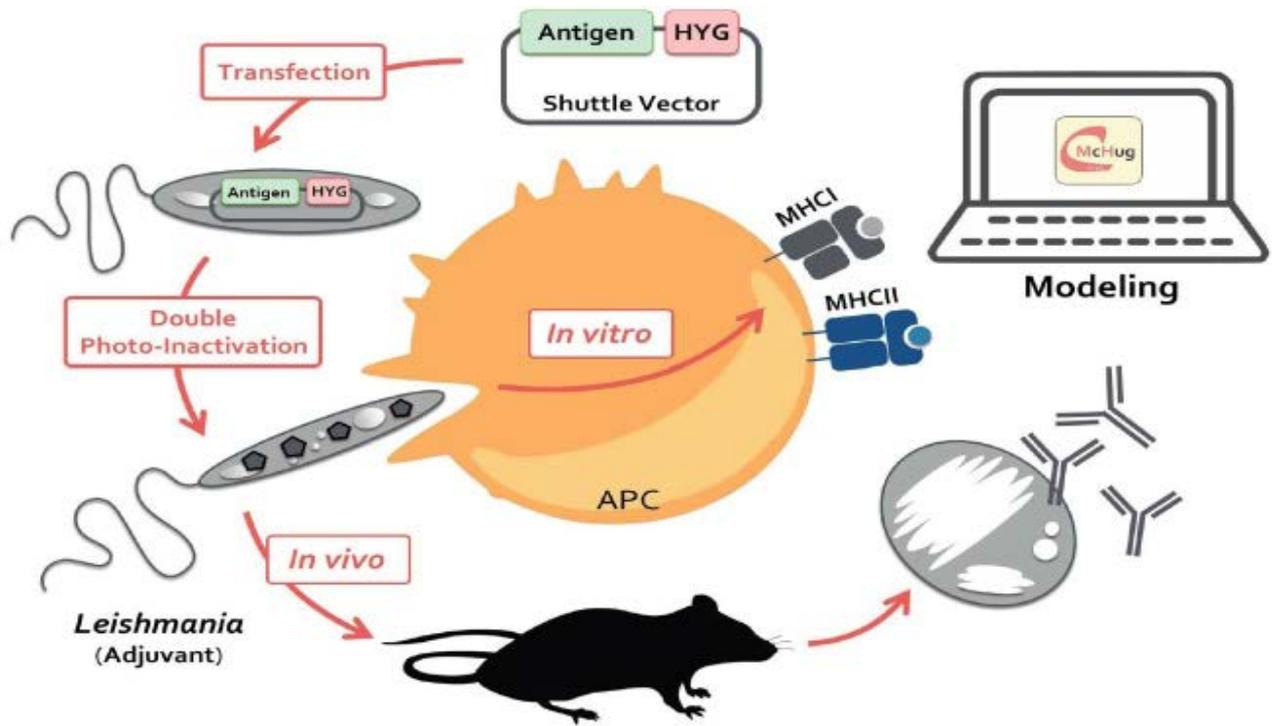
*What is the significance of this project?

Vaccine provides active adaptive immunity to a particular disease and most commonly used vaccines can be part into two big categories, live attenuated and inactivated vaccines. An inactivated vaccine consists of pathogens which are grown under controlled condition to kept them non-infectious and then killed to completely remove its infectiousness. Since inactivated pathogens tend to induce a weaker immune response than live ones, immunologic adjuvants are required to provide an effective immune response against the inactivated pathogens. Here we aim to introduce an effective, safe and pathogen specific adjuvant.

*What are the goals of the project?

Leishmania possess the advantages as a potential vaccine adjuvant, such as antigen presenting cells (APCs) recruitment, pattern recognition receptors (PRRs) activation, inflammasome activation and activation of MHC-presenting pathway. Genetically-engineered *Leishmania* that can be inactivated by light exposure acts as a safe carrier to deliver specific antigens to the APCs for T cells and humoral response. Based on this concept, we established a new model system to generate antigen-specific *Leishmania* adjuvant--Leijuvant. We will perform several experiments to check if it is a potential vaccine adjuvant.

Here we aim to design an *E.coli-Leishmania* shuttle vector constructed under biobrick standards to provide a standardized shuttle vector for our own experiment and for others' future application.



To test the efficiency of the antibody immune response of the photo-inactivated *Leishmania* as a vaccine adjuvant, we will co-inject ova recombinant protein and photo-inactivated *Leishmania* that is genetically modified to present OVA protein into mouse. Serum will be collected every 5 days after the second injection to test the antibody immune response with Anti-OVA ELISA. The outcome will be compare to the of Alum adjuvant.



Heinrich Heine University

DUESSELDORF



optoptosis

HHU Düsseldorf



@iGemHHU16



iGEM HHU



igem@hhu.de

This year we established the first iGEM-team at the Heinrich Heine University in Duesseldorf in Germany. Our group consists of 20 undergraduate members, who came together to found the team in late 2015.

Team

This year we established the first iGEM-team at the Heinrich Heine University in Duesseldorf in Germany. Our group consists of 20 undergraduate members, who came together to found the team in late 2015. We all joined iGEM with the motivation to apply and create synthetic biology. Already in the early stage of our project it was definite that we would emphasize on creating a mechanism to fight cancer.



Project

The essentials of OPTOPTOSIS

The global increase of cancer cases and the lack of highly specific therapies for various types of cancer states a clear message: New approaches have to be found, which allow a level of precision in cancer treatment that has been long-awaited. This is the goal of OPTOPTOSIS.

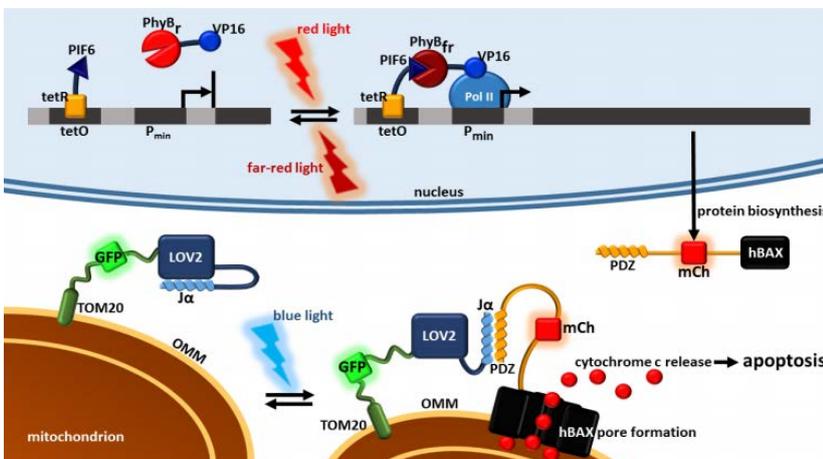
In contrast to human cells, cancer cells have lost the ability to recognize their own devastating impact on the organism. Thus they do not induce apoptosis, but rather show uncontrolled growth leading to severe consequences for health. With our optogenetic system we intend to reestablish the natural balance between life and death, proliferation and apoptosis, in impaired tissue. The future objective of this research is to implement our system in tumors via virotherapy. Therefore our

idea represents the perfect complement to the already innovative virotherapy.

Our approach aims at achieving high spatiotemporal control of apoptosis in tumor cells by applying an optogenetic double-killswitch. This system combines clean removal of cancer cells through apoptosis with the precision of light-controlled optogenetics. We utilize two optogenetic proteins, namely Phytochrome B and LOV2. The red light switch based on Phytochrome B controls expression, while the blue light switch LOV2 controls localization of apoptotic proteins to their target site.

The optogenetic induction of apoptosis in the budding yeast (*S. cerevisiae*) should serve us as a model and proof of concept for the future application in cancer cells. The application of optogenetic switches enables us to induce extremely precise and multiply regulated killing of cells. Thereby this

system will represent an improvement in comparison to conventional, less target-specific methods. Furthermore we aim to express this optogenetic system within mammalian cells.



The University of Edinburgh

ENDINBURGH_UG



BABBLED



@EdiGEM2016



EdiGEM2016



edinburgh.igem2016@gmail.com

BabbLED – The DNA Typewriter: Building a modular system to encode information in DNA

Team

We aim to develop a modular system for encoding any unit of information, into DNA. We will prove the validity of our concept by encoding Ogden's Basic English (a collection of 8501,000 words that can be used to express most concepts in the English language). Each encoded word; a BabbleBrick, will be stored in a different Phytobrick. Sentence assembly and unidirectionality is ensured by the stepwise addition of BabbleBricks that have alternating types of sticky ends; this prevents repeats and minimizes the occurrence of missing words. The whole sentence construct, or BabbleBlock can be melted off for easy retrieval and assembled back into a PhytoBrick for storage. Since the value that is assigned to each BabbleBrick is arbitrary, each one can be reused with any library or language. In this way, our encoding and assembly method can be optimized for many types of data. Furthermore, using a variety of error correction and encryption techniques we will provide an exceptionally secure and high fidelity storage medium.

*Team Members



Petar Iliev (aka Pepy):

I am a 2nd year Biological Chemistry student and in my spare time I do kickboxing and listen to exuberant music. I am from Bulgaria and I enjoy socialising, dancing and hiking. I am passionate about iGEM, because synthetic biology is a field where nature is utilised to complement itself and great ideas make a huge impact.

Nikita Lazaroo:

I am a 2nd year Psychology student from Australia, with a strong interest in biology and cognitive science. In my spare time I enjoy kickboxing, cooking and travelling. I am enthusiastic about participating in iGEM and exploring the field of synthetic biology as they are both aimed towards the advancement of biology as an interdisciplinary science that is accessible by all. The potential for iGEM projects to provide solutions for complex social issues is what initially drew me to the competition, and I'm thrilled to be a part of the Edinburgh team.





Azzurra Laura De Pace:

I am a 2nd year Biology student from Italy. I am very interested in Epigenetics and personalized medicine. In my spare time I love reading books outdoors and cooking. I am very happy to be part of the Edinburgh iGEM team. Taking part in the iGEM competition gives students the chance to reinterpret nature through the power of synthetic biology.

Catalina Rotaru:

I am a 2nd year Computer Science student from Romania and I am deeply interested in computer security and cryptography. In my spare time, I enjoy boxing, dancing, learning russian and watching CSI series. I am very excited to be taking part in such a massive engineering competition. iGEM represents the challenge to find solutions to worldwide problems. Moreover, it brings people of different backgrounds together to exchange knowledge and experience.



Freddie Starkey:

I am a second year Informatics student from Aberdeenshire. My main interests are in Bioinformatics and Artificial Life. I enjoy Nordic Skiing and Kayaking in my spare time as well as being a massive Doctor Who fan. I am excited to have the opportunity to work in the wet lab for a change during iGEM as well as to contribute to a project with such large real world applications.

Rosie Maddock:

I am a third year Biology student from Aberdeenshire. I am interested in biochemistry with a particular interest in disease research. In my spare time I play judo, go to the gym, and love to cook. I am really excited to be part of iGEM because it provides so many people the opportunity to develop and explore their synthetic biology ideas. The flexibility with iGEM projects allows input from different subject area backgrounds, which gives rise to many exciting lines of research.





Alexandra Bisia:

I am a student from Crete, Greece, who just finished her second year of studies in developmental biology. I am keenly interested in foreign languages, exploring the beautiful city I study in. It's an incredible experience being part of the Edinburgh iGEM team, as the competition has given us the opportunity to push back the boundaries of biology and apply our ideas to create novel solutions to current problems. I have also become familiar with different subjects and interdisciplinary collaboration, which I believe to be the future of research.

Brendan Largey:

I am second year Biochemistry student from New York. In my spare time I enjoy restoring antique furniture, teaching dance, and making use of the Oxford comma. I also enjoy playing the violin and other instruments, and can bake a mean kartoshka. I have been enjoying every moment with the team so far and love interacting with such a diverse group of individuals.



Patrick Lim :

I am a second year Molecular Genetics student from Bristol. I do a bit of programming and writing and am very excited to be helping out with Edinburgh iGEM. When I am not working with the team I do gardening at a hospice. During my time here I have learned a lot about science and have become a big fan of falafel wraps despite being skeptical at first.

Collaboration

Newcastle iGEM – We are collaborating over a simulation game designed to help people consider the ethics of both our projects. Additionally we will create DNA

'labels' for Newcastle's biocircuits and they will be checking our BabbleBricks for Biobrick compatibility.

Dundee iGEM – Cohosting a debate on GMO policy in Scotland between our respective university debating societies.

National Library of Scotland – We will be encoding Mary Queen of Scot's historic final letter before her execution into DNA format for the libraries archive.

Finally, we are always looking for new collaborators and suggestions on how to improve our project. In particular we are creating Biobricks for Dps to provide resistance to DNA damage from radiation and would love to hear from any teams interested in utilising this.

Human Practice

In 2014, over 10 sextillion bits of data were digitally stored worldwide. To put this in context, there are only 1 sextillion grains of sand on this entire Earth.

In 2020 data storage demand will reach 44 trillion gigabytes. In 2040[1] global memory demand will reach 3 billion billion million bits; the silicon required to store all this data vastly exceeds total silicon supply. Nowadays, data centres consume 1.52% [2] of all global electricity.

There is a dire need for new storage methods. DNA is a stable, dense and long-lasting molecule, making it perfect for storing data. Recently, researchers at Microsoft [3] have stored 200MB of data in DNA. However, their methods, using de novo synthesis are expensive, inflexible and inaccessible to the general public. We aim to create a cheaper and hence more accessible DNA storage method that will hopefully encourage the proliferation of this incredible technology in the future.

References

[1] <http://science.sciencemag.org/content/332/6025/60.full>

[2] <http://www.greenpeace.org/international/Global/international/publications/climate/2011/Cool%20IT/dirtydatareportgreenpeace.pdf/>

[3] <http://blogs.microsoft.com/next/2016/07/07/microsoftuniversitywashingtonresearcherssetrecorddnastorage/#sm.0000isbfbb6foeztu131v6ddlj9m4>

Discussion

To enable BabbLED to be as widely used as possible we have had to consider a number of important legal and safety points. Firstly due to the restrictions surrounding the use of GMOs outside of a laboratory we made the decision early on that our system should be cell free with our DNA messages instead being stored in solution. This means that we can send or store our BabbleBlocks without concerns over breaking these strict regulations. Additionally to combat the eventuality where one of our encodings needs to be used in an organism, for example in labelling a construct, we've added a 17bp stop codon region into every BabbleBlock to prevent the creation of any unexpected or potentially dangerous proteins.

Life Sciences at EPF Lausanne

EPFL



@EPFL_iGEM



igemepfl



igemmers2016@groupes.epfl.ch

We are a team of nine second year of bachelor students in Life Sciences at EPF Lausanne who decided to work on automating the design and construction of biological circuits with CRISPR-dCas9.

Team

We are a team of nine second year of bachelor students in Life Sciences at EPF Lausanne who decided to work on automating the design and construction of biological circuits with CRISPR-dCas9. Using our knowledge from our lectures in Life sciences, we started the iGEM competition with much zeal : first brainstorming, then the first steps in the lab and the results that followed... Even if they are not always what we hoped for, it didn't and will never get us down! If you visit us in Switzerland, between chocolate and mountains, you will find wild students from different countries

and even continents but with a single goal: to represent Lausanne and Switzerland at the Giant Jamboree.

When we are not in the lab we are often together on the banks of Lake Geneva making barbecues... Team spirit is key, and is as important as the rest.

To follow our summer journey through the iGEM competition, you can follow us on Twitter @EPFL_iGEM or Facebook @igemepfl.

See you in Boston !

Project

Introduction

CRISPR-Cas9 has already revolutionized synthetic biology. To build upon this development we aim to implement digital-like circuits in yeast using a CRISPR-associated RNA scaffold system (Zalatan et al, 2015). Furthermore, a recently



published study unveiled Cello, a modular program automating the design of artificial transcriptional circuits in *E. coli* (Nielsen et al, 20162). As a proof of concept we will modify Cello to use our dCas9 transistors in yeast for a so-called half-adder system, using AND and XOR gates, that we can experimentally assess. With this approach we aim to pave the way for even more complex biological circuits in yeasts.

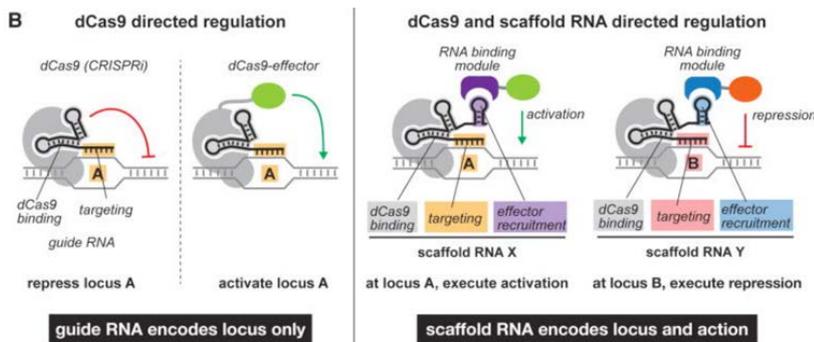
CRISPR/dCas9 use until now

The conventional method of using the CRISPR dCas9 system in the construction of biological circuits is to fuse dCas9 to VP64, an activation unit, in order to activate or repress gene expression. With this system, only one effector protein can be used, meaning that an activator or repressor will have to fill both roles functionally, its action being determined by its placement on the DNA. For example, a gene may be repressed by an activator-fused dCas9 purely through steric hindrance by placing dCas9 on the gene.

What's new with our approach ?

One of the novel aspects of our system, with respect to the "classical" dCas9 based DNA circuit, is that the effector proteins are recruited by a scaffold guide RNA (scRNA) through an RNA binding module.

In our system, the guide RNA bound to dCas9 serves both as a targeting sequence and an effector protein recruitment element. Contrary to the conventional system, this allows us to have both transcriptional activators and repressors in the same system. In addition, it has been shown that the activator VP64 was more effective when bound to the scaffold gRNA than when fused to dCas9 directly (Zalatan et al., 2015). Interestingly, the scaffold allows us to target two different effectors to the same DNA target, allowing the combination of different repressors, activators, or even activators and repressors together.



Comparison between classical dCas9 system (left) and scRNA-based system (right). Zalatan et al, 20151.

To go further

With the creation of our transistors, it will be possible to construct any logic gate, such as XOR, AND and NOR. By concatenation of these gates we can construct complex biological circuits, which could be used as biosensors or detection methods for diseases or substances. They could even act in coordination with circuits naturally present in organisms and react to external factors producing different proteins as a response.

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[1] Zalatan JG, Lee ME, Almeida R, Gilbert LA, Whitehead EH, La Russa M, Tsai CT, Weissman JS, Dueber JE, Qi LS, Lim WA. Engineering Complex Synthetic Transcriptional Programs with CRISPR RNA Scaffolds. *Cell*. 2015 Jan 15 ; 160(0) : 339-350.

[2] Nielsen AAK, Der BS, Shin J, Vaidyanathan P, Paralanov V, Strychalski EA, Ross D, Densmore D, Voigt CA. Genetic circuit design automation. *Science*. 2016 Apr 01 ; Vol. 352, Issue 6281. DOI: 10.1126/science.aac7341.

the University of Evry Val d'Essonne

EVRY



@iGEM_Evry



iGEM.Evry.2016



evryigem2016@gmail.com



igem_evry

We are 12 students from different nationality and different backgrounds (biology, biochemistry genetic, bioinformatics and law) from the University of Evry Val d'Essonne (France) and we are helped by two advisors.

Team



We are 12 students from different nationality and different backgrounds (biology, biochemistry genetic, bioinformatics and law) from the University of Evry Val d'Essonne (France) and we are helped by two advisors.

Here is some links:

To our Survey: <https://goo.gl/forms/O3qRm8V2fpWC83lK2>

Facebook: <https://www.facebook.com/iGEM.Evry.2016/>

Twitter: https://twitter.com/iGEM_Evry

Instagram: https://www.instagram.com/igem_evry/

E-Mail: evryigem2016@gmail.com

Project

*Key figures

280 million tons: it's the average amount of plastic produced each year worldwide (which represents 8,880 kilograms per second)

1,000 years: it's the average lifespan of a plastic bottle

Up to 450 years: it's the time needed for the degradation of a plastic bag in the environment. For these reasons, our team thought about working on a biological alternative to the plastic: the PLA!

*What is the PLA?

PLA, or Poly-Lactic Acid, is a totally biodegradable polymer and a thermoplastic. It's currently used as food packaging but also in other applications such as the sutures in surgery.

PLA has a lot of advantages. Indeed, it is considered that a bottle composed of PLA takes 80 days to be disintegrated instead of 1000 years for a classic plastic bottle. Moreover, PLA has good optic qualities such as transparency and brilliance, good properties for protection against oils and gases (O₂ and CO₂) allowing it to be an intermediate for different mass market polymers and an alternative to current plastics.

*Our objective:

Our objective for iGEM 2016, is to produce a significant amount of PLA solely through a biological way by engineering *Pseudomonas putida*.

PLA bioproduction presents several benefits compared to chemical synthesis: it uses simple carbon sources and it is inexpensive. On the other hand, *P. putida* is a safe organism reported to be efficient for polymerization, which gives advantages over other possible chassis. By modifying its metabolic pathways, we aim to improve PLA biosynthesis yields in a sustainable manner, and to determine the usability of our bioplastic by manufacturing a vesicle for drug delivery.

The University of Freiburg

FREIBURG



igemfreiburg2016



igem_2016@freigem.org

This year's iGEM team consists of 16 students from many distinct scientific fields including biology, chemistry, medicine, biochemistry/biophysics and informatics.

Team

We? Who is that?

16 students from Freiburg, Germany.

This year's iGEM team consists of 16 students from many distinct scientific fields including biology, chemistry, medicine, biochemistry/biophysics and informatics. Our team is supervised by Jun. Prof. Maximilian Ulbrich and Dr. Nicole Gensch, both being part of the scientific staff of the BIOS Centre for Biological Signalling Studies. We also get advice and support from Luisa Keilholz and Adrian Fischer, some former iGEM participants who are sharing their experiences and knowledge with us.

Project

Nanocillus 'cause spore is more!

The treatment of diseases while avoiding systemic side effects is still a major obstacle in modern medicine. After administration, conventional drugs are distributed throughout the whole body thus affecting both, diseased and healthy cells. Current strategies on targeted drug delivery are mainly based on the applications of antibodydrug conjugates or nanoparticles. However, both approaches revealed considerable challenges in their application due to short halflife and expensive production, respectively.

We develop a novel platform for targeted drug delivery by implementing highly specific nanobodies directed against surface markers of affected cells. The combination with an enzymatic functionality facilitates the local activation of prodrugs, thus preventing unnecessary side effects by systemic drug dispersal. By engineering the spores of probiotic *Bacillus subtilis*, a member of the human microbiome, we establish a lowcost carrier for welltolerated treatment.

Collaboration

Searching for collaborations

We are working with *Bacillus subtilis* and GFP nanobodies. So if you have any thoughts concerning those two components we are happy to collaborate. Also if you need any advice, we are eager to help you!

Human Practice

This year, Freiburg's iGEM team intends to spread the awareness of synthetic biology in the public. It doesn't only mean to speak to students and academic people about our project but also includes talking to people of non-scientific professions.

We've started a collaboration with the university's radio program to share our visions and ideas with students through a radio interview. We talked about iGEM, synthetic biology and the basics of our project. Our motivation is to get

the synthetic biology and the iGEM competition closer to public. We believe that the state of the art of scientific research is a rapid development that also concerns students from other majors. For example, do the history students of your university know anything about promoters, the bacteria used in drug development or even in their dairy products?

Furthermore, we want to find out what the people think about our application. While working in the lab is essential for

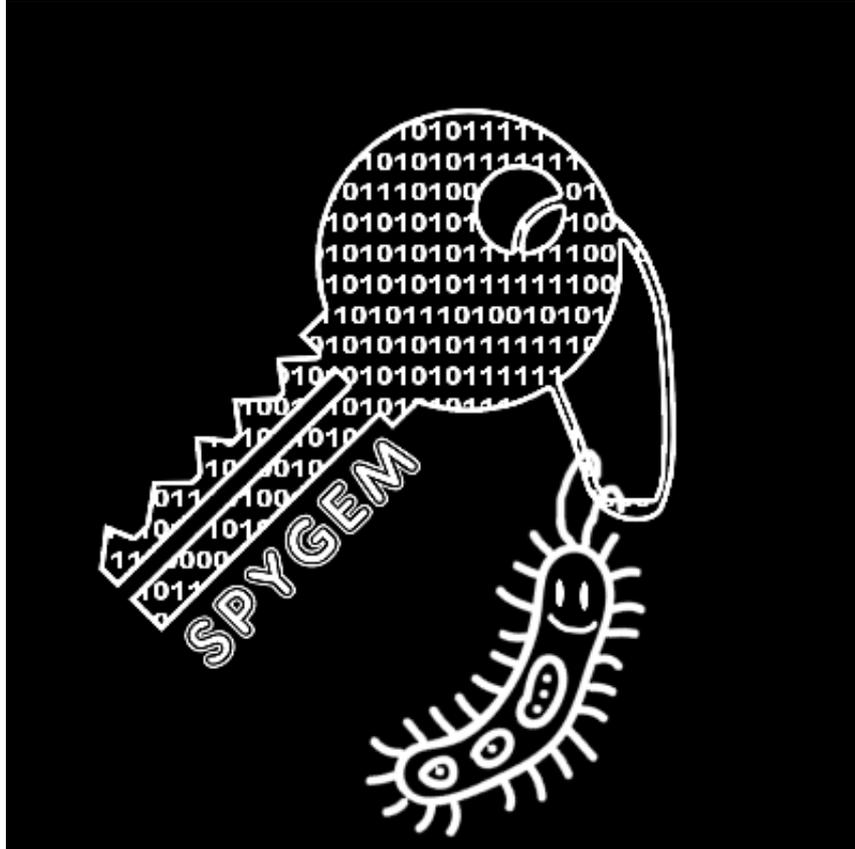
success and progress in your project, you might forget to think the usefulness for consumers and patients or about what your product will look like under real life conditions. If you develop an assay, will it be practical enough to be used in an efficient way? If you improve a drug, will it be affordable for the public? Would people have trust in a medical product with bacterial spores? Are there any drawbacks you didn't think of? Find it out – ask people who

are not professional scientists. Their opinions and suggestions will make your project fit better to public expectations. Since our project has a high medical relevance, we are also interested in the professional medical input doctors and scientific clinicians can give us. Therefore, we've conducted interviews with dermatologists and gastroenterologist who helped us to get new input.



University of Groningen

GRONINGEN



igemgroningen2016



ilonamager@outlook.com

We are the first iGEM team from Groningen!

Project

***Protecting data through encryption and storage in bacterial spore DNA.**

In 2002, the amount of information stored digitally had eclipsed information stored in analog format for the first time. Just five years later, only 6% of the world's data was still analog. In 2015, an estimated 2,500,000,000,000 megabytes of new data were created every day, and this number is growing at an increasing rate. It is not surprising that data breaches orchestrated by hackers are on the rise as well. Financial and legal records, military and government documents, these are examples of important information that must be preserved for a long time, but could cause great damage in the wrong hands. We have become a civilization dependent on information, and this information must be stored somewhere. As a result, we are faced with two problems: where do we store all of our data, and how do we keep it safe?

Storage of data in DNA has been proposed as early as the 1960's, but has only recently become a hot topic. This is in part due to the ever-growing demand for data storage, as well as advancements in DNA synthesis and sequencing technologies. Our goal is to create a system for long-term data storage and data transfer which cannot be hacked by digital means. Digital methods of encrypting information and converting it into binary code are well established, and data storage in DNA has already been demonstrated. Our project combines these two approaches by first converting information into binary code, encrypting it, and then storing it safely in DNA. Additional measures based on molecular biology will prevent unauthorized access, ensuring the safety of the stored information.

Our system will be useful for the kind of information that should be stored and transferred in a very secure manner, but does not have to be accessed quickly (within seconds). It will be possible to obtain the message in about 24-48 hours, however, this timeframe is likely to be reduced as new sequencing technologies are developed. For example, this system could be used to store patent and prototype information, genealogical records, legal and financial records, banking account details, login data or even top secret government documents. Given the stability and compactness of DNA, our system could also be adapted to serve as a time capsule for human knowledge.



***Advantages of data storage in spore DNA**

DNA is a far more stable data storage medium compared to magnetic and optical media, remaining intact for at least 700,000 years at -4°C . Even in harsh environments, DNA has a half-life of over 500 years. In contrast, current storage technology lasts only up to 30 years.

Spores are extremely resistant to aging, radiation, heat, and chemical damage. A viable spore-forming *Bacillus* strain was isolated from 250 million year old salt crystals.

The densest data storage medium commercially available today can hold up to 10 GB/mm³. DNA has a data storage density of up to 109 GB/mm³, 8 orders of magnitude higher.

Conservative estimates predict that based on global memory demand, the amount of silicon (required for flash memory) is expected to exceed silicon supply by 2040.

DNA storage will soon become a cheaper alternative for data storage as DNA synthesis and sequencing costs drop. It is estimated to become a cost-effective method for long-term data storage within approximately ten years .

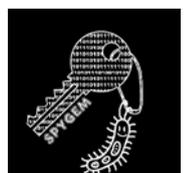
Data storage in DNA is more environmentally friendly than currently used digital data storage. In 2015, 416.2 terawatt hours of electricity were used by data centers worldwide. This is higher than the annual power consumption of the entire UK, and is responsible for approximately 2% of global greenhouse emissions, rivalling the airline industry.

Data stored in DNA cannot be hacked by digital means.

DNA data storage is an apocalypse-proof technology because DNA will be relevant to future civilizations. As long as intelligent DNA-based life exists, there will be compelling reasons to study and manipulate DNA.

***Our approach**

We use a layered approach with a combination of digital and biological security measures to ensure the information can only be accessed by the intended recipient. The first layer is digital encryption. The information is encrypted with the Advanced Encryption Standard (AES) algorithm, converted into a DNA sequence and



integrated in the genomic DNA of *Bacillus subtilis*, a safe, thoroughly categorized organism capable of sporulation. The binary data obtained after encryption will be encoded into DNA according to the following logic: since DNA consists of four nucleotides namely T, A, C, and G, every nucleotide will represent a binary pair (combination of a 0 and a 1). The T will be represented as 01, A as 10, C as 00 and G as 11. The decryption key and the encrypted message are integrated into two different *Bacillus* strains and are protected from unauthorized access with additional security layers.

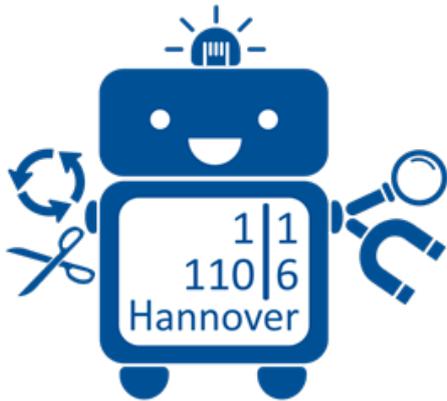
Once the message and key are encoded in *Bacillus* DNA, the cells are cultured in a sporulation-promoting medium. Bacterial spores are among the most resistant biological entities currently known, and thus represent an ideal substrate for long-term data storage. The spores containing the encrypted message and key are freeze-dried and embedded in separate filter papers (or any other porous material) for storage and transfer, along with a spiropyran-ciprofloxacin conjugate. The biological activity of this photoswitchable antibiotic is very low when the spiropyran photoswitch is in its stable closed form, but increases dramatically after irradiation with a specific wavelength of light (in our case, 365 nm) which brings the photoswitch into a less stable, open form. When the light source is removed, the compound slowly reverts back to its biologically inactive state. Irradiation with other wavelengths also results in deactivation. The strains carrying the message and key (which possess resistance to the antibiotic) are mixed with numerous decoy spores when brought onto the carrier material. The decoy spores are not resistant, and do not contain any encrypted information.

When the intended recipients want to access the stored data, they place the filter paper with key carrying spores and antibiotic in a culture medium, and irradiate it with the activating wavelength of light. This wavelength must be known by the recipient beforehand. The activated antibiotic kills the decoys but not our key carrying strain. After culturing, their DNA is sequenced and the key is found. The key contains information necessary to culture the message carrying strain, and to decrypt the message. Without activation, all the spores germinate and grow, including the decoys. This makes it impossible to find the key by sequencing. Once the key is obtained, the message carrying strain can be cultured. Their DNA is then sequenced and the message can be decrypted.



the Leibniz University

HANNOVER



Team
TALebots



@igem_hanover



iGEMTeamHannover



info@igem.uni-hannover.de

The second iGEM team from the Leibniz University Hannover consists of 12 highly motivated undergraduate and graduate students from very different fields of study: Biology, Life Sciences, Plant Biotechnology and even Computer Science.

Team

The second iGEM team from the Leibniz University Hannover consists of 12 highly motivated undergraduate and graduate students from very different fields of study: Biology, Life Sciences, Plant Biotechnology and even Computer Science. We all share the common interest in synthetic biology and passion for our project. This competition is the perfect opportunity to expand our scientific horizons and use our theoretic knowledge in the lab. With help from our leaders Prof. Boch and Dr. Reinard as well as the research group of Prof. Boch and several team members of 2014, we hope to transfer our ideas into practice.

In addition, we trust that we will form new friendships with teams from all over the world.

Good luck to all participating teams!



Project

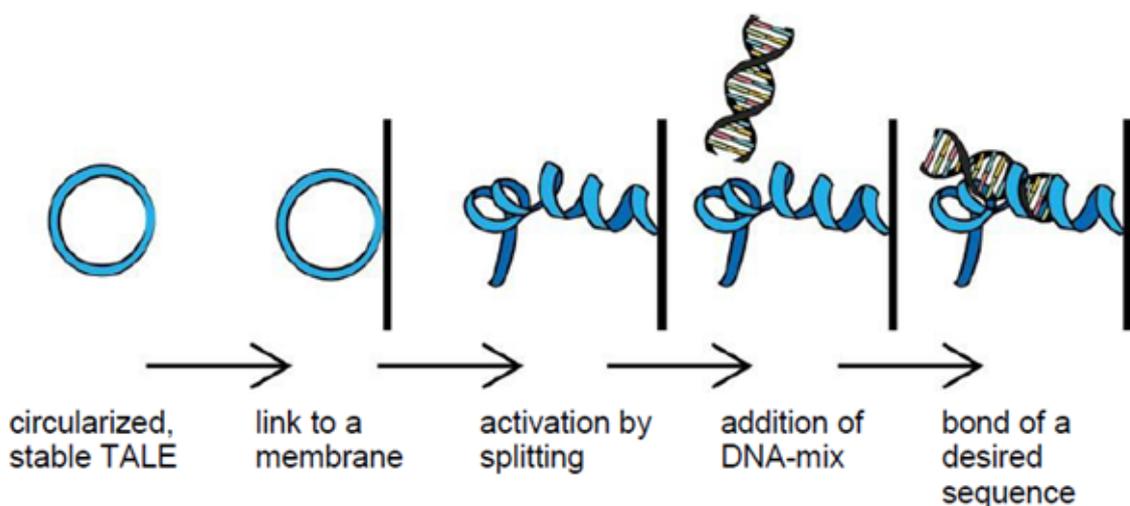
The vision to modify genomic DNA at any specific site has become reality by genome editing.

Scientists can now specifically cut DNA. Two cheap and easy strategies have been proven to be useful approaches: Crispr-Casg and TALEN.

This is exactly where we want to step in. We chose to work with TALE proteins that can be designed in a way that they recognize a changeable, but defined DNA sequence without further molecule classes like unstable RNA.

Typically, TALE proteins can be combined with various effector enzymes (e.g. restriction enzymes) which can carry out the desired operation. Nevertheless, the use of these systems is restricted to in vivo applications, since the enzymes are not stable in vitro. Proteases can attack and destroy the link between amino acids, which form the protein.

To circumvent these problems, we aim to stabilize TALEN via circularization of the protein. This modification should increase the stability of the protein and enables us to use it in vitro. To circularize the protein, we use linkers that bind both ends of the protein together. Combined with an effector domain, we want to establish a new type of recombinant protein that we call "TALeBot". Our



approach will permit many new applications for these proteins.

In the following, we will test our “TALeBot” for stability against heat, acid and storage. Hopefully, we can enable the use of TALE proteins for in vitro modification or aimed cutting of DNA in a cell-free environment.

Furthermore, we plan to immobilize a specially designed “TALeBot” onto a membrane and link a desired DNA sequence. With this technique, we image new applications like the detection of carcinogenic virus DNA from blood plasma or a one-step-purification of plasmids from bacteria solutions.



Collaboration

Do you want to work with us?

We are looking for other teams to collaborate with. Sharing ideas, knowledge, tips or even working together on part of the project would be great. If you are working with TALE proteins, CRISPR/Cas9 or you think that our project might help your research, don't hesitate to contact us or have a look on our website: <http://www.igem-hannover.de/>

We can also be contacted by Facebook (@iGEMTeamHannover), Twitter (@igem_hanover) or by mail (info@igem.uni-hannover.de).

Human Practice

What do we do outside the lab?

The iGEM competition is not limited to lab work only or transferring your dream project into practice. We are also applying our project into a social context.

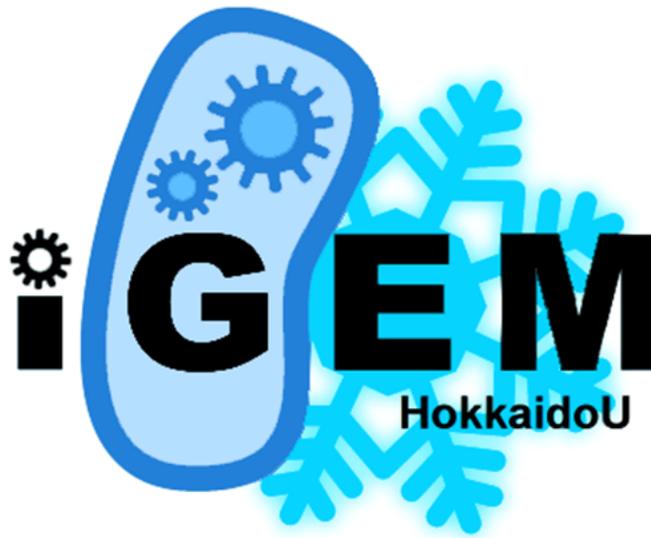
The enormous potential and the risks as well, are still not too much present in the media, and the general public. That is why we want to inform people that there are indeed silver linings. In fact, TALEs were already used to cure a girl from cancer.

Moreover, we want to address younger future scientists by giving talks at their schools and telling them about our fields of study and the iGEM competition.

Since we also want to enhance collaborate with other teams, we took part in the annual iGEM Meetup in Marburg and the European Meetup in Paris. Both weekends were great and we could talk to other iGEM teams about their problems and their ideas.

Hokkaido University

HOKKAIDOU_JAPAN



@igem_hokkaidou



iGEM HokkaidoU



yukinosbt@eis.hokudai.ac.jp

iGEM HokkaidoU is based at Hokkaido University and is the northernmost iGEM team in Japan. We have been participating in the competition since 2010 and it is our seventh time this year.

Team

iGEM HokkaidoU is based at Hokkaido University and is the northernmost iGEM team in Japan. We have been participating in the competition since 2010 and it is our seventh time this year. Hokkaido is vast and enriched with nature, and is especially famous for its numerous World Heritage Sites such as the Shiretoko peninsula. In the winter, it is a haven for snowboarders and skiers from other parts of Japan and also abroad, who gather in pursuit of powdery snow. Hokkaido University is no exception in terms of its blessed environment— its large campus is known as one of the most nature-enriched campuses in Japan. Here, students of various fields of study gather, such as literature, economics, law, engineering,

natural sciences, fisheries, agriculture and medicine to name just a few. iGEM HokkaidoU consists of members from a diverse field of study – most members belong to biology-related faculties such as natural science, agriculture and medicine, but we have had some years when members from faculty of engineering and literature participated and did a great job with modelling the experiments and designing the Wiki. We set no requirements for joining the iGEM team other than to have interest in synthetic biology, therefore we are able to assign specific work to each member in their strongest area to bring the best out of everybody. In the past 6 years of participating in the competition,



we were able to receive Best Wiki award in 2011 and Gold medal in 2012. Since not all team members major in biology, we hold study sessions to learn about basic molecular biology and synthetic biology essential for iGEM. We use the period of from after the competition up until around July to study and plan our project, and we get on with the experiments during the summer vacation in August and

September. Before the experiments kick off, we hold sessions in which one member uses the book 'The molecular Biology of the Cell' or a past iGEM Wiki of a different group to bring up some topics for discussion, which is followed by researching relevant papers and improving the project plan. In tandem with the experiments, we also work on modelling and Human Practice projects.

Project

Right now, at the end of August, we are getting on with the experiments and are about to reach the crucial part. This year, we are focusing on the properties of the self-assembling peptide, an amphiphilic peptide which self-assembles under physiochemical conditions, and are planning to use this for the circularization of proteins and other various purposes.

Circularization of a protein is one of the ways to enhance its stability against various temperatures and pH levels. We inserted cysteine residues in the linker which is essential for the circularization so that they form a disulfide bond. These two elements - the self-assembling peptide and the linker - enable the protein to be circularized.

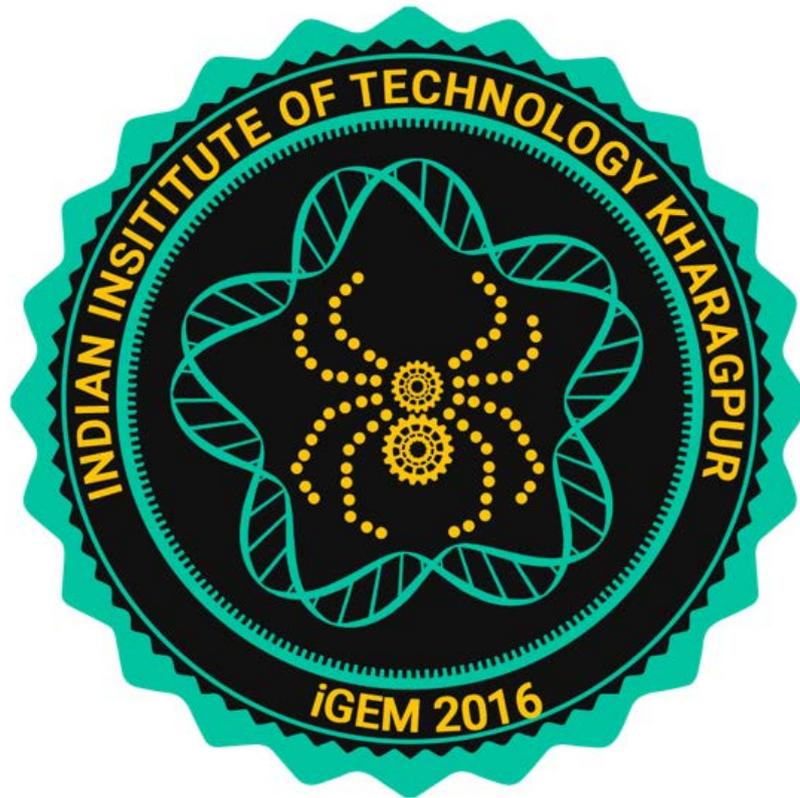
Human Practice

For our Human Practice, we organized a DNA-extracting booth at the school festival designed to introduce synthetic biology and iGEM to the public. Because there is still a negative impression of genetic recombination in Japan, we hope that this offered a good opportunity to spread the correct knowledge of what it is all about and how synthetic biology has a great potential and influence over other fields of study. Last year, other than the school festival, we also tried crowdfunding for the first time. Crowdfunding is a system where people present their project via a website to encourage the public to fund for them. We used a recently created crowdfunding website that specializes in academic projects to fundraise by promoting our previous iGEM project.

Thanks to the advance of such systems, the public and researches all over the world will be able to gain a better understanding about synthetic biology as well as other frontier researches. At the same time, beneficial or intriguing research projects will be able to gain more fund. We hope to continue engaging positively in such stimulating approaches to research.

The Indian Institute of Technology Kharagpur

IIT_KHARAGPUR



@igem_iitkgp



iGEMIITKharagpur



rhushikeshphadke@gmail.com

We are an 17 member team consisting of third and fourth year undergraduate students who look after different tasks of the project including wetlab, public policy, conducting surveys, keeping track of the updates .

Team

We are an 17 member team consisting of third and fourth year undergraduate students who look after different tasks of the project including wetlab, public policy, conducting surveys, keeping track of the updates . Our seniors who were a part of the team last year are our guides. Our team has two young and dynamic PIs as well.

Project

Silkotron: A genetically engineered machine for efficient production and export of spider silk

Our arena for this year's idea hunt was the use of available parts in its registry and past ideas. This was because it would take a bigger budget and higher expertise to create new parts and also the funding organization required us to use the previous parts only. Needless to say, there was only gold and diamonds and more inspiration to find in the old records. So, this year,

our aim is to have a fast and efficient production of spider silk using the synthetic biology model bacteria E.coli.

Spider silk is a very versatile material having its applications in various industries like medical, engineering, clothing, biomedical to name a few. But the spider silk cultivation is a very slow and cumbersome process. Our idea aims at synthesizing recombinant spider silk protein and produce it extracellularly in our model bacteria E.coli, the main motive behind it being



mass production of spider silk through rapid multiplication of bacteria. For the project, we are making two constructs to express silk outside the cell, all parts of which are already available in the parts registry. We also aim at developing a novel detection system using FRET dye pairs. This system will primarily be used in monitoring and modelling the cleavage activity of HIV protease used for selective cleavage and controlled release of the silk protein. CFP and YFP will be used as FRET dye pairs. The idea of cleavage and release of HIV protease followed by cleavage and release of silk

protein described above will be tested by the FRET based assay. The construct containing the MaSp2 protein will contain OmpA, HIV1 cleavage site, MaSp 2E. An HIV protease cleavage site will be introduced between the OmpA fragment and the silk protein assembly. A second construct containing the HIV protease will be similar to the silk construct, the only difference being replacement of the MaSp2 gene with the HIV protease gene. The FRET system using CFP and YFP will be used as FRET dye pairs.



Imperial College London

IMPERIAL_COLLEGE



@imperialigem



2016imperialigem



igem.imperial2016@gmail.com

Our team is composed of 12 members. Half of our team comes from the faculty of natural sciences, the other half come from the department of bioengineering at Imperial.

Team

Our team is composed of 12 members. Half of our team comes from the faculty of natural sciences, the other half come from the department of bioengineering at Imperial. Our chosen disciplines span from biochemistry to biomedical engineering. Here are some need-to-know facts about Imperial's 2016 team:

Biggest Challenge:

Not boiling cell cultures in the incubator (don't ask, it's been done before)

Team Communication Strategy:

Open lines of communication are important. We believe gifs, vines, and interpretive dances are better than words and talking.

Days since last seance: 10

Members who have to leave by 5 to get to their coven meetings: 4

Ideal Collaboration: A team which also enjoys long walks on the beach and candlelit dinners.

Things We Understand About iGEM:

Synthetic biology is exciting, GFP will verify anything, and building a wiki is insane

Supermodel Alter-Egos Present On The Team: Karlie Kloss, Kendall Jenner, Bar Refaeli, Gisele Bundchen, Naomi Campbell, Cara Delevigne, Adriana Lima, Priyanka Chopra, Gigi Hadid, Alexa Chung, Tiffany Trump, and Coco Rocha

Project

In nature, microorganisms live together and cooperate to accomplish complex tasks. As synthetic biology advances, the field will transition from engineering unicellular systems to multicellular systems. Products from complex genetic circuits that were previously too burdensome for a single cell will be split between



specialized populations.

It is difficult in co-cultures to ensure the stable coexistence of different cell types over time. Current co-culture technology suffers from its inability to accomplish this, making it unpopular in the scientific community.

This year we are developing a system called Genetically Engineered Artificial Ratio (GEAR) to control population ratios in microbial consortia. GEAR is composed of three distinct modules: communication, comparison, and growth regulation. The communication module will feature a bi-directional communication system composed of orthogonal quorum sensing systems. The presence of quorum molecules in the cytoplasm will activate the transcription of the RNA molecules. If the correct population ratio has been achieved, equal amounts of sense and antisense RNA will be produced, which sequester each other and prevent activation of the growth regulatory module. The growth regulatory module produces growth repressing proteins that aim to slow division of the cell, to achieve the desired ratio. In the future, we envision our GEAR system being used for distributed multicellular biocomputing, bioprocessing, and microbiome engineering.

In conjunction with our circuit, we are developing a software to facilitate the design and optimization of co-cultures. Our software will allow researchers and future iGEM teams to conveniently access co-culture protocols for the best practices in their own experiments. The software will provide information on appropriate pH, temperature, and inoculation ratios among other data. We will be verifying the information in our software through our own experiments, models, and collaborations with other iGEM teams who are utilizing co-cultures for their own projects.

Collaboration

Please contact our team at igem.imperial2016@gmail.com if you would like to collaborate. We are specifically looking for teams who are growing different cells in co-culture. However, we are open to other suggestions for collaboration.



Jilin University

JILIN_CHINA



@IGEM Jilin_China



jilin_china@outlook.com



iGEMxHU



+86 15754305638

Jilin_China team is a team that has fourteen undergraduates from four different majors and six teachers as PI, secondary PI, instructors and advisors.

Team

Jilin_China team is a team that has fourteen undergraduates from four different majors and six teachers as PI, secondary PI, instructors and advisors. Especially, professor Yongge Wu and Dr. Xin Hu provide us with excellent supports of lab equipment and academic consults. All members are from the School of Life Science at Jilin University.

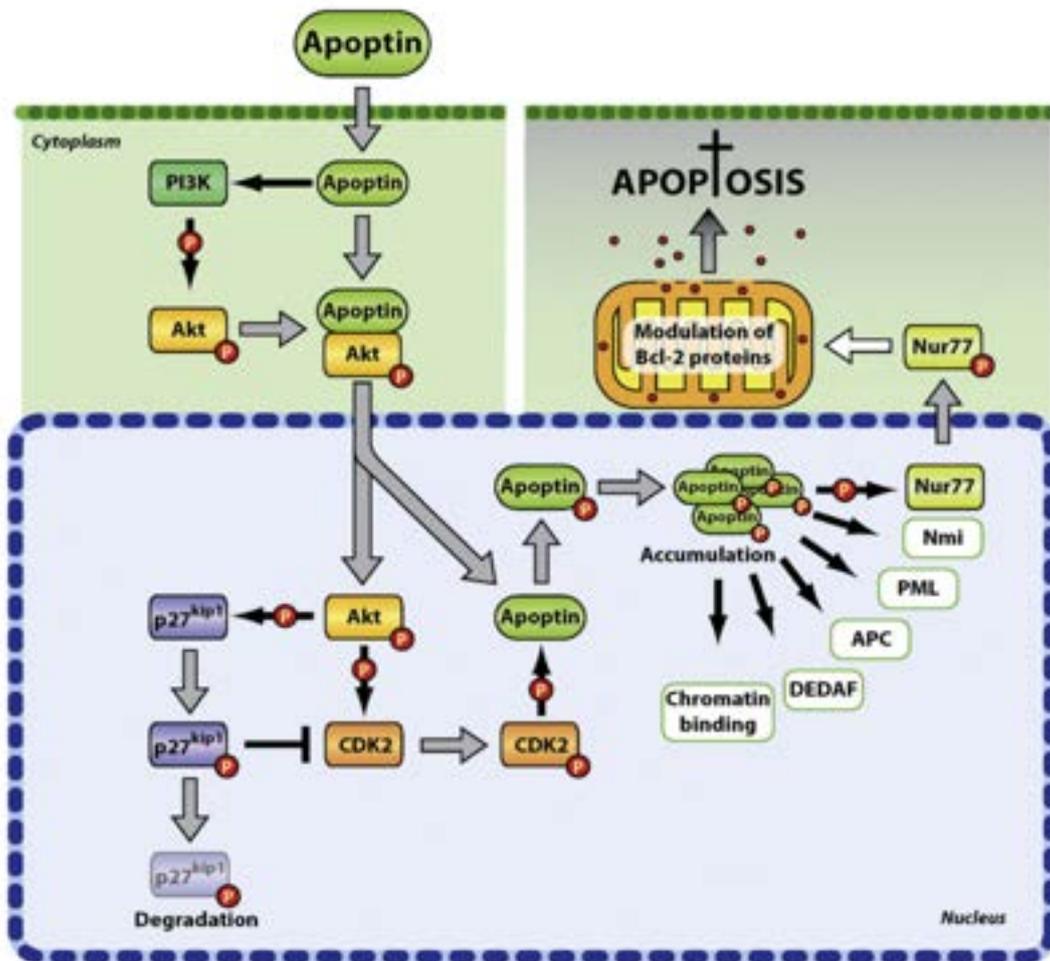
Project

Solid tumor cells can be considered as harmful mutants in the ecosystem in human body. We therefore propose a novel approach to eliminate tumor cells. Bifidobacterium has been proved to be non-immunogenic to human body, and has the ability to settle in hypoxic regions, such as solid tumors, preferentially. First of all, we plan to construct a recombinant plasmid that consists of DNA replication origins and related genes from pMB-1 and pUC18. In addition, we will clone HU (a Bifidobacterium promoter) and TAT-apoptin gene with a N-terminal secretion signal peptide into this recombinant plasmid. Apoptin is a small protein that can induce tumor-specific cell apoptosis independent of p53.

Thus it specifically kills tumor cells while leaving normal cells largely unaffected. TAT is a protein with a transduction domain named PTD that has the ability to help apoptin to transduce through cell membrane. Additionally, we plan to use two other signal peptides, Tmp-1 and Sec-2, to ensure the secretion of apoptin.

This plasmid can be amplified both in *E. coli*. and Bifidobacterium, and apoptin can be expressed specifically in Bifidobacterium. With the help of signal peptide and TAT, apoptin can be secreted by Bifidobacterium and be transduced into solid tumor cells successfully to achieve the ability of killing tumor cells.

Moreover, we will inject Bifidobacterium into tumor-bearing mice, and simulate the distribution and association of Bifidobacterium with tumor cells using ecological competition models. After the tumor-killing ability of this recombinant plasmid is proven, we will focus on enhancing the biosafety of our system through modifying the HU promoter activity to regulate the apoptin expression and applying a conditional regulatory mechanism to restrict the proliferation of Bifidobacterium.



Collaboration

To simulate the interaction between normal cells, tumor cells and our engineering Bifidobacterium, we have adapted a model based on classic predator-prey model. We hope that this mathematical model can fit in the vitro experimental data well. Therefore we need an ingenious design of vitro experiment that can mimic the environment in vivo.

Since there are both normal and hypoxic areas in the body of tumor-bearing mice, this vitro experiment must have both normal area and hypoxic area. In this situation, we will know whether our Bifidobacterium can migrate from normal area to hypoxic area or not. In addition, this hypoxic area needs to be similar to the environment in solid tumor, which will help us to evaluate the ability of our Bifidobacterium to kill tumor cells in vivo. If you have any suggestions or comments about the design of our experiments, please do not hesitate to contact us.

Human Practice



On April 4th we were invited by our local television through which the influence of our project is greatly expanded. On May 29th, we attended Changchun International forum of Oncobiology and Translational Medicine. We asked experts about new information and methods of treating solid tumors and their advice on our project. They provide us with lots of useful advice to improve our project. At the same time, we set up

our own WeChat Public Number and Twitter account to update our news and receive feedback from other teams and those who are interested in our project. The feedback will be used to improve our project. We were invited by Huazhong University of Science and Technology and attended HUST-Cheering meeting in Wuhan. We introduced our project to iGEMers from all over the country. Professor Wenliang Chen provided us with some insightful advice on our model and cell experiment. We also learnt lots of very valuable information to improve our project better.

Leiden University

LEIDEN



@iGEM_Leiden



igemleiden



igem@science.leidenuniv.nl

For the first year, Leiden University from the Netherlands participates with a team of its own in the iGEM competition – with a killer application for synthetic biology!

Team

For the first year, Leiden University from the Netherlands participates with a team of its own in the iGEM competition – with a killer application for synthetic biology! Our team consists of 13 highly enthusiastic and ambitious students with a broad variation of backgrounds: biology, life science & technology, statistics, physics, mathematics, astronomy and science based business, with bachelor as well as master students. If you have any ideas on how to improve our project, need help on any subject or just want to say hi – do not hesitate to contact us via igem@science.leiduniv.nl, Facebook or Twitter! We hope to get into contact with as many teams as possible!





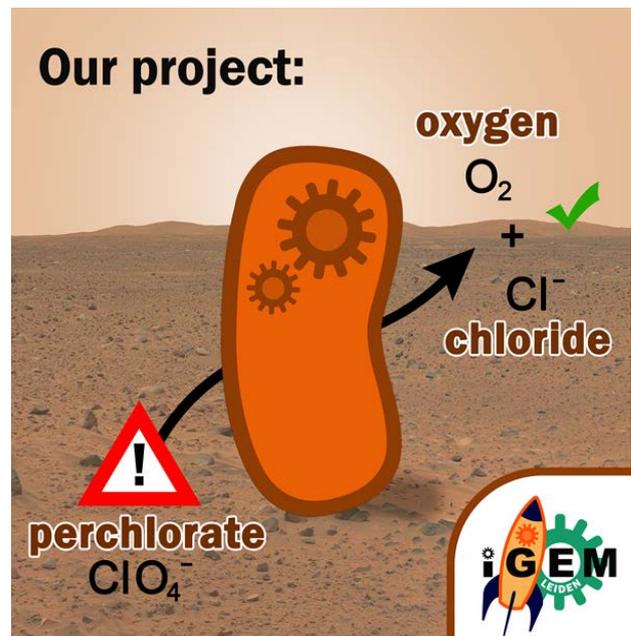
Project

E. colonizer: gardening on Mars! Have you ever thought of building a garden on Mars? Perchlorate (ClO_4^-) contamination of groundwater, surface water and food supplies is a widely spread hazard for the environment and our health on earth. However, it poses even a larger challenge when colonizing our neighbour planet Mars. Martian soil contains 0.5 - 1% perchlorate, which therefore needs to be remediated in order to cultivate edible crops.

The Leiden iGEM team will equip the bacterium *Escherichia coli* with the tools to convert the toxic perchlorate into chloride and oxygen (!) by introducing a set of codon-optimized genes from *Dechloromonas aromatica*, encoding for the perchlorate reductase complex and chlorite dismutase. In this way, the system can be much better understood, used in faster growing bacteria than the original ones and therefore optimized for use on a larger scale.

Besides, we will study *E. coli*'s gene expression under Martian gravity (0.38g) using a Random Positioning Machine, to make sure that our system will function in a bioreactor on Mars and find gravity-dependent genetic elements. Altogether, our system is widely applicable to remove perchlorate from contaminated soils on earth, while also being highly useful for future Mars expeditions. So collect your seeds, rake, shovel and watering can and join our mission to Mars!

More information on launch dates, selection procedures for Martian Gardeners and specifications of our



Mission can be found at our wiki: 2016.igem.org/Team:Leiden!

National Chiao Tung University

NCTU_FORMOSA



NCTU_Formosa-IGEM-team



nctu5168victory@gmail.com

Since the problem of chemical pesticide has long been existing, We, NCTU_Formosa, living in a high-tech era, created a novel safe and organic bioinsecticide. We utilized IoT talk coupled with automatic spraying system to achieve the goal of monitoring and operating remotely and solved the problem of chemical pesticide.

Project



Since the problem of chemical pesticide has long been existing, We, NCTU_Formosa, living in a high-tech era, created a novel safe and organic bioinsecticide. We utilized IoT talk coupled with automatic spraying system to achieve the goal of monitoring and operating remotely and solved the problem of chemical pesticide. A brand-new safe bioinsecticide "PANTIDE" which is a combination of two words, "PAN" and "Peptide". We hope that PANTIDE turns the environment into the primordial stage as if it were the god of shepherds, Pan, in ancient Greek mythology. The peptide is the essential substance of Pantide. Besides, sympathizing the laborious effort of farmers, we availed the technique of IoT and created a personalized automatic system that assists farmers to do the farm work remotely.

There are more than ten detectors in our device including tracking the image, the number of pests, temperature and so on. The data will be uploaded to the user's app by IoT technology, and it will set up the database in the cloud at the same time. So users can easily monitor the condition of the farm then decide whether to spray Pantide. We use IoT Talk to make a bridge between the device and the application on the smartphone. Moreover, the device is able to decide whether to spray Pantide or not according to the received data, simultaneously informing of the users when spraying.

Perhaps we do not have sufficient background knowledge as experts may possess, but we will do our best to make Taiwan and even the world a better place, and this is the blueprint of our dream. We are NCTU_Formosa.

Collaboration

To help the teams achieve their project, we give the iGEM distribution kits to Mingdao, NCKU_Tainan, and NTHU_Taiwan iGEM teams. Besides, our team and Mingdao also exchanged both our biobricks to test each other. Furthermore, we helped our partner, Aachen iGEM, to promote their survey in Taiwan area. And now we are focusing on one special idea for a Human and practice collaboration with Aachen, Israel, and Mexico iGEM teams. And we also helped Evry and Hannover to promote their surveys, too.

And Thanks for NYMU iGEM team provided us the Lepidoptera eggs. For our insect test, we needed lots of larvae to do experience. Additionally, they introduce the Pro. Hwang, who majors in agricultural science, and gave us many advises in our insect test. All in all, we thank for them lots.

Human Practice

This year NCTU_Formosa has designed an exciting board game, Pest Crisis. Pest Crisis is about scientists versus pests, fighting for crops. The player will separate into two groups, scientists, and pests. For scientists, players have to design biobricks to produce pesticides. Besides, pests have to evolve some traits to resist pesticides. The reason why we decide to create a board game is that want to make it easy to learn the knowledge of synthetic biology, and to make players know the importance of the pest problem and chemical pesticide issue.

We played our board game in many Conferences and meeting ups with other iGEMers. It's a great tool to let other iGEMers preliminarily understand our project and use it to promote synthetic biology. Everyone was so excited for our board game,

and we soon became good friends. We have designed both Chinese and English version of our board game and will reveal on the giant jamboree. Hope you will like it!



To make sure the opinions about our Project from the potential users in the future. We went to Organic Green Market in NTHU to have an interview. There are many organic farmers in this market and we visited them one by one. According to our product demonstration, except introducing insecticidal peptide PANTIDE only, we also ask them some questions about the device including a detector and automatic sprinkler to get farmer's recognition and recommendation. Most farmers really appreciate our project because, in the recent organic agriculture system, the pests of cruciferous vegetables still rage the farm. As a result, they are all willing to use our product. Additionally, the farmers are willing to let us go to their farm to grab pests for the intra-laboratory experiment. Moreover, they are even willing to provide their farm to let us have a real test for the field test.



To realize the professional opinion of pesticide residues, we went to the famous organic tea store – 飲川 consulted with the owner about organic knowledge including the effect of the pesticides residues on the tea leaves. He said though the cost of organic tea is high, it still has its market value. He also reminded us that if the upstream farm using pesticides, the downstream farm will be polluted by the pesticides residues too. So we think our project will solve this problem due to PANTIDE is biodegradable. All in all,



Besides, we consulted Dr. Huang and his postgraduate student in Taiwan Agricultural Research Institute Council



of Agriculture. He thought our device has its existent value because Agribusiness and related administrative departs need it to monitor farms. The device helps them detect the pests and analysis the related data such as temperature, soil moisture, and rainfall in real time and so on. Dr. Huang also told us the method of pest detection and interrelated knowledge. Other than conferring knowledge, Doctor Huang also gave us some remarks for our project, which makes us realize different aspects of agriculture in manufacturing and research. Even more, he let us visit the cultivated room of Oriental fruit flies.

In the process of feeding tobacco cutworm larvae, we face the problem of no larva dilemma because tobacco cutworm larvae we kept are all having diarrhea and finally dead. Later, the fodder we make also have several mistakes. At this time, to find out the reason and get the solution, we write letters to Professor 唐立正 at Department of Entomology at National Chung Hsing University to query insects feeding recipes and check with him that which step is wrong then improve it. We also interview Professor 黃紹毅 at Department of Entomology at National Chung Hsing University. We not only share our project with him but also inquire about the feeding method of tobacco cutworm. The professor is so enthusiastic about providing us tobacco cutworm larvae resource and give us a lot of advice related to raising pests.

To promote the concept of Synthetic Biology, we went to different senior high students' summer camp to popularize iGEM competition. We had a brief presentation to make them understand some knowledge about synthetic biology, such as EXSP digestion position, biology central dogma and so on. Also, we had Q&A time to clarify their confuse. In the presentation, we present this year's project to show how we use synthetic biology technology to solve nowadays problem, too.



This summer, fourteen high school students, who are interested in biology from National Taichung First Senior High School came to National Chiao Tung University learning about iGEM. In the two-weeks workshop, they have learned a lot of basic concepts of synthetic biology associated with laboratory practices.

On the last day, each group designed their own projects, which contained brainstorming, principles, and experimental designs. Their projects could be fully developed further. Through the series training, we think that they are the iGEMer-to-be and expect their projects in Giant Jamboree next year.



We were honored to have the opportunity to invite the captain of National Chung Cheng University. We exchanged a lot of experience of the operation of the team and academic advice. Besides, she also joined the two-week workshop with students from National Taichung First Senior High School. Hope this will help them in training new team members. We are looking forward to shining on the iGEM platform, too.



NCTU_Formosa and iGEM Aachen had a Skype video chat. In the online chat, we grasped the project bilaterally. iGEM Aachen designed an artificial enzyme with an enzyme-inhibitor that replaces boric acid in washing detergent.



Apart from exchanging academic opinion, we had a great chat in the history of team, agriculture, habits and also the mind-blowing table game “Pest Crisis.” RWTH Aachen University and National Chiao Tung University are sisters schools. Therefore, Aachen iGEM is one of a strong companion in Europe. Our partner also helps us to promote board game and survey in Germany area. There are already some people



very interested in playing our board game right now.

Discussion

For the safety part, we sonicate E. coli after IPTG induction and adds antibiotics into the solution then we heat it at 60 degrees for 2 hours so that we kill the E. coli but maintain the function as PANTIDE. It is crucial to build a safe and sustainable project and raise awareness among public.

University of Oxford

OXFORD



@oxfordigem



oxfordigem



oxfordigem@bioch.ox.ac.uk

We are interested in developing a novel probiotic therapeutic containing two copper-chelators, controlled by a copper-sensitive mechanism, with the aim of treating Wilson's Disease.

Project

Overview

We are interested in developing a novel probiotic therapeutic containing two copper-chelators, controlled by a copper-sensitive mechanism, with the aim of treating Wilson's Disease. Wilson's disease is a genetic disorder in which the body is unable to fully metabolise copper, resulting in copper accumulation and consequently liver and brain damage.

Our system involves a 6-step mechanism:

- i) Delivery of the bacteria into the small intestine via an encapsulation system
- ii) Establish a stable population of engineered bacteria
- iii) Transcription and translation of our chelators
- iv) Absorption of copper ions in the small intestine into the bacterial cells
- v) Sequestering of copper ions via copper chelators
- vi) Removal of bacterial population via egestion

Chelators

During our literary research we decided that the ideal copper chelator should have these properties:

- Be able to bind multiple copper ions per peptide to increase efficiency
- Be from the prokaryotic domain because eukaryotic proteins have expression issues in E.coli
- Be localised to the periplasm of the E.coli cell as E.coli naturally deals with copper toxicity by binding copper in the periplasm



Using these criteria, we found two copper chelators: Csp1 and MymT.

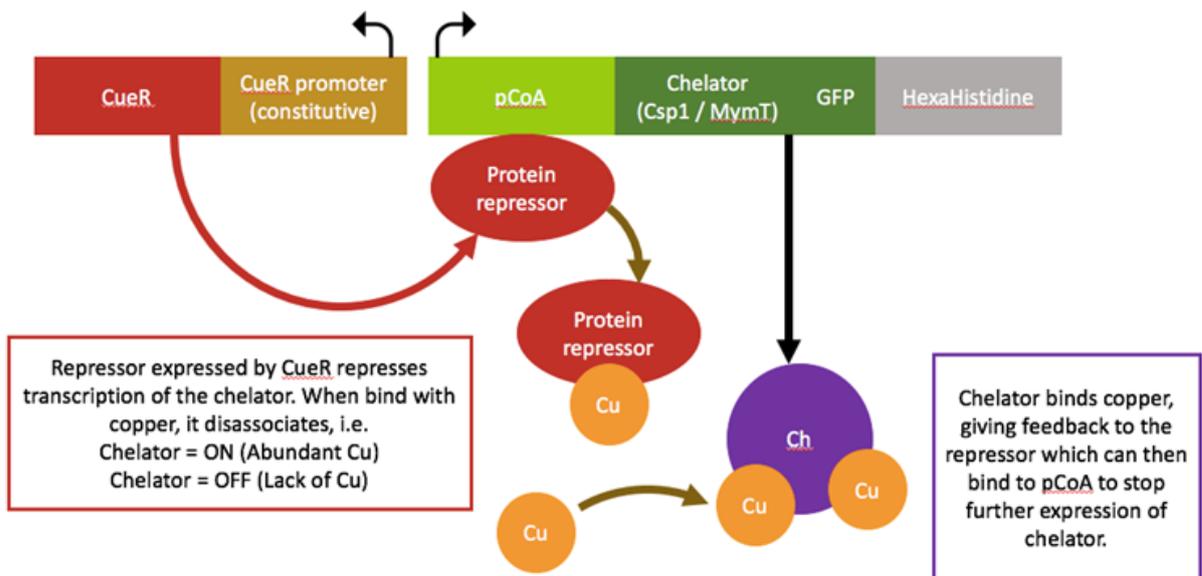
Csp1 is naturally found in a methane-oxidizing alphaproteobacterium called *Methylosinus trichosporium* OB3b. Each monomer within the tetramer structure can bind up to 13 Cu⁺ ions, giving a maximum binding of 52 Cu⁺ ions. MymT is a small prokaryotic metallothein discovered in *M.tuberculosis*. It can bind up to 7 copper ions in a solvent shielded core but studies show it prefers 4-6.

Feedback system

Our feedback system ensures the chelator is only produced at appropriate copper concentrations, to prevent overproduction of the chelator and hence the possibility of copper deficiency.

Potential applications

Although our project centres around the treatment of Wilson's Disease, the molecular mechanisms behind our system could potentially be transferred to a variety of other conditions. For example, hereditary hemochromatosis causes the body to absorb too much iron from the diet, leading to liver disease, heart problems and diabetes. Our model could be adjusted, altering the chelator and promoter to iron-sensitive versions, to provide a treatment for this condition.



Peking University

PEKING



|  pkuigem2016@126.com

The team of Peking iGEM 2016 consists of 15 students of various majors including biology, medicine and physics.

Team

The team of Peking iGEM 2016 consists of 15 students of various majors including biology, medicine and physics. Although the majority of the members are sophomores, they are creative and willing to learn. Over the past six months, Peking iGEM 2016 has become a professional and passionate team with the help of the leaders and instructors.

Project

Uranium, a heavy metal element, is weakly radioactive and poses a threat to both the environment and human health. A person can be exposed to uranium by inhaling dust in the air or by ingesting contaminated water and food. Long-term exposure to uranium increases the risk of various diseases and health issues including cancer, kidney problems and immune system damage. Uranium has become more commonplace due to nuclear accidents (the Chernobyl Accident, the Fukushima Daiichi nuclear power plant Explosion), uranium mining and the development of depleted uranium weapons.

To alleviate these problems, the Peking iGEM team aims to construct a novel functional biological material, which can absorb uranyl ion with the employment of a specific uranium-binding protein.

This novel material has numerous promising characteristics such as high specificity, high efficiency, self-assembly and self-reproduction. With some modification, the design can be applied to deal with uranyl ion in polluted water and soil, demonstrating its impressive potential. We believe that the material can effectively solve the increasingly serious uranium pollution in the near future.

Collaboration

1. Meet up

In August, we invited three other iGEM teams to attend the meetup in the College of Life Sciences of Peking University, where we met and exchanged ideas with members from Tianjin, BIT-China and UCAS.

We introduced our projects and built up mutual facilitation among teams. As one of the first teams of China to attend iGEM, Peking take the responsibility for guiding and helping other first-time contestants, UCAS for example.

Additionally, we suggested the members of Tianjin to use Golden Gate Assembly methods by considering their problem of low efficiency of assembling.

2. Visit

In the end of July, we visited the iGEM team BIT(Beijing Institute of Technology)-China. We shared our project design and gained some feedback which might promote us. Zhang Yihao, our instructor, shared with them his iGEM experience and insight to synthetic biology.

The process of BIT-China was limited because the result of RFP expression and asked us for help. By consulting literature and doing experiment, we suggested them to wait for a period to make the color displaying complete or have the protein exposed under UV light.

Human Practice

We assisted several iGEM teams of China by offering them plasmids and other experimental materials.

For instance, our team assisted Jilin-China with Interlab Measurement Study by sending them the plasmid J23117+l13504.

In addition, we mentored a high-school team: BHU(Beijing High School Union) from July to October. We supported them not only the guidance of fundamental molecular laboratory operating but the experimental materials such as expression vector pET28a and corresponding primers, as well as all enzymes and laboratory equipment.

Discussion

Our team aims to construct a biological material to absorb uranyl ion. As a result, uranyl ion is an indispensable part of our experiments, which may pose potential threats to lab safety. To alleviate the problem, all the experiments about uranium are carried out in a qualified Radiation Laboratory in Peking University. To assure the health of our members, we measured radioactivity in different positions of the laboratory. The data obtained proved that the radioactivity in every corner of the working area was close to the background value measured in grassy playground.

Consequently, we can affirm that our members are doing experiments in a safe environment.

Furthermore, in the laboratory with potential radioactivity, all the students would be well-equipped with special protective garments, lead protective aprons, respirators, gloves and caps.

University of Peshawar

PESHAWAR



iGEM Peshawar 2016



sidra.usman.biotech@gmail.com

We are the first iGEM team from Pakistan!

Project

Air pollution is a major global problem, a significant amount of which is caused by vehicular emissions. This problem is particularly prevalent in developing nations, where regulations regarding vehicle emissions are either not present or not properly implemented. Carbon monoxide and oxides of nitrogen are two dangerous constituents of exhaust fumes. Our aim is to produce a portable, quick and easy to use Biosensor device for environmental law enforcement agencies and even consumers that can detect for levels of carbon monoxide and oxides of nitrogen in vehicle exhausts and express corresponding chromoproteins as a result. The system is based on two separate gas-sensing mechanisms that work together to give a range of results. The mechanism sensing for carbon monoxide uses the CooA transcription factor and two CooA dependent promoters: CooF and CooM. The NO_x sensing mechanism uses the NsrR repressed promoters, YeaR and NirK, as well as the NorV promoter.

Life Sciences at EPF Lausanne

PRETORIA_UP



@iGEMPretoria_UP



iGEM Pretoria_UP



PretoriaUPiGEM@gmail.com

The Pretoria_UP IGEM team was first established in 2015 as part of the Forest Molecular Genetics (FMG) research group of the University of Pretoria.

Team

The Pretoria_UP IGEM team was first established in 2015 as part of the Forest Molecular Genetics (FMG) research group of the University of Pretoria. The team has since grown to have 11 students in different study disciplines under the supervision of an instructor and advisors who are experts in their respective fields in 2016.

Dr Steven Hussey
Instructor



Prof Zander Myburg
Advisor



Dr Tjaart Kruger
Advisor



Dr Eshchar Mizrachi
Advisor



Dr Musa Mhlanga
Advisor



Prof Ncholu Manyala
Advisor



Dr Michal Gwizdala
Advisor



Dr Marco Weinberg
Advisor



Pieter
BIS Multimedia



Bernard
BSc Microbiology



Brad
BSc (Hons)
Medicinal Plant
Science



Modjadji
BSc Microbiology



Thabang
BSc (Hons)
Biotechnology



Simon
BEng Electronic
Engineering



Stephan
BEng Mechanical
Engineering



Maryn
BEng Industrial
Engineering



Vaughn
BSc (Hons)
Genetics



Ricu
BSc Genetics



Nomakula
BSc Microbiology
and Genetics



Project

Our project Watts-Aptamer aims to Design and construct an optimized photo-bioelectrochemical cell using an RNA aptamer synthetic biology strategy.

The world population consumes approximately 3500 kWh/y/capita, increasing the demand for clean alternative energy. Recent improvements of photo-bioelectrochemical cells (PBEC), which harness electrons from photosynthesis to generate electricity, include synthetic attachment of chloroplast thylakoids to graphene electrodes. However, current attachment techniques require costly chemically synthesized linkers and PBECs are not yet efficient enough for industrial energy generation. In this project, DNA aptamers were designed and evaluated as low-cost biological linkers to tether plant photosystem II (PSII) complexes to graphene foam electrodes. Systematic Evolution of Ligands by EXponential enrichment (SELEX), together with software developed by team Heidelberg 2015 (MAWS and JAWS) were used to develop PSII- and graphene-binding DNA aptamer candidates. This project aims to improve the attachment and orientation of the PSII complex to the graphene electrode for higher electron transfer efficiency, and serves as a prototype for the in planta expression of RNA aptamers for self-assembling thylakoid attachment.

We are also looking into improving a part in the registry that has been submitted by another team. A project description video will also be available on YouTube soon.

Collaboration

We have been in contact with the Aix-Marseille and Egypt teams and hope to possibly collaborate with them and contribute to their respective projects in a valuable way. We are also looking for other teams we can collaborate with for our project. the teams can either contribute towards our project as a whole or the human practices part of the project.

Human Practice

Last year we visited two high schools, one in a developed area and one in a previously disadvantaged area. The aim was to introduce the field of synthetic biology and to present our 2015 project to the learners in the two schools. This was followed by the learners filling out a short questionnaire (see more under surveys in this document). From the results we realised that as synthetic biology is a relatively new field, especially in South Africa we need to make people aware of it. This applies not only to academics and university students but also to the public in general.

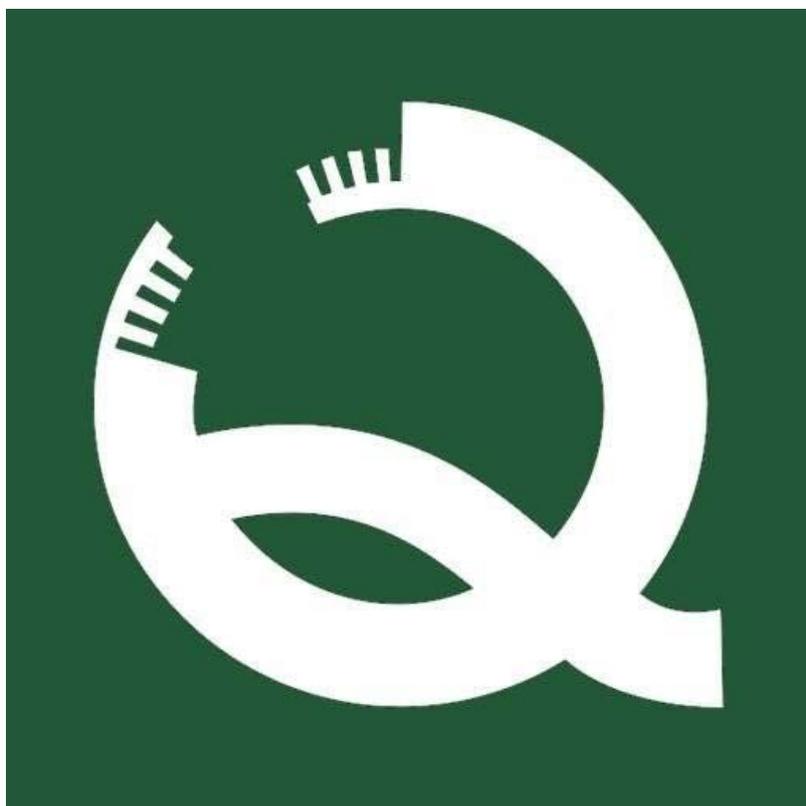
For our human practices this year we want to focus on awareness and education. We aim to make people in south Africa aware of synthetic biology and its many applications. We plan to do this through social media (Facebook, Twitter, YouTube). We will also be presenting to a group of secondary school learners as well as at a synthetic biology symposium which the team will be involved in planning. We will conduct radio, newspaper and hopefully TV interviews where we will also be introducing the field of synthetic biology and making the public aware of its applications. To gain more knowledge that we can apply to our project we will survey and interview experts in the photo-bioelectrochemical cells and Energy field in general. We have already interviewed Dr Karen Surridge-Talbot, centre manager of the Renewable Energy Centre of Research and Development (RECORD) at the South African National Energy Development Institute (SANEDI). Her input has proven to be very valuable and we hope to talk to other representatives from stakeholders in the energy sector such as the National Energy Regulator and Eskom. All human practices events will be documented on our wiki with links to the videos as well as extra information if necessary.

Discussion

We hope to get conversations going and provide a platform where synthetic biology and its applications will be discussed. Such discussions will be around how synthetic biology can be applied to other fields such as energy, engineering, environment, health, medicine, food and nutrition in South Africa. People can also use these discussions to voice out their opinions and concerns regarding this. Experts in the field will also have a chance to share their opinions and answer any questions that people might have. We hope this will bridge the communication gap between the scientific community and the public in addition to the education and awareness parts. These discussions will form part of our human practices.

Queen's University

QUEENS_CANADA



@igemQueens



QGEM



director@qgemteam.com

We will be constructing a pipeline that will allow us to modify NRPS products which are difficult to synthesize.

Team

Executive Members



Dragos Chiriac
Co-Director
Wet Lab Lead



Jia Tanwani
Co-Director
Policy & Practices Lead



Danai Topouza
Computational
Modelling Lead



Yifei Wang
Technology &
Administrative Lead

Wet Lab Volunteers



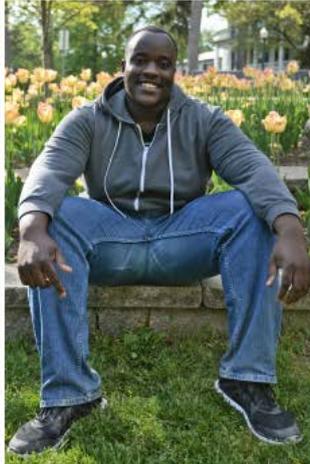
Matthew Bentley



Simran Sharrma



Linor Berezin



Kennedy Ayoo



Policy & Practice Volunteers



Hillary Chan



Danielle Ciren



Julia Grein



Rajiv Tanwani

Computation Modelling Volunteers



Anna Ilina



Sabrina Quazi



Vinith Suriyakumar



Project

The current drug market is dominated by drugs of biosynthetic origin. Greater than 70% of these essential compounds are commonly found in the bacterial kingdom and a significant portion are produced by nonribosomal peptide synthetases (NRPS). NRPS are large multimodular enzymes that create a variety of structurally and functionally diverse peptides. NRPS have catalytic domains to carry out these functions such as condensation domains (C) to couple amino acids together, adenylation domains (A) to activate amino acids, peptide carrier proteins (PCP) which use a thiol arm to swing the substrates between active sites, and the last domain is a thioesterase domain (TE) which facilitates product release.

These large synthetases are composed of a string of modules where each module is responsible for the addition of one amino acid to a growing peptide chain. These amazing cellular machines produce only one major product through efficient enzymatic steps. The shortcoming is that often times the products are cyclic and thus hard to reproduce through total chemical synthesis. Therefore, it is difficult to optimize potential drug leads.

We will be constructing a pipeline that will allow us to modify NRPS products which are difficult to synthesize. The pipeline will consist of 3 phases.

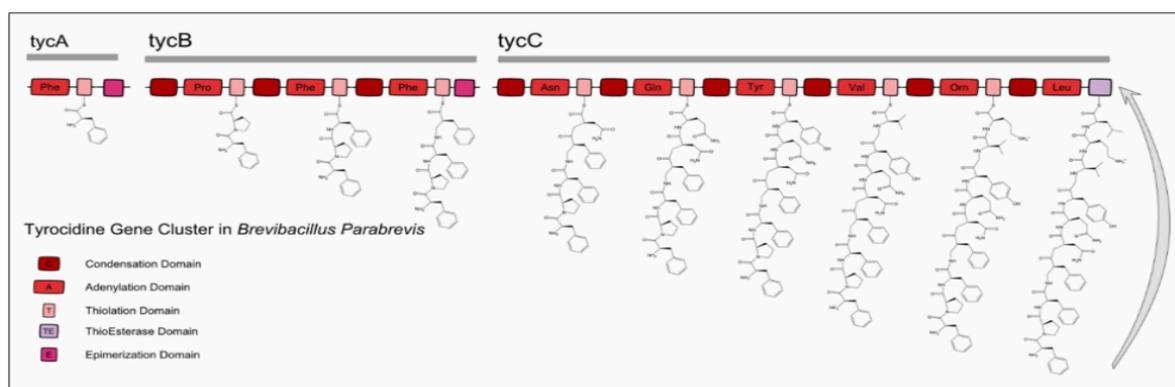


Figure 1. Tyrocidine gene cluster containing C, A, T, TE catalytic domains. Epimerization domains change L amino acids into D amino acids. The final product is a head to tail cyclized peptide which has anti-bacterial properties. Unfortunately, it is also hemolytic.



1. Creating a robust tagging system to visualize the production of NRPS compounds.
2. Modifying domains through genetic and protein engineering in order to change their substratespecificity and append additional modifications onto the substrate.
3. Develop a homologous recombination system which takes advantage of conserved sequences in NRPS domains to construct novel pathways.

Phase 1: Tagging

Indigoidine is a NRPS product that is produced by a single module and produces a brilliant blue colour when it dimerizes. In 2013, Heidelberg's iGEM team demonstrated that it was possible to use the IndC module which produces indigoidine and couple it to a valine module to form a dipeptide which

dimerizes to produce a blue colour. Strangely, no paper was published on the ability to tag NRPS products leading one to be skeptical of their work. The most likely explanation was that they had failed to engineer the robustness of the IndC module and were unable to tag larger preceding peptides which were sterically bulky. Indigoidine dimerizes through a nonenzymatic reaction to produce a blue pigment. A cyclized glutamine that is oxidized triggers the dimerization event. Evidence for this dimerization mechanism can be found if one looks at another NRPS product known as Padanamide B. Padanamide B contains a cyclized glutamine as its final amino acid but lacks an oxidation domain hence the cyclized glutamine cannot be oxidized. Without the oxidation of the cyclized glutamine, no dimerization event occurs in padanamide B while indigoidine is able to dimerize.

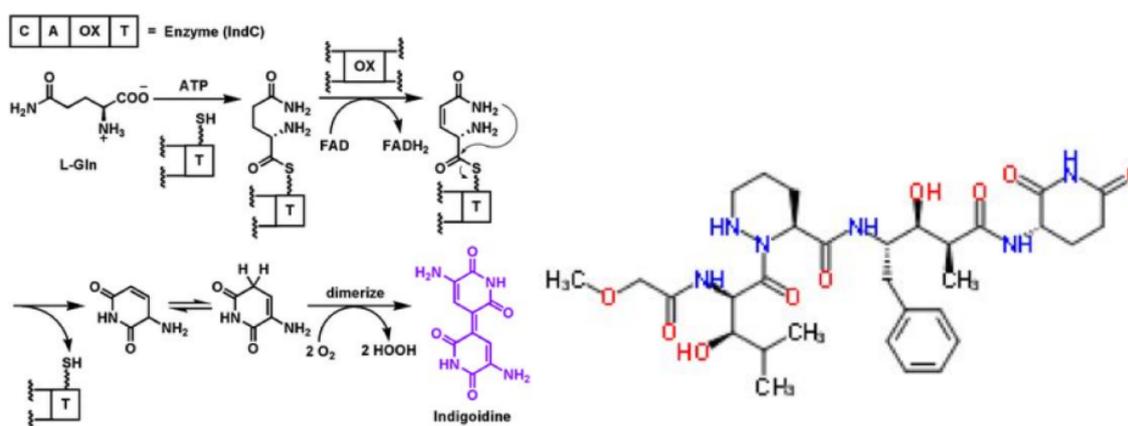


Figure 2. (Left) one of the proposed mechanisms for indigoidine synthesis. Oxidation occurs first then cyclization, opposite of what we suggested. (Right) Padanamide B structure, the final amino acid is the unoxidized version of cyclized glutamine.



This led us to another interesting observation. The mechanism of glutamine cyclization in IndC was proposed to occur through oxidation of glutamine which forced its cyclization, but padanamide lacks an oxidation domain. Therefore, one can conclude that it is the TE domain that is responsible for its cyclization (Figure 2).

We aim to demonstrate that the oxidation domain oxidizes the cyclized product rather than glutamine to induce dimerization. Given that this is true, one would then be able to modify the specificity of the oxidation domain through protein and genetic engineering to visually label any preceding peptide sequence. The visualization of NRPS products will allow us to carry out high throughput screening of small peptide production which is essential for Phase 2.

Phase 2: Module Modification

While previous research shows that one can swap out whole modules in NRPS gene clusters its extent is limited since production of the biologically active product is greatly diminished. Instead research suggests that adenylation domain substrate specificity is not as robust as once believed. Stachelhaus et al. were able to demonstrate that there are ten residues on the adenylation domain which dictate substrate

specificity. Taking both preceding statements into consideration it has been shown that through genome engineering it is possible to change the specificity of adenylation domains to incorporate different compounds. This bypasses the pitfall of low compound yield due to whole module swapping.

Changing adenylation domain specificity is a huge step towards being able to optimize potential drug compounds, but QGEM 2016 wants to introduce more flexibility and control of adenylation domain substrate modifications. It has been observed that modules often contain domains which modify their substrate such as ketoreductases, and oxidases embedded within the adenylation domain. Therefore, the site of these domain additions is a great target for introducing engineered auxiliary domains which can add additional functionality to adenylation domain substrates.

Using the IndC module tagging system discussed in Phase 1, it would be possible to apply high throughput methods to screen for modified modules. During the summer QGEM will create an automated software pipeline which uses supervised logistic regression to reduce the search space for determining the genetic modifications needed to change the substrate specificity of adenylation domains. We also aim to



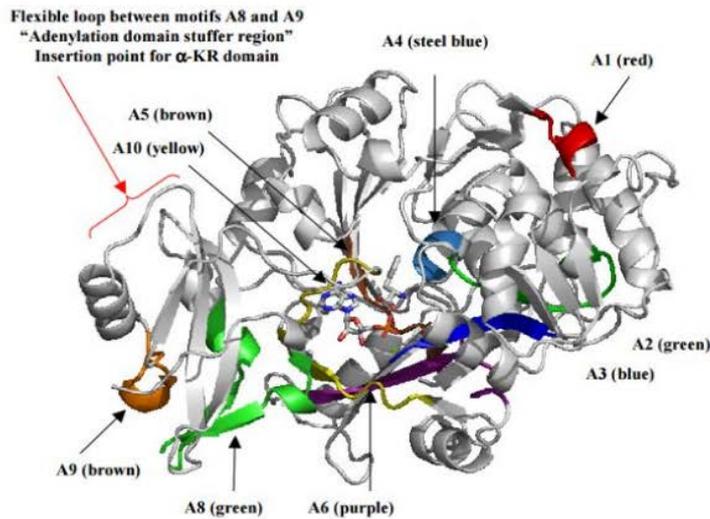


Figure 3. Ribbon diagram of the GrsA phenylalanine adenylation domain crystal structure. Adenylation domains are highlighted and hypervariable loop region for auxiliary domain insertion is highlighted.

design software which easily creates primers needed to insert auxiliary domains into modules through Gibson assembly by targeting the non-conserved region between A8 and A9 motifs in the adenylation domain.

We will demonstrate the potential of the software by creating a genetic construct using the tagging system previously mentioned. A module of interest (MOI) from a NRPS system will be isolated

and joined to an IndC module in order to create a construct where the MOI can be easily optimized through high throughput screening. Modifications of the MOI will be carried out in concert with the guidelines provided by our software pipeline. Lastly, once the MOI has been optimized to incorporate the substrate with the modifications of choice, we will be modifying the NRPS gene cluster as discussed in Phase 3.

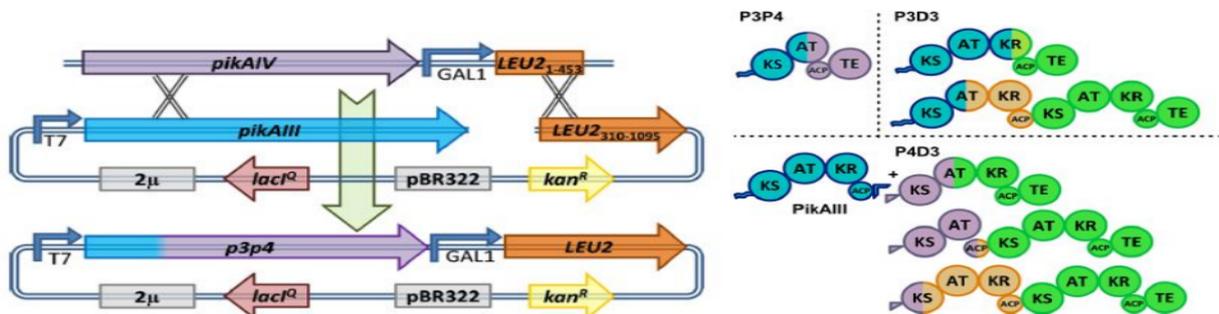


Figure 4. (Left) Proposed method of using homologous recombination in yeast to construct chimeric NRPS systems similar to those produced with PKS. *S. cerevisiae* Leu(-) mutants were grown on Leu(-) media. Successful recombination resulted in recovery of Leu synthesis. (Right) Chimeric PKS systems created using homologous recombination. Possible recombination sites are demonstrated.



Phase 3: Homologous Pathway Synthesis

In order to construct novel pathways and compounds, we are turning to homologous recombination. This technique is inspired by the evolution of NRPS systems. By studying the modules of NRPS domains as well as conserved regions in their structure we hope to use a homologous recombination system for stitching together different NRPS modules and letting nature do the heavy lifting for us, similar to a technique used for polyketide synthesis (PKS) (Figure 4).

Due to a limited number of characterized genetic clusters, combining modules in an order that preserves essential domain interactions such as docking domains and critical proteinprotein interactions may prove difficult. Hence we will construct a NRPS system that is as close as possible to the desired sequence using homologous recombination and then change the specificity of certain adenylation domains as well as add auxiliary domains.

Multiplex automated genome engineering (MAGE) will be used to efficiently change the Stachelhaus residues of the adenylation domain of interest found in the gene cluster. This has been done effectively before in literature and has been shown to

preserve biosynthetic compound production when compared to whole module swapping [4]. Additional auxiliary domains may need to be added to the module. In order to insert the domains, we will use MAGE to insert a protospacer adjacent motif (PAM) site in between the A8 and A9 domains and use a CRISPRCas9 system to insert the engineered auxiliary domain(s) into the module of the NRPS gene cluster.

By accomplishing the proposed three phases, we will be able to create a pipeline to optimize drugs produced by NRPS systems as well as produce novel drugs. This would allow one to rapidly construct and modify cyclic peptides which are difficult to chemically synthesize. This proposal has huge implications for the pharmaceutical industry by helping to create biosynthetic drugs which are safe for consumers, therapeutically effective, and more economically sound than chemical synthesis. The creation of this pipeline has the ability to revolutionize the way we look at biosynthetic drug design and optimization.



Human Practice

1) Second Annual Synthetic Biology Research Seminar

On September 28th, 2016 Queen's iGEM will be hosting its second annual Synthetic Biology Research Seminar. We believe that it's important to educate undergraduate students, high school students as well as the general public about synthetic biology technology and what it can offer to their daily lives. Since synbio is an emerging field, we are compelled to define synthetic biology to the public and give them examples of successful synbio technologies that have improved society in some way or another. Furthermore, at these seminars we also discuss out research findings from our studies over the summer. This is a fantastic opportunity to promote Queen's iGEM and the research that we do as well as engage with the public on the ethical facets of synthetic biology and the benefits the technology can offer to us.

2) Course Development APSC 100: Engineering Practice (Module 3: Engineering Design)

APSC 100 is a first year engineering practice course taught at Queen's University. The design module, a twelve week project, offers students an introduction to teambased work. A team is formed of four to five students who are paired with an upper year advisor and a community client. Students are guided through the design process as they work to solve openended design challenges which often emphasize prototype development and system modeling, two critical overarching concepts in engineering. QGEM partnered with APSC 100 again this year to offer an engineeringrelated project with a focus on biology. While engineering is a key component of the design and modelling aspects of our project, the team has traditionally struggled to breach the gap between the biological sciences and engineering within the university. This partnership is QGEM's way of strengthening the relationship and bridging the gap between these two disciplines at Queen's University. By incorporating synthetic biology and the considerations of our work into an engineering project, we hope to introduce the students to



the possibilities of merging applied science with more traditional forms of research, with the aim of promoting interest in the biological applications of engineering. This year, QGEM entrusted APSC 100 students with the task of designing and producing either an incubator, a thermocycler, or a transilluminator in the most efficient way possible, integrating skills from their other courses.

3) Interview Series

To supplement our project this year on novel drug discovery and optimization, as well as related projects engaged by other iGEM teams, we've decided to film a three-part interview series. This interview

series aims to address overarching questions related to biosynthetic production, aspects of drug design, metabolite reprogramming and much, much more from a variety of unique perspectives. We welcome you to join us through each of the three segments as we explore perspectives from research, the industry, and academia! See our blog at queensigem.ca for more updates on the interview series as well as some short clips from the interviews we've conducted!

4) Synthetic Biology Summer Enrichment Program

This summer, Queen's iGEM decided to hosted three, weeklong synthetic biology courses for middle school



Students fascinatingly observe DNA extracted from bananas.





A fun picture of our grades 7&8 students.

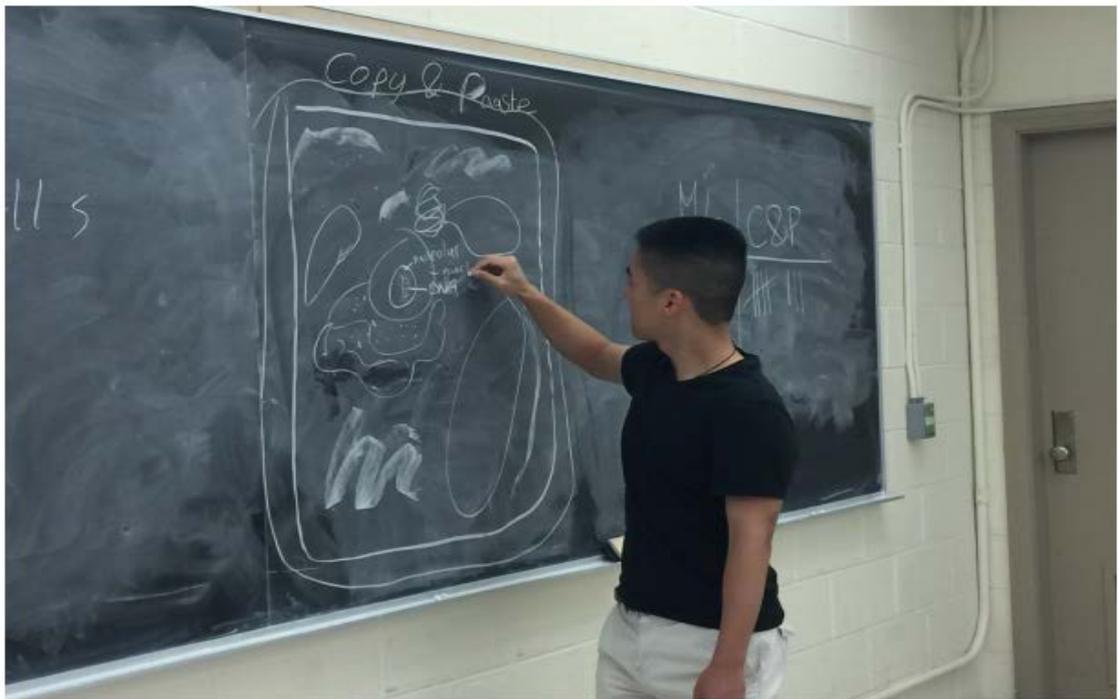


Our grade six students thoroughly enjoyed extracting DNA

and high school students through the Summer Enrichment Experience at Queen's (SEEQ) program hosted by the Enrichment Studies Unit (ESU). We believe that synthetic biology is an emerging field that students are seldom educated about whether that be from elementary school, high school, and or even postsecondary institutions. We believe that synthetic biology is the future, and therefore educating young people about it is a crucial component to raising public awareness. It is also an excellent way to promote iGEM to high school students on the verge of attending postsecondary institutions, for which they will have gained prior exposure that can potentially influence them to join an iGEM teams.

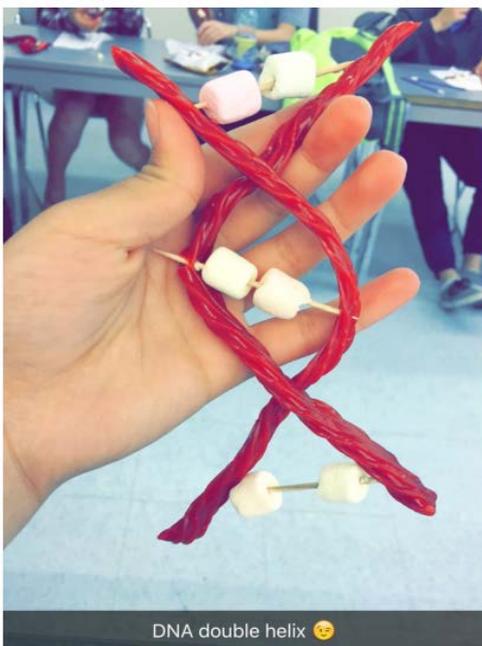
Our lesson plans were modified from the William and Mary 2013 iGEM team's Synthetic Biology Curriculum. We included many interactive activities (such as making cell models using dessert treats, DNA helices using Twizzlers®, and extracting DNA from bananas) to ensure that the learning process is hands-on, engaging, interesting, and drives the important points home. We were fortunate to have run these courses in partnership with the Queen's Enrichment Studies Unit (ESU) who helped us with much of the advertising and enrolment process for our synbio course.





Instruction Yifei Wang draws a diagram of a plant cell.

Our synthetic biology course ran for 3 weeks in total. The first week featured grade 6 students, the second week included students from grades 7 to 8, and the final week was taught to high school students. Overall, the students were lots of fun to teach, and consisted of very bright young individuals and budding scientists who asked very advanced and probing questions for their age. We received great reviews from the students on the course and it was a pleasure to teach students so eager to participate and learn!



The students constructed very elegant DNA helices and cells made from dessert treats!



Sun Yat-sen University

SYSU-CHINA



SYSU-CHINA 2016



@iGEM_SYSUChina

Team SYSU-China has been established since 2011, then recruits undergraduate students from School of Life Sciences, Department of Mathematics and other institutes.

Team

Team SYSU-China has been established since 2011, then recruits undergraduate students from School of Life Sciences, Department of Mathematics and other institutes.

Team SYSU-China 2016 invokes experiences of their own as well as those of their predecessors to address some issues related to the administration, management, and leadership of an iGEM team.

Project

A lack of techniques to figure out cells undergoing different number of cell-cycle in their lineage has limited our ability to evaluate the efficiency of stem cell therapy and investigate the mechanism behind it.

Here, we SYSU-China, describe Cyclebow, a system for labeling cells undergoing different number of cell cycles after a specific state in the lineage based on cyclic promoters combined with recombinases and fluorescent proteins.

We intend to demonstrate imaging of up to three cell cycles in a specific lineage, which can help tracking the proliferation, differentiation and migration of stem

Sun Yat-sen University

SYSU-MEDICINE



@iGEM SYSU-MEDICINE



IGEM SYSU-Medicine



SYSU_MEDICINE@163.com

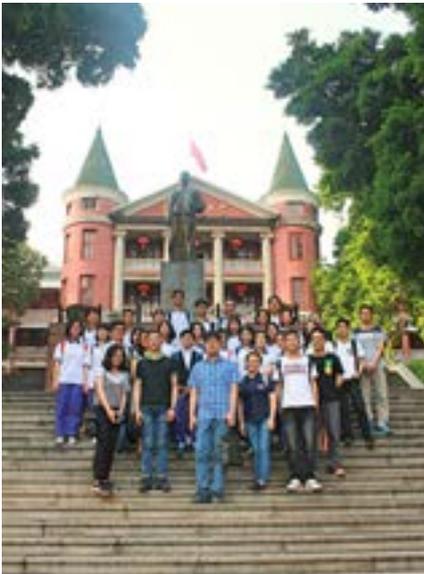


SYSU_MEDICINE

This is a fresh team starting from November, 2015. Unlike most teams on the platform of iGEM competition, the leading players of SYSU-MEDICINE are medical students.

Team

This is a fresh team starting from November, 2015. Unlike most teams on the platform of iGEM competition, the leading players of SYSU-MEDICINE are medical students. The whole team, jointing forces of students from school of medicine, school of public health, school of communication and design, school of mathematics and school of software, is a diversified team. With the power of synthetic biology and the valuable chance of taking part in the iGEM, we are looking forward to enhancing the existing treating method as well as producing therapeutic tools of better efficiency and pertinence.



Project

With great power to suppress adaptive immune system as well as innate immune system, mesenchymal stem cells (MSCs) are promising candidates for cell-based therapy to treat inflammatory diseases, such as IBD, diabetes, encephalitis, etc. However, clinical trials of MSCs have demonstrated that only a few MSCs can indeed arrive the inflamed tissue after systematic administration and exert their immunomodulatory function due to the inefficient homing ability of MSCs.

This year, next generation of MSCs are coming. In our project, we will

1) Empower MSCs with a series of chemokine receptors in order to ensure its effective homing.

2) Introduce several kinds of positioning system, such as luciferase to locate in vivo MSC and assure their arrival at the inflamed tissue.

3) Design a switch to kill MSCs when they differentiate into other types of cells.

Finally, we will confirm our modified MSCs in animal models, such as IBD and DTH.

Collaboration

KERWORDS about our project:

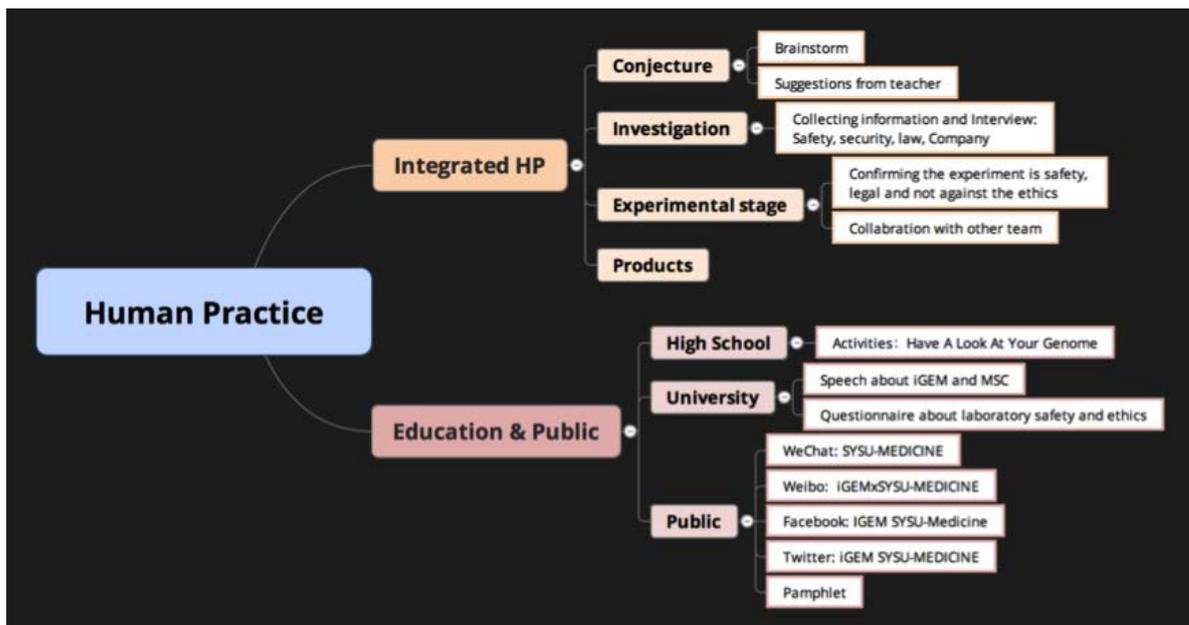
- (1) Mesenchymal Stem Cell (MSC), stem cell
- (2) Chemokine receptor
- (3) Fluorescence protein: luciferase, dTomato, eGFP, eBFP

- (4) α -SMA and its promoter
- (5) Apoptotic gene of MSCs
- (6) IBD and DTH animal model
- (7) Mathematic modeling
- (8) Gateway, single point mutation and other experimental skills etc.

If your project has any connection with these key words, or you just want to know more about us, do not hesitate to contact us.

Human Practice

The human practice of our team can be divided into two parts. One is integrated human practice, which is bounded with experiment together, serving the experiment group. The other one is about public education, which is to help the public to know more about our project, synthetic biology and iGEM.



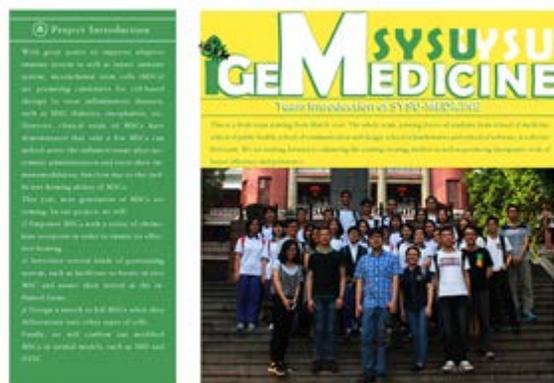
Connecting with the experiment, all the team members had a brainstorm about what we were going to do. Several members of our team were having pharmacology course at that time and knew some traditional anti-inflammatory agents. At the meantime, the instructor of our team experts in Mesenchymal Stem Cells (MSCs), who told us about the anti-inflammatory effects of MSCs. However, during the discussion, we found that both traditional anti-inflammatory agents and existing MSCs have effect on the whole body instead of working just at the inflamed site. Then, an idea come to our mind-modify the MSCs to home to the inflamed site. For HP, the second step was to search all the information about MSCs like ethics, law, safety, etc. To learn more about the background of MSCs, apart from collecting information and literature reading, we had interview of doctors, patients, companies and legislators. Experiment started at the third step, during which HP group had to confirm whether the experiment is safe, legal and not against the ethics, and moreover, asking for collaboration with other teams. The fourth step—products, is what we are going to do when finishing the experiment.

"Human Practices is the study of how your work affects the world, and how the world affects your work."— Peter Carr, Director of

Judging. As what I mentioned above is "how the world affects our work", then the next part is "how we are trying our best to affect the world". For offline activities, for example, "Have a Look at Your Genome". The objective of this activity is to help high school students to know more about synthetic biology.



Additionally, we sent out pamphlets.



For online publicity, we have set several official accounts and share a lot about our project.

Official Accounts of WeChat: SYSU-MEDICINE

Weibo: iGEMxSYSU-MEDICINE

Facebook: IGEM SYSU-Medicine

Twitter: iGEM SYSU-MEDICINE

Official Accounts of WeChat: SYSU-MEDICINE

Official Email: SYSU_MEDICINE@163.com

Discussion

With great power to suppress adaptive immune system as well as innate immune system, mesenchymal stem cells (MSCs) are promising candidates for cell-based therapy to treat inflammatory diseases, such as IBD, diabetes, encephalitis, etc. However, clinical trials of MSCs have demonstrated that only a few MSCs can indeed arrive the inflamed tissue after systematic administration and exert their immunomodulatory function due to the inefficient homing ability of MSCs.

(1) Safety requirements. For lab work, we have always complied with the provisions of the laboratory safety in the Laboratory Biosafety Manual made by World Health Organization. Laboratory equipment, facilities, supplies, reagents, and animals are in accordance with the requirements of the processing and operation in Laboratory Biosafety Manual. According

to rating provisions in Laboratory Biosafety Manual, our laboratory consists of a laboratories of first safety level and secondary safety level, respectively for different experiments.

(2) In our first three weeks, we accept the training of laboratory safety and some experimental operation and in the end of the third week we have a

test, which consists of written test and practical operation, in order to ensure that our team member can operate in accordance with the laboratory safety manual and the relevant provisions. The training we accept mainly includes three aspects:

① How to protect themselves. At first safety level laboratory we should wear the lab-gowns, ready to wear gloves and masks. Slippers, shorts are not allowed to appear in the laboratory to minimize the exposed part of the body in the laboratory air;

② How to protect the experimental materials. Firstly, experimental materials - animal. Animals (mainly murine) are fed in the animal experimental center, and we strictly obey the regulation of the animal husbandry and breeding in Sun Yat-sen University and we properly deal with the corpses of mouse according to the relevant regulation in the Laboratory Biosafety Manual. Secondly, experimental material - cells. All cells experiments are operated in the clean bench or biosafety cabinet and who

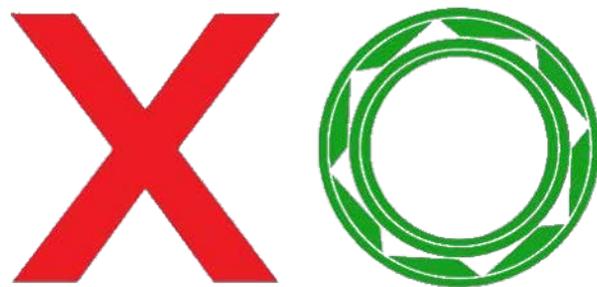
conduct the cell experiments have been specially trained to ensure that the cells won't be polluted and the experiments can go smoothly.

③ How to protect the environment and others. The reagents we use, animals and cell experimental materials have been harmless treated and will not be directly released into the environment. Animal experimental materials will be transported to specific department in Sun Yat-sen University and handled properly by them.

(3) Ethical issues. There are mainly two parts of the ethical tissues, the source of the MSC and the animal experiments. Our research is in accordance with Helsinki Declaration and the relevant policies and regulations in China, and we fully respect for the rights and interests of the donors, we communicate with the donors, get the permission of obtaining, and they sign the informed consent; As to the animal ethics, we submit the application for ethical animal experiments, and our application passed the examination of Sun Yat-sen University ethics committee

Instituto Tecnológico y de Estudios Superiores de Monterrey

TEC-CHIHUAHUA



MYXOBACTERIA
TEC-CHIHUAHUA



@iGEM_TecChih



iGEM ITESM Chihuahua



itesmigem@gmail.com

Our team is composed mostly by biotechnology students from different semesters but has also the collaboration of law, mechatronics and computer science students.

Team

Our team is composed mostly by biotechnology students from different semesters but has also the collaboration of law, mechatronics and computer science students. We all have different areas of interests but the thing we all have in common is that we are eager to learn and do something of value with that knowledge.



Project

Our Project: Myxobacteria as biological control in cultivations

The state of Chihuahua is the 2nd producer of alfalfa in Mexico and as we all know, phytopathogens are a great problem concerning agriculture, and frequently lead to great economic losses. Although chemical pesticides and fungicides have been used against these pathogens, they often result in the accumulation of toxic compounds or increase the resistance of the pathogens. This is why biocontrol using microorganisms has become an effective alternative of controlling plant pathogens.

For this alternative we decided to use a type of soil bacteria known as Myxobacteria, which are a common and diverse group of bacteria largely fed through predation and able to produce a wide range of secondary metabolites. We isolated this bacteria from our region and got proof that they are able to inhibit fungus and other bacteria. This may be due to competition for nutrients or the production of antifungal compounds. Therefore, we saw a potential project on enhancing Myxobacteria's properties to attack some specific endemic fungi.

To achieve this, we intend to create a BioBrick™ that can give this bacteria the ability to resist extreme weather, and enhancing its antifungal capability. For this investigation we took samples from damaged alfalfa crops found nearby Chihuahua city from which we pretend to isolate phytopathogenic organisms and prove the efficiency of our modified bacteria making confrontations between them. This will also help us broaden the impact in other crops as well, such as chili and potato.

Human Practice

Our Human Practices team works on an analysis of the implications of Synthetic Biology from the Ethical, Economical and Legal perspectives, with a focus on the Latin-American dimension, for the diffusion of Synthetic Biology in the light of a responsible progress.

We intend to give Synthetic Biology diffusion by contacting local farmers and stakeholders from the agricultural sector, in order to learn about their positions, targeting their main necessities, as well as knowing how would our project be perceived.



Tel-Hai academic college

TEL-HAI



iGEM Tel Hai



igemtelhai@gmail.com

Cystic Fibrosis is an autosomal genetic disease. It is the most common lethal genetic disorder affecting mostly Caucasian populations, but affecting populations worldwide as well essentially.

Project

Cystic Fibrosis is an autosomal genetic disease. It is the most common lethal genetic disorder affecting mostly Caucasian populations, but affecting populations worldwide as well essentially.

It is most known for affecting the lungs (through accumulation of mucus, causing difficulties in breathing) but it also leads to problems in multiple additional organs, such as the liver, kidneys and pancreas. In addition, mutations correlated with Cystic Fibrosis have been seen to be in correlation with male infertility.

Currently, there is no cure for Cystic Fibrosis, and most patients with the disease pass away by the way of approximately forty, though multiple medicines have been used or developed in order to alleviate multiple symptoms the patients have been

found to suffer from, aiming to improve the quality of life of those for as many years as possible.

Of the hundreds of mutations affecting the CFTR protein and indicating the existence of CF amongst patients, the most common is $\Delta F508$. This mutation leads to a damaged CFTR protein due to the loss of one Phenylalanine amino acid on the 508th place on the protein. The protein folds incorrectly, and leads to an imbalance in osmosis, therefore leading to salty skin, and negatively affecting sweat, mucus and digestive juice accumulation.

Our aim in this project is to correct the CFTR gene in the epithelial cells of Cystic Fibrosis patients by using the CRISPR technology. The conjugation of the CRISPR/Cas9 Plasmid/ B Subunit of the Cholera Toxin will allow tissue specific delivery, and therefore assure a reversal of the $\Delta F508$ mutation.

Human Practice

actice work is essentially divided into a few smaller categories:

1. Raising awareness with the help of celebrities
2. Classes for younger students on Cystic Fibrosis and entrepreneurship as a whole in the field of science and genetic engineering specifically.
3. Lectures at local bars and restaurants

We worked with the Israeli Cystic Fibrosis Organization in order to raise awareness country-wide for the disease and the affects it has on patients and their families medically and economically.

We met with multiple patients with the disease in order to hear their stories, and gain a deeper understanding of the daily battles they fight, and the fields which still need much more improvement, with the hopes of meeting their needs to the largest extent we can as a group.

Discussion

Just A little Food for Thought...

Ethics and Genetic Engineering- friends or enemies?

For years this debate has been a heated one and has attracted scientists, religious figures, social scientists and more.

From the religious perspective- how dare us play God and interfere with the divine work. On the other hand, if we can help save lives, could it be an exception?

From the scientist perspective- if we have the technology and ability, why not essentially?

From the social scientist perspective- At what expense will this all come? Does using embryos mean denying the birth of a potential baby in the future? Are we researching in the safest and most responsible way possible?

Does beginning genetic engineering necessarily mean we will begin creating "designer babies" or will we know when to stop, so as to utilize technology only for necessities and life- saving instances?

Tianjin University

TIANJIN



@iGEM_Tianjin



yuanma@tju.edu.cn



iGEM2016Tianjin

iGEM Team Tianjin is one of the first batch of iGEM team in China, which was found in 2007 and this year is our 10th time to participate in the iGEM competition.

Team

iGEM Team Tianjin is one of the first batch of iGEM team in China, which was found in 2007 and this year is our 10th time to participate in the iGEM competition. Look back on the history of Team Tianjin, we have won 4 gold medals totally, and last year we won the Best Energy Project Prize.

Project

Our project this year is about the biodegradation of PET, a widely-used plastic material. We use two enzymes, PETase and MHETase, which were found this March and certified to have much higher activity than any PET degrading enzyme found before in room temperature. We express the enzymes by *Saccharomyces cerevisiae*, a well-researched and generally-regarded-as-safe (GRAS) microorganism. The products of the degradation reaction are terephthalic acid (TPA) and ethylene glycol (EG), which are also toxic to environment.

We construct a co-culture system and introduce the *Rhodococcus rhodococcus*, which can use TPA as its carbon resource and *Pseudomonas putida*, which can degrade the EG. If possible, we also want to introduce Cyanobacteria to our system to transform optical energy to chemical energy and make our system autotrophic, so that we can apply our system in degrading the PET products in nature.

What is impressive is that the *Pseudomonas putida* can produce PHA, a kind of biodegradable plastic, so we apply the particular promoter from former iGEM project to control the VanX gene expression, which can cause bacterial lysis and release PHA.

Another direction of our project is directed evolution of the key enzyme, PETase. Since we do not know the 3D structure of PETase, we turn to other enzymes which can degrade PET and analyze the possible amino acid sites which is likely to affect the degradation activity and then we design 21 possible mutant and realize them by site-directed mutation. We also apply error-prone PCR to randomly mutate our PETase gene and we are looking forward to obtain mutant which has higher activity than wild-type enzyme.

Human Practice

We have helped the Tianjin University of Science and Technology to establish their own iGEM team, two students from there have studied with us for a whole month. Team Tianjin, BIT, BIT-China, Jilin_China, and CGU_Taiwan have set up a league and we can easily communicate our project and seek for help from each other. Apart from these teams, we also seek for collaboration everywhere. For instance, we help the HFUT-China to test their software. We have communicated with other university and asked for bacterial or plasmids we need. We will go to the Dagang No.1 middle school in Tianjin next month to give a speech about synthetic biology and help them establish their own iGEM team. We also collaborate with some famous corporation for financial support for our experiments, competition and travel. We also conduct our human practice online. We have our own Tweet account and WeChat public platform to inform our recent work to other teams and students. We have made a questionnaire and spread it online and we have received 686 effective results totally. (Turn to appendix file for details).

Tongji University

TONGJI_SHANGHAI



will wang/Tongji Shanghai



+86 15821399962

The up-conversion nano particle(UCNP) is a luminescent material which converts 980 nm light into 670 nm light. The carbon dot serves as photosensitizer intakes 670 nm light and brings heat effect, helps lysosome escaping.

Project

The up-conversion nano particle(UCNP) is a luminescent material which converts 980 nm light into 670 nm light. The carbon dot serves as photosensitizer intakes 670 nm light and brings heat effect, helps lysosome escaping. We joined two materials with SO₂ and transport the complex into the breast tumor cell, start apoptosis by ROS from sodium copper chlorophyllin on C-dot. To amplify the heat therapy, we designed a virus expressing P53 in the tumor cell by cancer specific promoter. During the therapy, we load the virus into all cells, express P53 only in tumor ones. Afterwards the UCNP in injected, guided by the light to trigger heat therapy. This treatment is considered to be a better targeted however less toxic method to cure cancer.

Human Practice

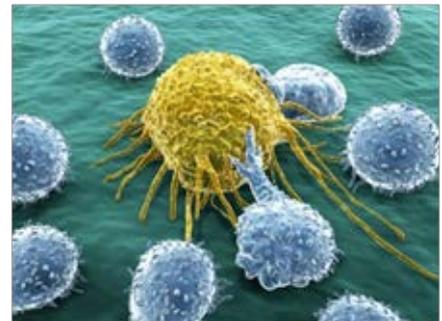
•exhibition in Shanghai technology museum

Our team delivered an exhibition at Shanghai science and technology museum at 8/16/2016 together with three other schools. At the scene we use a computer to broadcast the acknowledgement of synthetic biology and IGEM. Further more, we built our own post to help illustrate what our work is all about to visitors. It is a very interesting experience!



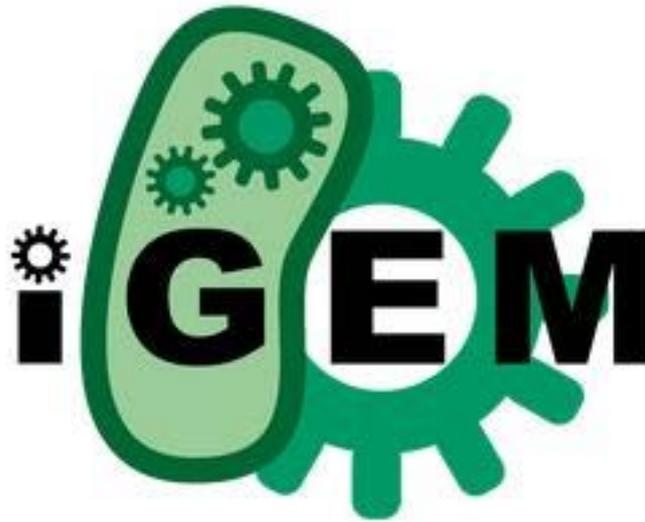
Discussion

What is cancer? For many years people considered it a terrible word to be heard. Meanwhile, if we see at the angle of the homo sapiens, we may understand that cancer is the weapon to promote evolution. There are typically two reasons to trigger cancer, one of which is defective gene before birth, the other is bad living environment causing the expression of cancer gene or inhibit of cancer inhibitor gene. Never the less, both triggering factor can lead to one consequence, which is defective gene passing on. In order to prevent that, cancer is developed in our body to destroy an individual from passing his gene. As most of the immune system can't avoid sacrificing, if we consider cancer a kind of eternal immune system, it sacrifices single body to protect the human gene, and in the end leads to better and healthier human.



University of Chinese Academy of Sciences

UCAS

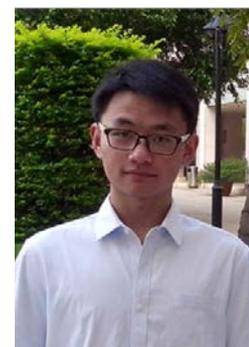
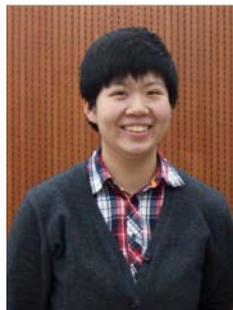


ucasigem@163.com

UCAS iGEM is devoted to degrading antibiotic residues in waste water treatment plants(WWTPs) or hospitals. We mainly focuses on tetracycline, which is one of the most abundant antibiotic detected in Chinese city rivers.

Team

This year, our team have fifteen members from different majors, three advisors, two instructors and four Principle Investigators. Ten of the fifteen undergraduates are junior students. They are Boyi Wang, Qingyang Tan, Wenbo Wang, Ye Liu, Ye Yang, Yiqun Dang, Yongming Li, Yue Xu, Zepeng Mu and Zhuoning Zou. The others are sophomeres. They are Cai Ruiling, Huang Jianyi, Jiang Yueren, Liu Yushan and Mo Yajin. The four Principle Investigators are Chunbo Lou, IM-CAS, Jianguyun Wang, IBP-CAS, Xianen Zhang, IBP-CAS, Xiaoguang Sheng, UCAS. The two instructors are Hao Jiang, IM-CAS, Xiaohong Liu, IBP-CAS. The three advisors are Cheng Hu, IBP-CAS, Hua Li, IBP-CAS, Sha Wei, IBP-CAS.



Project

The discovery and mass production of antibiotics has saved the lives of hundreds of millions of people. However, antibiotics residues in nature and industrial products make people exposed to antibiotics on a daily basis. The consequence is not clear yet, but some researches have shed light on its role in antibiotic resistance or obesity.

UCAS iGEM is devoted to degrading antibiotic residues in waste water treatment plants(WWTPs) or hospitals. We mainly focuses on tetracycline, which is one of the most abundant antibiotic detected in Chinese city rivers. Screen of enzymes including TetX makes the system MORE EFFICIENT, whilst the design of a sensor with a positive feedback and a TA module-based kill-switch makes the system SAFER and SMARTER. Our engineered bacteria will not express degrading enzymes unless antibiotic is presented, and it effectively kills itself after leaking into wrong places.

Collaboration

As for collaboration, we provided the gene of toxin CcdB for ShanghaiTech University. They are also working on toxin genes as we are doing and we are glad to share what we have. We are also participating in this year's CCiC held at SYSU in September. As one of the biggest events in China, we are looking forward to meeting iGEMers from all over the country. We are having some problems with modeling, and we would be grateful if other team could help us with this.

Human Practice

For human practices one of our focuses was in students' education.

We delivered a speech to the students from many different high school, aiming to explaining the students what the igem was and the current situation of antibiotics. After our speech we had a communication with them, not only they gained some knowledge about igem and started to concern about the antibiotics but we were inspired by their questions and opinions.



Beside delivering speech, we are also helping the igem lab of Beijing National Day School analyze their data and lending equipment to them.

Another important part of our human practice is talking with different igem team or experts of synthetic biology.

Up to September 2016, we have taken part in conferences with Peking University, Xiamen University, Tianjin University and Beijing Institute of Technology. We also met Mr. James Schroeder, who





is a Penn Alumni & Synthetic Biology Advisor, at World Financial Center in Beijing. He has collaborated with leading Synthetic Biology organization SynBERC. Mr. Schroeder discussed the emerging synthetic biology market in China, showing us how works done in labs can serve society and make profits.

Discussion

Antibiotics has played an important role in people's health and saved many lives. However, the misuse and overuse of antibiotics can result in some harmful consequence to human health. It may kill the normal bacterial colony in human body. It may cause anaphylaxis or obesity. The most dangerous consequence is that it can result in antibiotic resistance. With long-term use of antibiotics, the majority of sensitive strains have been killed, but the resistant strains has survived and multiplied. Hence, more and more bacteria can protect themselves form antibiotics. Resistant microbes are increasingly difficult to treat, requiring alternative medications or higher doses—which may be more costly or more toxic. Examples of drug-resistant bacteria are: methicillin-resistant *Staphylococcus aureus* (MRSA), vancomycin-resistant *S.aureus* (VRSA), extended spectrum beta-lactamase (ESBL), vancomycin-resistant *Enterococcus* (VRE), multidrug-resistant *A. baumannii*(MRAB). Antimicrobial resistance is on the rise with millions of deaths every year. A few infections are now completely untreatable due to

resistance.

In China, ~147,000 tones of antibiotics were produced annually. Waste water from factories producing antibiotics is heavily polluted with antibiotics. It is estimated that the amount of antibiotics discharged at rivers in China is even higher than that consumed in some northern European countries. We hope that we can degrade antibiotic efficient, safely and smart by our well-designed biological system. What's more, we hope to make more and more people realize the possible harm of misuse and overuse of antibiotics.

In our design, the degrading enzyme will be expressed only in the presence of antibiotics, like tetracycline. Being expected to stay in given working condition, the bacterium effectively kills itself when leaks into the unwanted places, which lack a chemical provided in the working environment. By putting the toxin protein and degrading enzyme on the same plasmid, we intend to restrict the spread of antibiotic resistant gene by preventing horizontal-gene-transfer.

To make our system more efficient, we screened several potential enzymes, including tetX. We utilized varied methods like LC-MS to evaluate the activity of enzymes both qualitatively and quantitatively.

Concerning biosafety and bioethics, we designed a TA module-based kill-switch, where the toxin is neutralized by its cognate antitoxin when antibiotic is abundant. In low antibiotic concentration, the express of antitoxin is repressed, leading to a higher toxin/antitoxin ratio in the cell, thus kills the bacteria.

University College Cork

UCC_Ireland



@ucc_igem



UCCIgem



corkigem@gmail.com

UCC iGEM 2016 are the third team from UCC, Ireland partaking in the iGEM competition. In 2014 our university entered the first Irish team into this competition.

Team

WHO ARE WE?

UCC iGEM 2016 are the third team from UCC, Ireland partaking in the iGEM competition. In 2014 our university entered the first Irish team into this competition. Our team this year is composed of 7 members. There are 2 medicine students (James Meeke and Aoife O'Brien Horgan), 2 pharmacy students (Kevin Ryan and Donnacha Fitzgerald), 2 genetics students (Amy Bergin and Seema Subedi) and a biomedical science student (Regina Walsh). We are all third year students and our team leader Brandon Malone is in 3rd year Pharmacy.

In order to become a member, all applications had to be submitted by the beginning of February. There were three components to our initial application, these included: (i) Curriculum Vitae (ii) An original idea on how synthetic biology can be used to solve a particular problem in the world (iii) A brief illustration on why we wished to partake and which skills and abilities we could add to the team.

After narrowing down applicants, several were invited to partake in an interview. We were informed of our acceptance to join the team by the end of that month. To learn more about our team check out our Youtube channel where we have a Meet the Team video!

Our project name this year is Limitless Lactis, as we are utilising *Lactococcus lactis* as a protein delivery platform for disease treatment. Potential applications which we have investigated include vaccination strategies and macrophage modification. This modification is enabled via the use of CRiSPRi. CRiSPR is a unique genome editing tool that enables precise genetic manipulation. This platform may be employed to modify the phenotype of other phagocytic cells associated with diseases such as cancer.

We have also developed a vaccination platform against leishmaniasis, a neglected tropical disease increasing in geographical distribution. LJM11 is an immunogenic salivary protein of the

sandfly vector, *Lutzomyia longipalpis*. Our inexpensive platform, through simple oral administration, has the capacity to deliver this protein to antigen presenting cells, and potentially immunise against the lifecycle of leishmaniasis.



Left to Right: Seema Subedi (Genetics), James Meeke (Medicine), Brandon Malone (Pharmacy), Donnacha Fitzgerald (Pharmacy), Amy Bergin (Genetics) Regina Walsh (Biomedical Science), Aoife O' Brien Horgan (Medicine)

Project

Vaccines prevent 3 million deaths and 10.5 million cases of infectious diseases every year. One of the major components of our project involves developing a protein delivering platform to act as a vaccine. This can be used as a prevention for a neglected tropical disease, namely leishmaniasis. Our inexpensive platform through oral administration has the capacity to deliver an immunogenic protein to antigen presenting cells. This could potentially immunise against the life cycle of leishmaniasis. This disease affects approximately 1.3 million people every year. It is most prevalent in the Americas and Asia.

Leishmaniasis is caused by a protozoan parasite and is transmitted by the bite of the female sandfly. There are 3 forms of leishmaniasis ; visceral, cutaneous and mucocutaneous. The strain we are targeting is *Leishmania infantum* which is transmitted by the sandfly *Lutzomyia longipalpis*. This strain is particularly prevalent in Honduras, which is a country in Central America. There are 1500 cases reported annually

in Honduras. The treatments which are currently available are lengthy and onerous. Hence, our vaccine could present a welcome alternative to current elimination strategies.

The cases in the Middle East are also of particular note due to the Syrian refugee crisis, which has been prevalent in the news over the past year. *L.tropica* (a species of *Leishmania*) which originated in Syria now accounts for 90% of cutaneous leishmaniasis cases. As Leishmaniasis is a communicable disease, the vast number of migrants entering EU from Syria could present a risk to public health. Although with our project, we are not targeting this particular strain, a similar basis could be used to produce a vaccine for EU citizens in the future. How are vaccines regulated in the European Union?

Discussion

EU VACCINE REGULATION

In general there are six main stages in development of new vaccine; exploratory stage, pre-clinical stage, clinical development, regulatory review and approval, manufacturing and quality control. In the exploratory stage, an understanding of the disease is obtained. The epidemiological data is collected and the correct antigen or protein (which in our case would be LJM11) to use in prevention or treatment is identified.

The second stage involves identifying the relevant antigen via screening, creating the concept and evaluating if the vaccine is efficient in vitro and in animals. Following this, the production procedure must abide by Good Manufacturing Practice standards.

This leads to the Clinical development stage, in which the vaccine is tested in humans. There are four stages involved over a number of years. Phase 1 trials involve determining if the vaccine is safe to use for humans and studying the nature of the immune response the vaccine evokes. Leishmaniasis is a disease of poverty. Thus, 1a trials would involve

European volunteers and subsequently there would be 1b trials with populations in developing countries

Phase 2 trials are on a larger scale. They assess how effective the vaccine is against artificial infection, then the clinical disease. The safety of the vaccine along with side effects and immune response are all evaluated during this stage.

Phase 3 trials involve hundreds of subjects in an array of sites. Efficacy can thus be determined under the natural disease conditions. A license can be applied for after a defined successful period in this stage. Phase 4 trials involve post market surveillance i.e. detecting any rare long term effects and long term efficacy.

Other than the extensive regulatory procedure, another consideration when bringing a vaccine to market is whether public opinion will be conducive to its introduction.

TO VACCINATE OR NOT TO VACCINATE, THAT IS THE QUESTION

There have been many controversies in the past in relation to the introduction of particular vaccines such as the HPV (human papilloma virus) vaccine and MMR vaccine. Personal beliefs and fear of side effects are two of the most significant reasons the public may be opposed to vaccine introduction.

Gardasil and Cervaxil which are HPV vaccines utilised in Ireland have been associated with 856 reports of side effects, since introduction in 2011. These reports have deterred many parents from giving permission for administration of this vaccine to their children.

In relation to the MMR vaccine, in the late 1990's a researcher released a paper attempting to illustrate the link between administration of MMR and incidence of Autism. Although the contents of this paper have been since retracted, the publication of the article caused a significant decrease in uptake of the MMR vaccine over the past 20 years. In the US alone the MMR vaccine rates decreased by almost 3% in one year (2008:93.5%, 2009: 90.6%).

From this controversy, it becomes apparent that having factual information available to the public is vital. This ensures people can be protected against preventable diseases. Misinformation as with the MMR controversy can lead to preventable fatalities.

Undoubtedly, vaccination is a societal bone of contention however immunisation is one of the most cost effective public health investments. Vaccines have helped to cut child deaths in half over the last two decades. By facilitating medical research and distributing factual information this trend may continue and hopefully completely eradicate the most common preventable diseases.



Univesity of Copenhagen

UNIK_COPENHAGEN



@CosmoCrops



CosmoCrops



igem16.cph@gmail.com

Univesity of Copenhagen has had an iGEM team for the past four years, with last year's team called SpaceMoss as the most successful with a gold medal and two nominations. This year, University of Copenhagen will be represented by ten highly dedicated students from several different life sciences programs of the university - These students are also commonly known as CosmoCrops.

Team

Assembling of the team

The headquarter for the iGEM team at University of Copenhagen is Center for Synthetic Biology and they believe that the best team will be assembled if the students have the best basis for what iGEM is. Therefore, they believe that it is the students from last year's team that can assemble the best team, because they have the knowledge to explain what iGEM is and what is the best way to think synthetic biology when thinking of iGEM.

In February members of last year's team held an introduction event where all interested could come and hear more about SpaceMoss and iGEM in general. Shortly hereafter, they opened for applications, so all interested could make an application to join this year's team. Because last year's team was very diverse, including students from Biotechnology and molecular biomedicine to physics and bio business, they wanted to make this year's team as diverse. Therefore, the application should include what program we are studying, what contribution we can make in an iGEM team, here meaning

laboratory experience, applying for funding or having contacts to the media. Furthermore, we should also take a personality test and include the result for the application.

After the deadline for handing in an application, members of last year's team sat down and looked all the applications through. Out of all the applicants, they choose the ten students they thought would make the most diverse team, seen from competences, studies and personality.

All of this added together gives CosmoCrops - we are ten students that represent 8 different programs from the faculty of Life Sciences at University of Copenhagen ranging from Biotechnology and Biochemistry to Computer Sciences and Mathematics. We believe that because of our diversity, we are able to work together as one team and everyone gets to use all their competences, instead of one person controlling everything. Therefore, we can definitely suggest making as diverse team as possible!



CosmoCrops that is representing University of Copenhagen at the iGEM competition 2016

Structure of the team

To make a team work most efficiently, we believe it is important to have some structure within the team so each person know what tasks one is responsible for. Therefore, we decided on five sub-groups where each person should belong to at least one depending on the interest and competences of the person. These sub-groups were made to include all areas within an iGEM project and they were called Organisation, Outreach, Funding, Green lab and Red lab. To make sure everyone is up-to-date we believe that meetings are important and should be often.

Organisation

The Organisation has the full overview of the project, and make sure that we will reach all the deadlines within the

time. Furthermore, they are the ones that will book flight tickets and hotels every time we are attending a meet-up or conference outside Copenhagen.

Outreach

The Outreach group will be the face of the team. The outreach people will be contacting the media to get articles out about the project and control the social media such as Facebook, Twitter and LinkedIn, and make sure those are updated. Furthermore, the outreach will also be those who are presenting the project at most of the lectures and conferences we are attending, and make deals with museums and such to get out to the public.

Funding

The Funding group is the people who will be applying for money so we can attend different conferences abroad and the Giant Jamboree in Boston. Furthermore, they are also the ones that will make sure that we get sponsorships from relevant companies. This could for instance be Eurofins or QiaGEN so the lab work will be cheaper.

Green lab

The Green lab consists of the biology-related people, and they will be doing most of the wet-lab work of the project. This could be the cloning and transformation work and setting up the bioreactor.

Red lab

The Red lab is on the other hand the physic lab of the project. This lab will be doing all the testing of the viability of our organisms in extreme environments.

Project

After the new team was assembled we should decide on a project. Very soon we all agreed on keeping the space theme that last year's team started. To get some ideas, our supervisors had an inspirational lecture during one of our first meetings to come up with ideas on what is trending and what topics that could be interesting. This gave a lot of ideas that the people in the Green lab started looking into. This was done by using literature search to find what is already known for the different topics and TedTalks to find some inspiration as well. After a few weeks where Green lab along with Red lab had brainstormed different projects, it was narrowed down to three; primarily continuing on last year's project, working with a co-culture or working with tardigrades. These three project and the ideas behind each project were then presented to our supervisors and the rest of the team, and feedback was giving on each of the projects. Hereafter, the team decided on a project - the co-culture!

Then it was up for Green lab to make a project description to define the scope of the project. This included finding the problem we want to solve and how to solve it. Space exploration is a costly affair but important for the life here in Earth. Most of the things we use every day were invented because of space exploration; therefore making space exploration cheaper will not only help exploring space

and maybe find life out there, but also push the innovative ideas forward here on Earth. Therefore, our aim is clear - Make space exploration cheaper!

The way we want to do this, is by decreasing the amount of material that is needed on-board on the space shuttle before launching, hereby the amount of fuel can be decreased. So we thought - why can't we produce the material we need out in space when we need them? Therefore, we are developing a biological system that can produce almost every kind of material and product that can be needed on space missions or on future settlements on other planets. This biological system consists of two organisms; *Synechococcus elongatus* and *Bacillus subtilis*. The *S. elongatus* will make sucrose out of sun light and carbon dioxide from the air. The sucrose can be used by *B. subtilis* to produce whatever compound we have modified it to produce. In a proof-of-concept, we are producing the bioplastic P(LA-co-3HB). The bioreactor with these two organisms are designed in such a way that they two organisms are physical separated, meaning that one of the organisms - *B. subtilis* - can be removed without disturbing the other organisms. This provide a bioreactor with a high modularity where one of the organisms can be removed and replaced with another engineered strain, so instead of production of bioplastic we get production of vitamin B12 - all of this without starting a new bioreactor!

Furthermore, the two organisms, *S. elongatus* and *B. subtilis* will be tested for their viability in low pressure, low temperature, high UV and microgravity - all parameters that are found in space. Hopefully some beneficial mutations can be induced that makes the organisms more able to survive those parameters. This is in short what we try to obtain this summer!



The Skype meeting with the iGEM team from Leiden, the Netherlands seen from the point of view of Leiden.

Collaboration

At CosmoCrops we are really eager to get to know so many people as possible, and therefore we would like to hear from so many different iGEM teams as possible! Collaboration is an important part of iGEM, because it is not only a competition but it is a place to meet new people and share knowledge within the field synthetic biology. Also, it is a possible to have fun with people you otherwise will not have met - therefore, we love having Skype meetings with people all around the globe.

We already have several different collaborations going, but could always use some more! We are working closely with the Leiden iGEM team, because we both work with space exploration. We are doing some low pressure and high UV experiments on their Escherichia coli strain and in return they test the viability of our B. subtilis and S. elongatus in microgravity.



Our first Skype meeting with the iGEM team from Imperial College London.

In the collaboration we have going on with the Imperial iGEM team, we are providing them with some data on how our organisms are growing in our co-culture under certain growth conditions. They are using those data to make a model on how different organisms are working together and when it is best to grow them together and how the amount of one does not out-grow the other organism. They will then do some modelling work for us, because we do not have time to do the modelling on how much sucrose our *S. elongatus* is secreting.

The last collaboration we already have going on is with the iGEM team from Istanbul, Turkey. We have a monthly skype meeting where we discuss the progress of the projects, and if we have any problems we talk about them and try to solve them together with our combined knowledge. And here we have not mentioned the two other teams from Denmark that always will be there to help if something happens or we have any questions, and the same goes the other way around.



Group picture of the teams that participated in the Danish Meet-up at the Technical University of Denmark.

We are looking for all kind of collaborations - this could be sharing of data, helping each other with making experiments or just monthly Skype meetings to discuss how the project is going. In short, we are working with co-cultures, *Bacillus subtilis*, *Synechococcus elongatus* and space explorations - so if your team is working on something similar then feel free to contact us. If not - then contact us anyways, maybe we have something in common we can discuss!

We can be contacted on our mail, Facebook, Twitter or LinkedIn - We are looking forward to hear from all of you .

Human Practice

iGEM is a competition within synthetic biology to promote the field of synthetic biology among students and to make synthetic biology more accessible by making it an open source. However, one important part of iGEM is not only the research and the BioBricks, but also the Human Practices. This is the things that go beyond the lab bench (which many of us are stuck at during the summer, where we work on iGEM), such as ethics, information to the public or intellectual property rights (IPR). Explained in other words; we need to think about what societal impact our project can have on the society and for the general public.

In CosmoCrops, we are working with something most people have a hard time relating to - space exploration. However, this does not hinder us in thinking and applying the Human Practices to our project, in fact we do think that Human Practices is one of the most important parts of our project. This is also why we for the first time ever, hosted a Human Practices Workshop in Copenhagen for the teams from SDU, DTU and Gothenburg-Chalmers, to help them see the beauty in Human Practices as well. Our Human Practices consists of ethics, collaborations and lectures/informative speeches for the public.

Assembling of the team

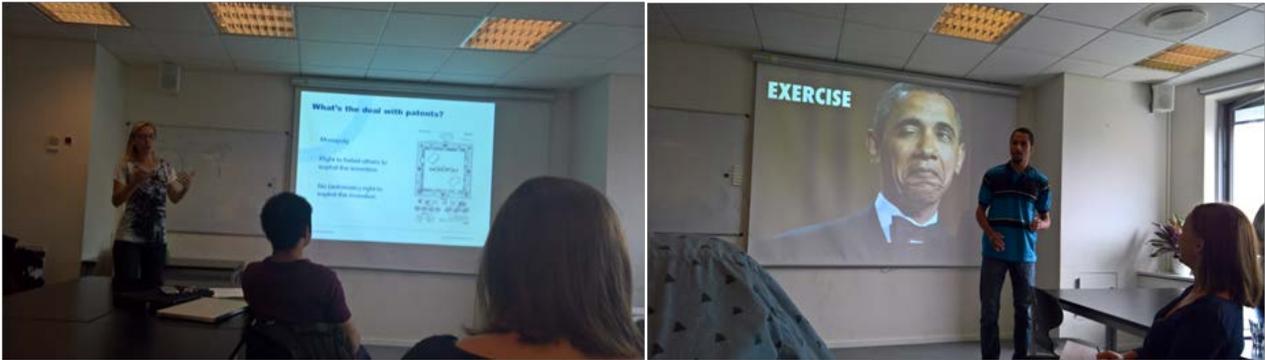
When doing a science project there is a tendency that scientists forget about the ethics - which that is very important. The importance of ethics cannot be stressed enough because it says something about the project and if it is morally right or wrong. But what is morally right and wrong? This is a subjective decision, because some people think it is okay to do something, while others think it is a wrong decision.

In our case, we work with a co-culture we want send to space. However, is it ethical correct to send living organisms into space and use them out there or on settlements on other planets? This is just one of many ethical questions we have, and there is no right or wrong answer to it. By sending living organisms to for instance Mars we potential disturb the ecosystem, which can be ethical wrong - the big problem is, what if the living organisms we bring up there by accident is set free? This could potential destroy the whole ecosystem on Mars or the two organisms we bring up there could outcompete all of the already present inhabitants of the planet hereby destroying a whole planet. This is also why it is illegal by law to introduce new organisms to new planets.

Like for every colonization, one of the big ethical and political issues is, who are going to own the place of interest? This has been seen several times during history, and it might be repeated if we begin to colonize Mars or other planets. Should the planet of interest be giving to the first to arrive or should it be divided up so every country that thinks they have a claim of the planet can get a piece of the planet? This might end up in a war - not on Earth but on the planet of interest, and the history will begin to repeat itself again.

Besides the ethical problems regarding space exploration, comes the whole ethics around synthetic biology in

general. Are we, as humans, allowed to change living organisms just because we can? You can argue that the nature have had millions of years to evolve and make the best organisms as possible, so if an organisms cannot make a certain compound, it is because the nature does not want that compound to be produce in that organism. As scientists and consequentialists we do not see this as a problem. By engineer *Bacillus subtilis* to produce bioplastic, we do not only make space exploration cheaper, which can make life on Earth easier as well, but we do also help the environment that already contain too much conventional plastic. This will help to maximize the overall good on the planet. However, we do acknowledge that not all have this point of view - especially not in the public. In Denmark there can be a very skeptical view of synthetic biology. This skeptical arise from different things, but one of the main reasons can be lack of knowledge - the Danish population does not simply know what synthetic biology is, and how it is even possible to combine synthesis with biology. Therefore, we think that it might be needed to give the population a bigger knowledge on synthetic biology and why it can maximize the overall good for the world. One way of doing this is to make public event where we go out and talk about synthetic .biology and our project, so the general public can get an inside to what synthetic biology is, and why it is actually important for the way they are living their life today.



Pernille Gojkovic from Højberg giving a talk about patents, and Joel from SDU doing a storytelling for the public speaking activity

Lectures/Presentations for the public

As mentioned above, one way to give the general a bigger knowledge of synthetic biology is to give public speaking about synthetic biology. This is something we regard as important for the future of both the iGEM teams here in Denmark, but also for the scientific community as general. Therefore, we have been out to several Danish media to get them to tell our story about synthetic biology, and how it can be used to help the future generations on Earth. This is for instance videnskab.dk and ing.dk there both is big scientific platforms in Denmark. Furthermore, we have been giving several different talks, for instance for the FantastiCon in Valby, Denmark. FantastiCon is a Sci-fi convention, where science fiction and reality was discussed, and how those might be mixed more and more in the future by persons like us. This was mostly for Science fiction fans that actually think that this topic is interesting, but it also important to come out with the message to younger minds that then might see the how fantastic synthetic biology can be, if it is used correctly. This we will do in the middle of September, where we for four days will give a presentation about our work and synthetic biology for around 500 high schools students. This will help to get the attention of the future generations and might get them interested in synthetic biology, just as we are!

Besides that, we will have presentations at the Planetarium in Copenhagen where all interested in space and space exploration can come by and hear more about our work among other space related things. We will also be found at the European Astrobiology Symposium in Athens, Greece in late September, where we are going to present our project to astrobiologists.

Collaborations

In part 2 of this issue, we mentioned our collaborations and why it is important to have collaborations in iGEM. However, this is also important for the Human Practices, because the collaborations will enable you to get some feedback on your project and what you can make better to make it more feasible for the general public to accept it.

Besides the more scientific collaborations we mentioned in the previously part, we have a close collaboration with the teams from the University of Southern Denmark, Technical University of Denmark and the University of Gothenburg and Chalmers in regard to the Human Practices. From the 19th to the 21st of August we were host for the very first Human Practices Workshop in Copenhagen, where the above teams participated. During this workshop the teams went through ethics and why it is important for synthetic biology, intellectual property rights and how they can be used to make your project into a start-up and what should draw your attention when talking patent and patentability. We also went through public speaking and how a good presentation should look like, and how it should be presented. This workshop gave some ideas in regard to Human Practices to all the teams that participated and how to think when thinking about Human Practices. The workshop ended with a presentation by Ana Sifuentes from the iGEM Headquarters who talked about Human Practices and why it is important for synthetic biologists and iGEM.



The teams from DTU, SDU, Gothenburg-Chalmers and KU eating dinner after the first day of the Human Practices Workshop.

Conclusion

In conclusion, we have considered the ethics as a big part of our project and the Human Practices. However, during the workshop we were made aware of the patents and the patent landscape which we also need to consider. A short look into this by using a few keywords of our project showed that there is a lot of patents within the scope of our research. This means we need to consider to either change the project if we want to make it into a start-up, because we cannot use already patented inventions or to buy licenses from the patent holders, which might be an expensive affair. Including the ethics, this is something we need to consider during the last two months of the iGEM competition and how we can apply it to our project.

UPO-SEVILLA



@iGEM_UPO



Igem Upo



igem.upo@gmail.com

Our team is divided into four parts dry lab, wet lab, communication and informatics. We are a multidisciplinary group from two universities that are willing to do its best to produce an amazing project. Do you want to meet us?

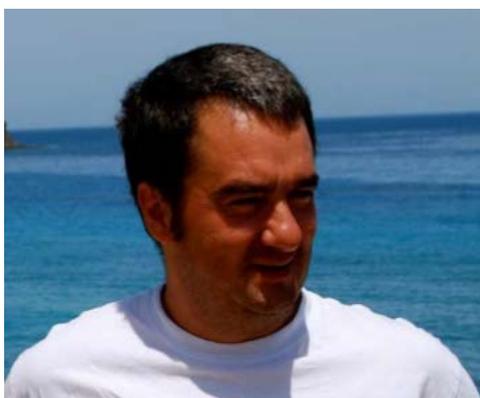
Team

Who we are: Team UPO-Sevilla

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Dry Lab

Alejandro Cuetos



Hi, I'm Alejandro Cuetos, lecturer on Chemical Physics Area. I am not sure why I am working with Bacteria, but I am finding it very funny. I am the supervisor of the modeling part of the project, where we are learning very interesting things about how to predict the biofilm behavior.

Francisco Javier Lobo



Hi, there!! I'm Francisco Javier Lobo, one of the members of the dry lab. I love Microbiology and Programming, so when they asked me to join I jumped in with both feet (and a computer). I think the iGEM competition is an opportunity to interact with people from all over the world. See you all!

Wet Lab



José Martín

Hello everyone! I'm José, a recent biotech graduate, and I'm really amazed by the potential of synthetic biology. I'm a member of the wetlab group and responsible for the propionate module. Let's make iGEM 2016 an unforgettable experience together!

Laura Claret

Hi! I'm Laura Claret, a graduated student of the Biotechnology Degree in our University. I'd never thought I would be selected for such a cool thing as iGEM, but here I am, in charge of half the workload of the Micro Wet Lab. Hope you find our project as bewildering as I do!



Álvaro Escobar



Hi iGEMers and current people! I'm Alvaro and I'm a biotechnologist. I think I'm a funny and peaceful guy, and I love making jokes. I cannot yet believe that I was chosen to have the opportunity of being part of the iGEM contest. I'm excited for being in touch with teams of the whole world, and I'm looking forward to explaining our fantastic project to everyone. See you at Boston!



Rafael R. Daga

Rafael R. Daga was born in Malaga (Spain). He studied Biology in the Faculty of Science, University of Malaga and graduated on 1991. He received his Ph.D. in 1999 in Genetics in the same University. He did his postdoctoral works in the CSIC/University of Salamanca (Spain), Columbia University and Rockefeller University (New York, EEUU). In 2006 he

joined the University Pablo de Olavide in Seville (Spain) where, since 2009, he is full professor. Since 2008 he has his own laboratory in the Andalusian Centre for Developmental Biology (CABD) in Seville. In 2010 he participated as instructor in iGEM with the UPO-Sevilla team and presented the project entitled Bacterial Crowding.



Gabriel Ruiz

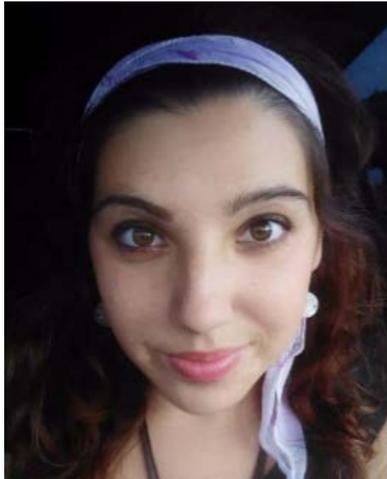
Hi! I am Gabriel Ruiz. I am a 21 years old student of Biotechnology at Pablo de Olavide University. When our teacher told us about making an iGEM team, I would have never thought I would be selected to be part of it! I'm so grateful and excited, as synthetic biology is one of my two passions, apart from cinema. I would love to direct a movie about iGEM!

Fernando Govantes

And this would be me... Over the years evolved to be called Fernan and then Fer... I may end up losing my name altogether. I am the grandfather of the team and I run the Biofilms subteam. Best known for my secular knowledge on bacterial strain genotypes, transposons and other genetic gimmicks, and for being the keeper of legendary protocols and media recipes.



Communication



Maite González

My name is Maite but everyone calls me May. I am studying Journalism at the Sevilla university since 2013. I just finished my third year of my career and I am very passionate about it. My hobbies are the photography and audiovisual creations like short-films and spot. I really love the literature, the videogames and the Science fiction too. I am interested in the popularizing of science so I think it is something very important in the last years to communicate new scientific advances. I consider myself a creative, fun and very sociable person so I am always delighted to participate in new projects. Nice to meet you.

Informatics



Israel León (Advisor)

Hi! I am Israel León, a student of the Information Systems Engineer Degree in University Pablo de Olavide. I don't quite understand what biotechnology is, but I love this project and its complexity. I hope you find it as funny as I do!

Project

Bacteria are an amazing field to work on. They can develop a wide variety of functionalities and properties in a very short time, being easy to work with and extremely versatile. On this basis, we wanted to take advantage of one of the most remarkable characteristics of the bacteria we work with, *Pseudomonas putida* – the ability to form biofilms. These structures are composed of communities of bacteria which are integrated in an extracellular polysaccharide matrix that protects them from several kinds of stress. In addition, they have an increased metabolism, a feature that is often required in industrial production.

Therefore, we thought that it would be a good application to try and do some bioremediation with these strong and

powerful bacteria. First, we thought about glyphosate, an herbicide that has been proved to be toxic to human. But due to technical difficulties, we could not continue with that idea. Instead, we found out about glycerol. It is being overproduced in the biofuel industry, and it is starting to become an environmental problem. So we started to model how our bacteria would eat that glycerol, using biofilms for that. But, is there anything else we can do once the bacteria are eating glycerol? We decided that there could be a compound we could produce with these biological reactors, and our choice was propionate. It is widely used in a large variety of fields, easy to excrete from our bacteria and, according to the model, easy to produce. Let's get started!

Human Practice

Through the human practices we want to show people that we have an important project that can improve problems in the environmental field, and produce some advances in industrial production. We are aware that those goals are not enough. Therefore, if we want to ours become a great project, we have to believe in our team and effort. We believe that science should be closer and more interactive and for this we have participated in science fairs and exhibited in several schools. Interacting with young students has been a unique experience for us, and has made us better communicators and better people. In science the only goal should be to make the world a better place. This is the compromise that all of Spain iGEM teams have decided to become real, and so we work together for the same purpose.

Science Week, Universidad Pablo de Olavide

This was our first contact with teenagers that wanted to learn a bit more of "real science". It was funny to explain them what synthetic biology was, and how did it work. They seemed fascinated about this, and they were bewildered when we explained them how our project was going to develop. We also showed them a software developed by a team from Paris, in which they had to read some text and give answers. They all told us that it has been a good experience, so we were extremely happy!



Lab practices with students from IES Martínez Montañés

At the beginning of the summer, some students of this high school came to the university to do some practices with us. We prepared a practices lab and we show them how to do an electrophoresis gel or a digestion. They were all fascinated to be able to perform some real experiments, that otherwise they had only seen in text books. Some of them



Science fair

The 14th Science Fair celebrated at the Exhibition and Conference palace at Seville on 5, 6 and 7 May 2016. This year the theme was climate change and environmental pollution. Because of that we decided that our project fit perfectly in this edition, and had our own stand in which we could talk about the project. During the fair we made contact with people from very different fields, from teachers of different Spanish and foreign universities to children of primary school. It was a

decided that wanted to study a biotechnology or biology degree, what made us feel very proud. During those days, we also learned lots of things, as it usually happens when you are the one that has to give the information to others. In summary, it was instructive and fun!





great experience to show our project, and to answer the questions of all the people that wanted to ask. For children we were able to turn our project into an interactive game, and for young students our idea was inspiring for their future, as themselves told us. As we were also explaining how synthetic biology works, people instructed in some technical careers could also understand us.



On August 18 we traveled to a meeting with other Spanish iGEM teams to the city of Valencia. Throughout the weekend the 3 groups (UPO-Sevilla, Barcelona and Valencia) discussed constructively on different projects and applications and contributions to science. It was a wonderful way to collaborate and help each other. All projects were very applauded and congratulated by the others. We are really delighted and integrated when interacting with the different disciplines that make up the total of equipment, from biotechnologists and biomedical or computer, to a journalist. Thanks to this meeting different teams have created good friends who made us now a great family that helps each other.



different Edmonton high schools

URBANTUNDRA_EDMONTON



UrbanTundra Edmonton - IGEM HS 2016



urbantundraedmonton@gmail.com

Our team, Urban Tundra Edmonton, is composed of twenty-one students from four different Edmonton high schools. The majority of us are from the graduating class of 2016, attending the University of Alberta this fall. Nonetheless, each and every one of us shares a keen interest in the STEM fields- science, technology, engineering, and math.

Team

Our team, Urban Tundra Edmonton, is composed of twenty-one students from four different Edmonton high schools. The majority of us are from the graduating class of 2016, attending the University of Alberta this fall. Nonetheless, each and every one of us shares a keen interest in the STEM fields- science, technology, engineering, and math. Our primary investigator is Professor Mike Ellison (Ph.D., Dept. of Biochemistry at the University of Alberta). This year, we will be presenting our project at two sites: aGEM in Calgary mid-September, and iGEM in Boston late October.



Project

Our project came together from learning about iGEM through the “Fusion” Science Expo last fall; and the theory behind the movie, “The Martian”. With these two components in mind, Urban Tundra Edmonton has committed the bulk of our summer to lab work.

The goal this year aims to colonize Mars by creating oxygen through the metabolic pathways of e. Coli; as well as provide rocket fuel. Martian soil has been reported to have a toxic perchlorate (ClO_4^-) concentration of 1%. However, ClO_4^- can be broken down into oxygen (O_2). Using this information, our team is genetically engineering e. Coli DNA with the enzymes perchlorate reductase and chlorite dismutase. This allows for the bacteria to biodegrade ClO_4^- firstly into ClO_2^- ; and later biodegrade ClO_2^- into Cl^- and breathable O_2 . As for rocket fuel, when perchlorate is reacted with ammonia, it produces ammonium perchlorate. With these two essential elements- oxygen and rocket fuel energy- human life and activity can be sustained on the Red Planet.

Collaboration

Searching for new collaborations

If you want to find ideal partners, do not hesitate to tell everyone!

We would love to collaborate with other iGEM teams on a local and global scale. Yielding the technology we have today, our most efficient way of collaboration is through face-to-face video conferences. By getting to know other iGEM teams, we not only have an insight of each other’s projects- but each other’s ways of life outside the lab. A video call with Urban Tundra Edmonton generally entails getting to know project details, protocols, and backstories, as well as cultural customs and personal quirks.

Human Practice

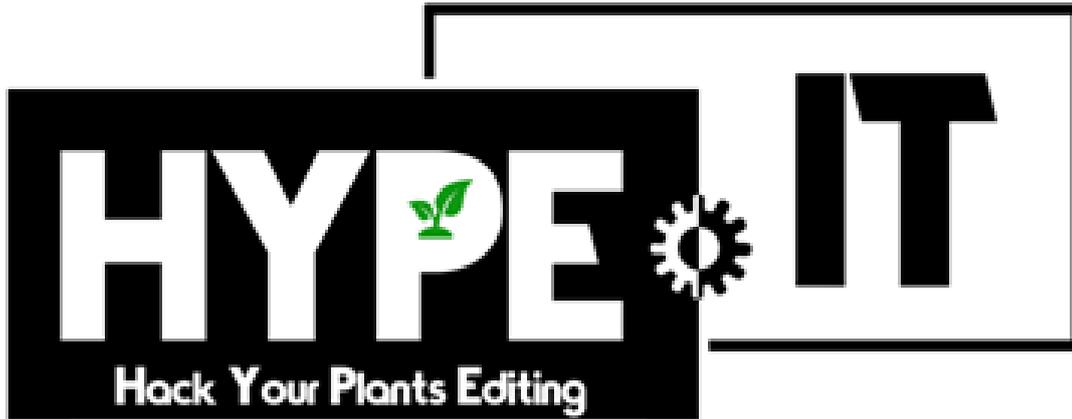
Our team takes special precaution when working at the University. All of our team members diligently wear lab coats, latex gloves and closed toed shoes when working in the lab.

Discussion

Regarding the ethics and safety of our work, remediation of Martian soil does not directly violate human nor animal life. However, the mission to provide sustainability of life on Mars brings up controversial discussion as to whether it is ethical to create a manmade habitat on a different planet. Due to the exponential growth of the human population, Earth and its resources are becoming less sustainable. In order to protect life as it is, we must think outside the box quite literally and figure out how to sustain fundamental elements, starting with oxygen.

the Polytechnic University of Valencia

VALENCIA_UPV



@UPVigem



Valencia UPV iGEM



Valencia.upv.igem@gmail.com

Valencia UPV team has been participating in iGEM the last ten years. In our team it has always been valuable the multidisciplinary, so it is composed by biotechnologists, industrial, electronics and biomedical engineers and computer scientists from the Polytechnic University of Valencia (UPV), Spain.

Team

Valencia UPV team has been participating in iGEM the last ten years. In our team it has always been valuable the multidisciplinary, so it is composed by biotechnologists, industrial, electronics and biomedical engineers and computer scientists from the Polytechnic University of Valencia (UPV), Spain. The team members are all undergraduate, between 19 and 23 years old, including iGEMers from iGEM 2015 that advise and support the new members.



Project

Our project is supported by IBMCP-CSIC (Plant Molecular and Cellular Biology Institute-Superior Council of Scientific Research). The development of our project was mostly inspired by the innovative genome editing technique CRISPR/Cas9 and our wish to apply it as a new plant breeding technique. The obtaining of new crop varieties usually takes a long time and a high economical investment. Additionally, the mutations that characterize the phenotype of the new variety are not known, except if it was obtained through transgenesis. However, transgenics are not socially accepted and in some regions like Europe they are practically forbidden.

We found that the solution to these problems was to improve the accessibility of plant genome editing with CRISPR/Cas9. We aim to create a system that allows local plant breeders to obtain their new varieties in a simplest and fastest way than the current techniques. This includes making the information needed to work with CRISPR/Cas9 easier to use and to understand.

Our project is based in making knockouts in the selected genes to obtain new plant varieties. The

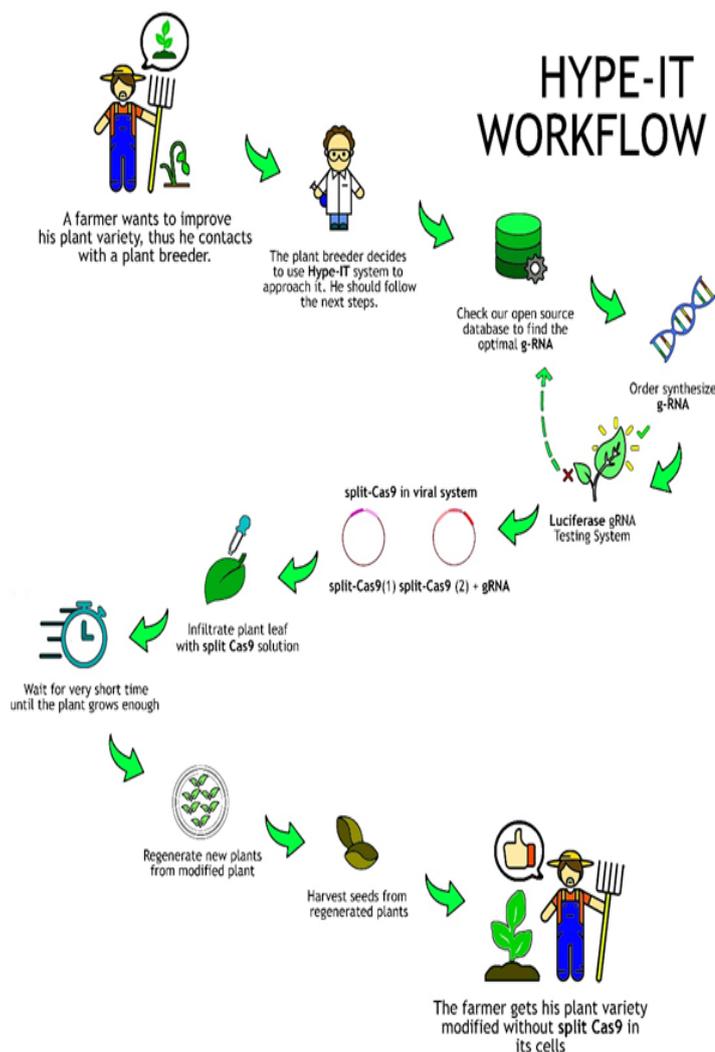
knockouts are possible by using CRISPR/Cas9 and the necessary gRNA, which are inserted in the plant with viral vectors. But, how do we make the process accessible to local plant breeders? We propose four methods to achieve this:

1)Data processing software: we firstly will create a database which connects desired plant traits with the gene that is necessary to knockout to obtain the trait. This database will include genes found in bibliography and predicted genes that can be knocked-out in other plants to obtain possibly the same phenotype. Using this database, the data processing software will be able to obtain the optimal guide RNA to knock-out the gene that the user selected.

2)gRNA testing system: the database provides gene consensus sequences. However, plant breeders might use a plant variety different from the one which is sequenced. Given that, it is necessary to test the gRNA for the specific variety that they want to improve. Our modular and standard testing system will inform the user if the gRNA works on his plant and which efficiency does it have compared to other gRNA. The reporter system is based on luciferase, so the breeder will

obtain luminescent signal if the gRNA works and hence the CRISPR has cut.

3) Split Cas9: one of the keys of our project is the way to make more efficient and faster the obtaining of the new variety. The traditional technique for plant genome editing uses *Agrobacterium* as vector to insert the CRISPR/Cas9 in the plant. This makes the plant transgenic and has low efficiency, given that a low proportion of the plant cells get infected with *Agrobacterium*, lacking the Cas9 and the gRNA, making impossible the editing. The alternative is using viral vectors, which have higher infection rate and don't insert transgenes in the plant. However, the insert size admitted by viral vectors is



2.2kb. Cas9 measures around 4kb. For that reason, we will divide Cas9 in two, insert each part in a different viral vector and once inside the cell, both subunits of Cas9 will bind through inteins.

4) Lab-case: finally, to be sure that actually anyone who wants can have access to this editing technique, we will design and build low-cost laboratory equipment specific to make plant genome editing with CRISPR/Cas9. This will include centrifuge, electroporator, thermal cycler, luminometer and electrophoresis module. The cost could be around 500\$ for all this equipment, considerably cheaper than the traditional equipment.

With these strategies we expect to improve accessibility to plant genome editing with CRISPR/Cas9, allowing the global enhancement of agricultural economy, from the small local plant breeders to the biggest companies. In the human practice section, we discuss why is this important and how the problem of food distribution and quality should be solved.

Collaboration

Our team is one of the few working with plants as chassis. We always wish to make contact with other plant teams to exchange experiences and ideas. We would like to make a call to these teams, so we can establish a solid collaboration that will benefit both parts.

The objective of our collaboration is firstly to test our gRNA testing system with other teams. We have designed it in a way that it can be standard for plants, and the only way to be sure about this is that other teams around the world try it in their labs.

We are open to help other teams in anything they need, from testing their models to try their parts in a different chassis. If some team is working improving the accessibility of synthetic biology technologies, we could make together a solid study for human practices, taking in consideration the general variables that affect this issue.

Human Practice

One of our basic needs as living beings is nutrition. However, it is a luxury good in many areas around the world. Countries in development are regions in need of basic food resources because of its lack of access to developed technologies.

Nevertheless, people in countries that are classified as first world suffer similar situations every day. World population is going to increase exponentially the next years, but natural resources won't be enough for provide minimal calorie

necessities to each person. For that reason, providing staple food to everyone has become one of the main social problems to solve in this century.

There are several problems related with this issue. First, food is lopsided distributed. There are areas without enough provisions to supply necessary nutritional intake to less than half of its population while others waste several tons of edible food. Also, available food often does not provide the correct amount and proportion of nutrients. Second, there are several problems with food expiration date. For example, fast ripening after fruit harvest avoids a correct distribution and remote areas are not able to consume them.

Scientific research has the possibility of solve these kind of social problems by managing their biological and engineering tools. However, different barriers limit the expansion of scientific advances and the possibilities of take advantage of them to solve basic social challenges. Some of those barriers are related to social opinion about new technological progresses, caused mainly by misinformation. Additionally, basic laboratory tools are too expensive for average users, so most of them usually struggle to obtain enough funds to start a seedbed or to buy new equipment.

Bearing that in mind, we want to develop our project, HYPE-IT, to knock down those barriers that avoid scientific advances to arrive to those who have actual problems. Farmers need new plant varieties to have more efficient crops, as well as products with enhanced properties. Consumers need new plant varieties that arrive to the market, with better flavor, less expensive and more durable. The world needs plant improvement to cope with the increasing demand of food amount and quality.

However, the obtaining of a new plant variety is expensive, and requires a long time, usually around 10 to 15 years. The population increases 1 billion each 12 years. The time to obtain any new variety is unacceptable if we take in account these figures. Our team aims to offer an affordable technology that allows plant breeders obtaining desired enhanced crops by blocking gene routes, making the editing faster and more efficient. New plant varieties could be obtained in 1 or 2 years. If every local plant breeder improved its local varieties, a new green revolution would help the world to overcome one of the biggest problems of the century.

Discussion

GMO, transgenics vs gene editing.

Nowadays, food and nutrition are considered as necessary goods. Food production should be increased in order to satisfy all the nutritional needs in developing countries. In order to improve the production of the basic crops, transgenics play an essential role.

Transgenic plants are obtained when DNA of a specific plant variety is genetically modified by introducing genes from another species, which confer new traits to the variety. These traits could have an important agronomic interest. Transgenic plants could have significant benefits. For example, herbicide resistant plants allow the farmer to improve yield production, avoiding the direct damage to its crop with the herbicide. Other example is modified plants that can be resistant to bacteria that introduce toxins into the genome of the plant, producing lethal effects. This modified plant includes a foreign gene that blocks the bacterial toxin.

However, the advantages of transgenics go beyond this. Plants can be modified to introduce nutritional benefits such as vitamins or hormones. With this application, nutritional deficiencies may extinguish in the whole world. Furthermore, therapeutic proteins produced in transgenic plants are used in many treatments. Antibodies against hepatitis B and vaccines against infectious diseases are examples of how transgenic plants are necessary.

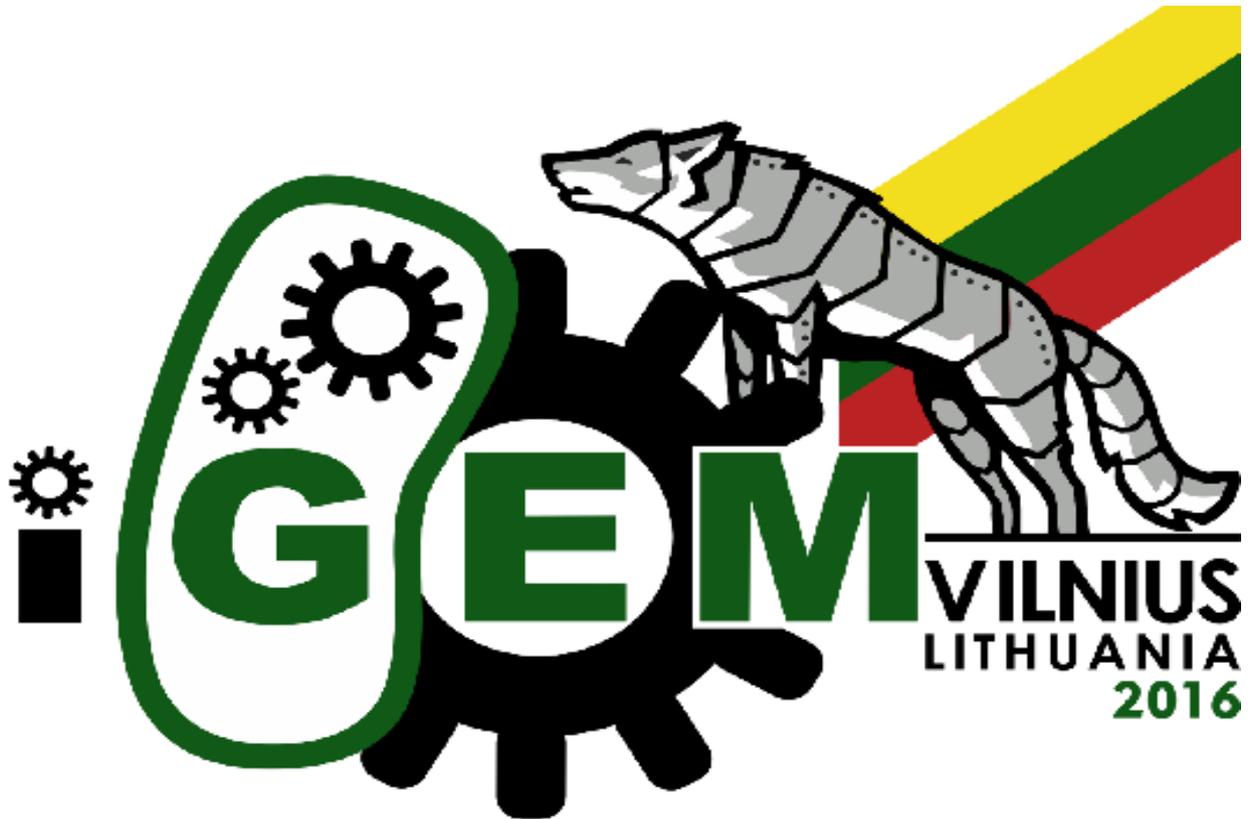
Nevertheless, some people think that transgenic crops could cause allergies. They also believe that they are no natural, and hence transgenic crops must be rejected. In general, people do not like eating food in which foreign DNA has been inserted. Moreover, people think that transgenic crops may damage the natural environment and themselves, so they prefer not to consume them.

Enhanced production of crops through transgenesis could end with all the nutritional and health problems around the world. However, public opinion plays an important role in the future of the transgenics and this is not very optimistic.

For this reason, in this project we consider that the genome editing technique CRISPR/Cas9 is an alternative to obtain new crop varieties. With this method, the mutations that allow the obtaining of new traits are known. These crops are more probably to be socially accepted because the genetic researchers do not “play God” crossing genes of different species. Moreover, there are not any risks to the environment or even to the consumers. It is the same plant variety with a specific modification in the genome which allows us to obtain a new and interesting phenotype. The same mutation could have been obtained through the more common and traditional technique of random mutagenesis. CRISPR/Cas9 technique is a faster and simpler method compared to random mutagenesis, and can be used in an easy way by anyone when comparing to transgenesis. We hope that more countries, and particularly the European Union, will recognize the benefits of gene editing and will approve its usage as plant breeding technique, not stopping this time the revolution that brings scientific innovation in the synthetic biology field.

Vilnius University

VILNIUS-LITHUANIA



@Vilnius_iGEM



VilniusiGEM



vu.igem@gmail.com

Our team consists of 18 students from different tracks of life science – biochemistry, genetics, molecular biology, medicine and bioengineering.

Team

Our team consists of 18 students from different tracks of life science – biochemistry, genetics, molecular biology, medicine and bioengineering. To make the teamwork more efficient, we have divided our team to several subgroups – PR group (6 people), finance and marketing group (5 people), research group (4 people), Interlab group (6 people) and human practices group (5 people); however, as it is seen by looking at the numbers, team members have overlapping responsibilities. Such partition enables more organized project development, hopefully leading to better performance at Giant Jamboree.

Project

The aim of Vilnius-Lithuania iGEM 2016 team is the treatment of a genetically inherited condition called phenylketonuria (PKU). The condition is defined by person's inability to metabolize an essential amino acid phenylalanine due to the mutation of phenylalanine hydroxylase (PAH) gene. As a result, the phenylalanine hydroxylase enzyme is not produced and phenylalanine derivatives accumulate in the brain of the affected person. This, in turn, causes different neurological symptoms ranging from depression to epilepsy and severe mental retardation. At the moment, there is no known cure to this condition; however, since phenylalanine is found in almost any protein-rich food, the only treatment is a low-protein diet. This diet excludes the most common foods consumed by the majority of the population – bread, meat and dairy products. Although the patients are being supplied with a special phenylalanine-free powdered food, the disease might have a negative effect on the quality of their lives leaving them with the struggles of food choice in the public catering.

Vilnius-Lithuania iGEM team came up with an idea of a possible treatment for phenylketonuria – a probiotic, which would absorb phenylalanine and metabolize it in the intestinal tract of the patient. To be more precise, the team has two approaches to this idea.

The primary approach is a phenylalanine ammonia lyase (PAL) producing bacteria – the PAL enzyme will break down phenylalanine. To fulfill the second approach, the team created a new synthetic gene, which consists of a large amount of phenylalanine codons. During the process of translation, the excess phenylalanine will be incorporated into the synthesized protein. Such protein is expected to form inclusion bodies inside the bacterial cells. Lastly, since the constitutive synthesis of phenylalanine-rich protein is highly disadvantageous and possibly even harmful to bacteria, the team is thinking of developing a riboswitch-based posttranscriptional gene expression regulation. The riboswitch will consist of a phenylalanine aptamer and a ribozyme, providing the translation only in higher phenylalanine concentrations in the environment.

Collaboration

We would gladly appreciate any help regarding probiotics and metabolic diseases whilst we too are capable of providing any help in this field.

Also, since our project includes the creation of an artificial aptamer, we are seeking help in computer modeling of the aptamer. We have found software written by Heidelberg iGEM 2015 team (MAWS and JAWS); however we did not succeed in launching it.

Since our life sciences research centre brings together specialists from various fields, we can be of any use to other iGEM teams. What is more, we have a strong CRISPR/Cas research background, especially with prof. dr. Virginijus Siksnys as our PI, so we can offer help to the teams working with this system. Do not hesitate to contact us if you want to discuss collaboration opportunities.

Human Practice

As a part of iGEM initiative, we are very dedicated to the outreach activities. One of the greatest achievements in this area is the foundation of Synthetic Biology Organization in Lithuania. The organization is aiming at spreading the knowledge of life sciences to the public. To this day we have visited a large number of schools where we introduced the possibilities of synthetic biology and gene engineering to the students and launched an art competition. Adding to that, this summer, a high school student has been practicing life science research in our laboratory.

Also, we have launched a cycle of lectures called Café Synthetique, during which we hosted different professors and specialists in the fields of our interest to discuss the most intriguing science-related topics of the society in one of the city's coffee shops. We have also appeared on the news pages and participated in several Facebook and Youtube live translations. In autumn, we are planning to launch even more outreach activities, such as bio-breakroom, bio-art exhibition and a quiz.

We are also keeping in touch with the Lithuanian PKU association – we have attended several meetings, interviewed the patients and participated in several workshops.

Discussion

Due to the fact that the project is aiming at producing a probiotic, it was very important to consider the opinion of the society of this kind of medication. The outreach activities showed that a lot of people lack basic knowledge of what probiotics are. Also, the fact that probiotic would be a genetically modified bacteria disseminated skepticism and insecurity in whether it would be safe to consume such product.

We had to do a lot of research on the available data on this subject to ensure that our engineered bacteria will not have any negative effect as on the patient, as on the environment. At the same time, we feel the urge to educate other people on this topic and we are fulfilling this desire during our outreach events.

University of Washington

WASHINGTON



UW IGEM Team



uwigem@uw.edu

The University of Washington iGEM team was founded in 2008, and is run by students from all over the world.

Team

The University of Washington iGEM team was founded in 2008, and is run by students from all over the world. With a diverse set of skills from bioengineers, computer scientists and aspiring entrepreneurs, our team has been able to tackle problems with biofuels, paper-based diagnostics and complex multi-enzyme pathways. This has proved especially helpful with this year's project which wouldn't be possible without a significant amount of drylab work. With an innovative approach, our research will serve as a building block in the expanding field of synthetic biology.



Project

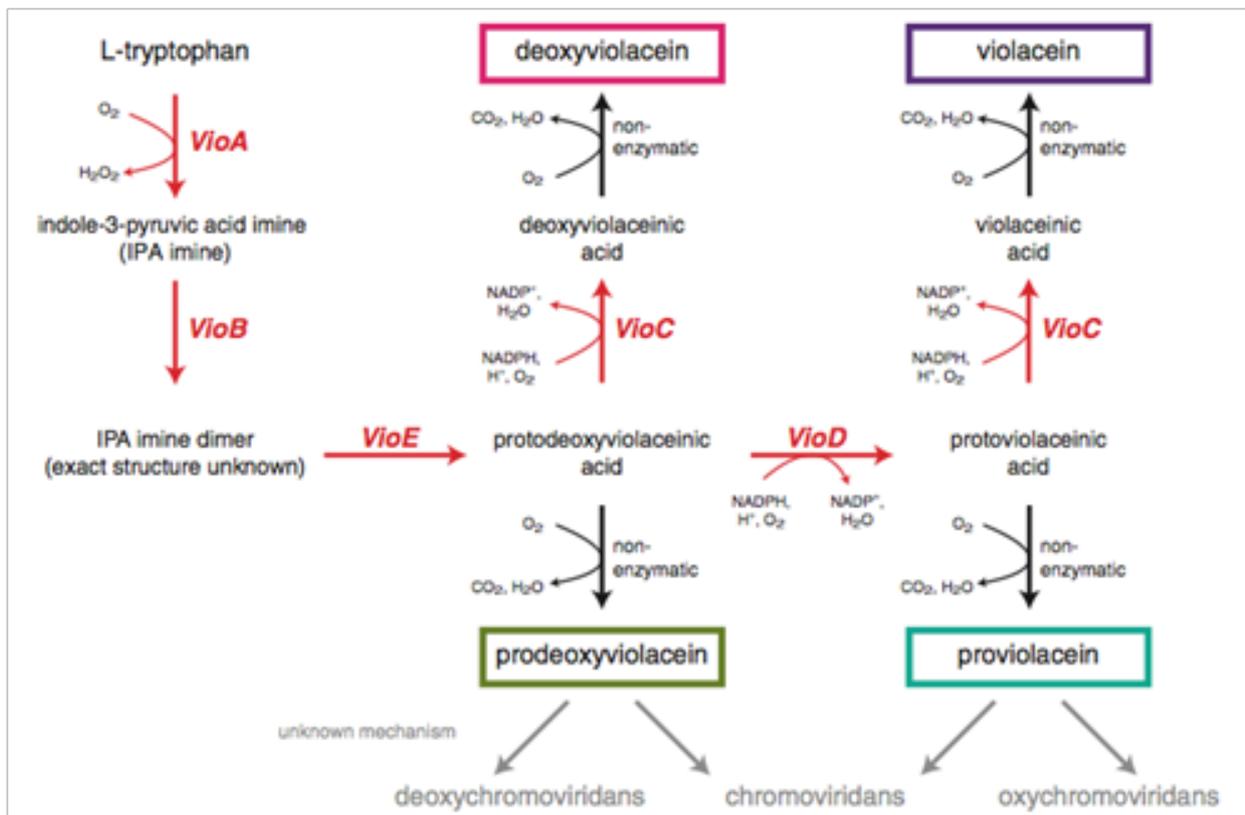
Viva la Violacein An Autonomous Control System for Yeast Cultures

Managing cultures is a vital task in synthetic biology, but constantly measuring and adjusting culture conditions is both tedious and labor intensive. Our project



aims to reduce the amount of time and effort needed to maintain cultures through the creation of an affordable image analysis system that reads visual data to measure the current state of a culture and then determines whether to release inducer chemicals based on user input.

Our project utilizes the violacein pathway to simulate other metabolic pathways with colored signals. By regulating gene expression in this gene set with two different inducible promoters, we are able to yield up to four different color outputs.



[4] The violacein biosynthetic pathway

These outputs are then measured by an open-sourced Raspberry Pi setup, which captures visual data via camera, measures the culture's RGB value, and then directs the gradual release of inducer chemicals to maintain or change the culture's color over time.

To control the violacein pathway in *S. cerevisiae* we are using the CUP1 inducible promoter, a concentration dependent yeast promoter that responds to CU_2+ ions, as well as the GAL1 yeast inducible promoter which responds to galactose. However, switching out the inducible promoters might lead to a number of new applications.

Human Practice

For years scientists have been observing the Violacein Pathway. Originating from the Amazon and marine protozoa, the pathway produces violet pigments (Violacein) in common bacteria. Today, this pathway is produced in laboratories to dye textiles and is studied for its medicinal properties. Violacein is unique as it is both antibacterial and tumoricidal, and thus at the frontier of cancer research. Despite numerous studies, little is known about the capabilities of Violacein and its full effect on protozoa and eukaryotic cells.

Our diagnostic culture imaging system will increase efficiency of those studying the Violacein pathway and other pathways with colored signals. An analysis of the RGB value will improve accuracy and precision in measuring the amount of Violacein, Prodeoxyviolacein, Proviolacein and Deoxyviolacein present in a culture. Our project will also add to the current information on violacein and eukaryotic organisms as we are using yeast cultures.

L-tryptophan, the first step in the Violacein pathway, is an amino acid which increases serotonin in the brain and also functions as an antidepressant. High doses can lead to chronic muscle pain and cognitive deficiencies. Violacein's effect in eukaryotic cells on the molecular level is also still unknown. Therefore, additional testing should be conducted before humans receive high exposure. It is also imperative to wear personal protective barriers between the cultures and human skin.

What is the effect of the release of this waste into systems?

A major concern is the waste that is generated from an increased production in Violacein. Fifteen percent of healthcare waste produced annually is biohazardous waste. Like most non-sharp biohazardous waste, the end result of the Violacein Pathway will be incinerated or autoclaved.

While the waste produced from the cultures will be a fraction of the global output, it is necessary to preserve limited resources. By measuring cultures with our diagnostic technology, scientists will incur fewer errors from solely



human observation, resulting in the production of less biohazardous waste. Often in the lab trials are repeated due to calculation errors or imprecise readings. Our computer simulation will also improve accuracy and reduce the time in the lab analyzing the cultures or repeating trials. With the information stored online, sharing data will be more feasible and thus promote a global education and awareness of the Violacein Pathway.

Further questions to explore:

Is this commercially feasible?

This system will be able to be downloaded onto computers for easy access. Licenses have the potential to be sold online or instore. By selling our product online, we would increase the overall net profit as less money is spent on the assets required to manufacture the software.

What is the effect of an increased electrical output? Can the screen be dimmed when not in use?

Laboratories consume on average between thirty and one hundred kilowatt hours of electricity per square foot. However, if our system averages 100 kW of power for one hour, it will only consume 0.1 kWh per day. Taking into consideration the other devices used in synthetic biology labs, our diagnostic technology is an energy efficient option. It is also important to consider the amount of resources used to create the cultures, such as electricity for the laboratory, microwave, and other devices necessary to produce L-tryptophan. If our system reduces the need for repeated trials, our system is the most cost effective and eco-friendly approach.

How much would it cost to produce?

Not only is our system orders of magnitude cheaper than most commercial bioreactors, taking into account it may cost \$0.10 per kWh, it would cost the laboratory \$3.65 annually. Considering most lab equipment and electrical bills, this is not only energy efficient but a cost effective solution to repeating trials.

Would this take the position of workers in the lab?

This would not affect the labor required in the lab, but would potentially save time for each worker. Fewer trials would be repeated, but not to the extent that someone would be laid-off. This is assuming that the initial analysis of cultures is mostly accurate.



Discussion

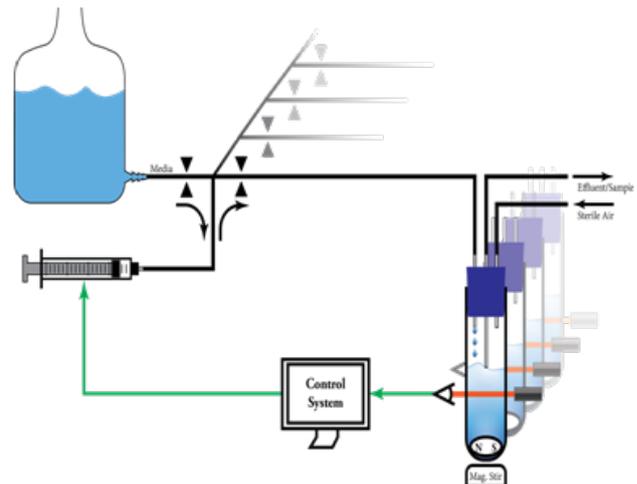
Education:

Our iGem team not only focuses on the work of students at the University of Washington, but we aim to inspire the next generation of bioengineers through community outreach. The team has traveled to venues across the Puget Sound region to share our love of research and synthetic biology with kids K-12. Interactive demos such as a strawberry DNA extraction [left] provide kids and parents an opportunity to ask questions about biology and our team's research.

Our booth at Engineering Discovery Days at the University of Washington

Want to extract Strawberry DNA in your community?

Search this link: <http://uwigem.org/post/127224577381/cool-outreach-ideas-for-igem>



[8] Basic design and operation of the Klavin's

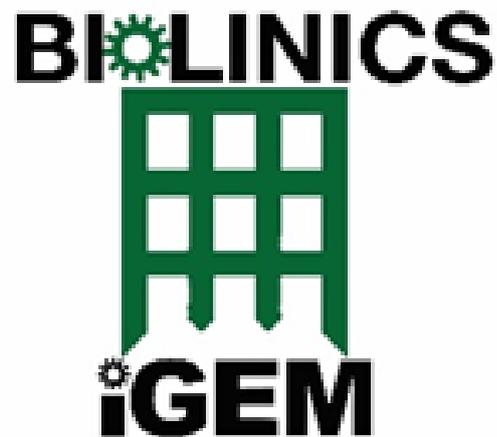
The iGEM registry is evidence of the open-source nature of iGEM. However, there are many other ways iGEM teams are contributing and benefiting from open-source science.

In our project we benefited from open-source access to the design of a mixture controlled turbidostat created by the Klavins lab at the University of Washington. We used the turbidostat's wiki to help build our image analysis system, a significant part of our project. This is an example of how important access is to our team and to other iGEM teams. We plan to use the wiki to share our image analysis system and research for others to use and benefit from like we did.



the University of Westminster

WESTMINSTER_UOW



@WestminsteriGEM



westminsterigem2016



uow.igem@gmail.com

We are the University of Westminster iGEM team: the Biolinics. "Bio" from biosynthesis and "Linics" from amino-levuLINIC acid.

Team

Who are we?

We are the University of Westminster iGEM team: the Biolinics. "Bio" from biosynthesis and "Linics" from amino-levuLINIC acid.

Our team is consisted of students from a range of different subjects making us a very multidisciplinary team, subjects include: Biochemistry, Microbiology, Molecular Biology and Genetics, Pharmacology and Physiology, Biomedical Science, Biotechnology, Computer Science and Contemporary Media Practice. Degrees of expertise vary between 1st, 2nd and final years (who have now graduated). We are very proud to have an international team as you can see in our team picture with the flags below.



If you are interested in knowing what else we are up to, follow us on Twitter and our Facebook page! Any questions about our project or interested in collaborating contact us via our email!

Looking forward to meeting you all in the Giant Jamboree and good luck with your projects!

Project

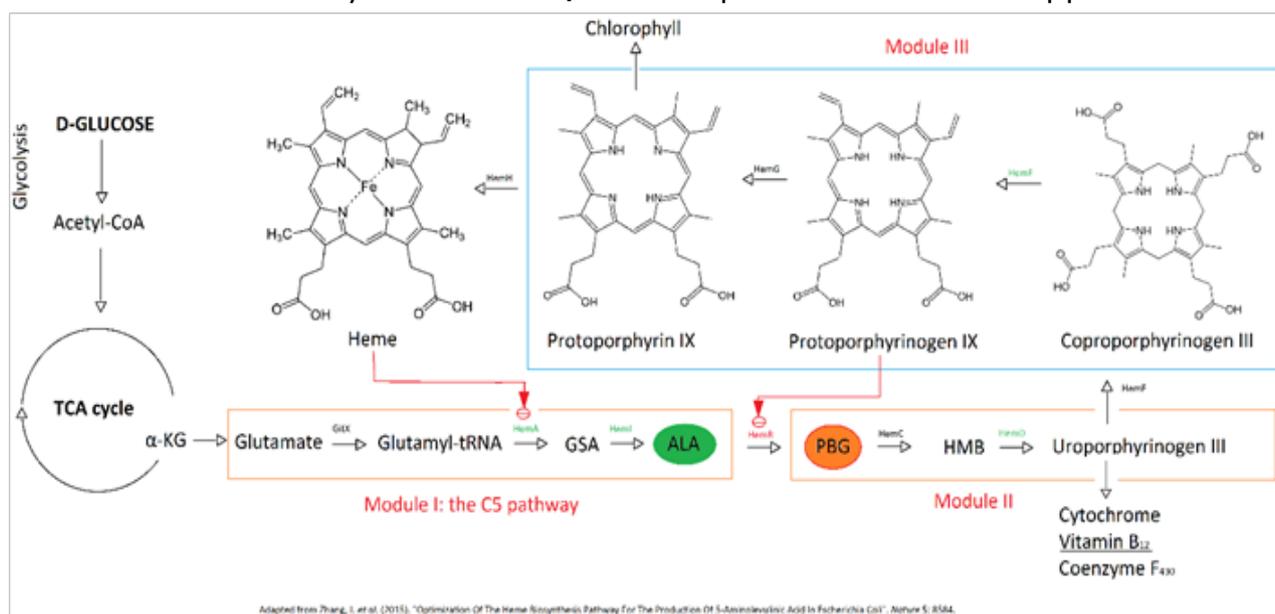
METABOLIC ENGINEERING OF E.COLI TO PRODUCE A NON-PROTEIN AMINO ACID WITH ANTI-CANCER AND BIOHERBICIDE PROPERTIES

We are metabolically engineering E. coli to biosynthesise amino-levulinic acid (ALA), a non-protein amino acid which is a key intermediate in the tetrapyrrole pathway – see figure below. This pathway is naturally occurring in bacteria, archae and plants. Since ALA is quickly used up, our main aim is to upregulate the genes involved with ALA production and adding a membrane transporter protein to pump ALA out of the cells. We, therefore, picked HemaA, L, D and F as the responsible genes for ALA accumulation and have built our constructs. After successfully transforming the E. coli cells with the new plasmids, we are going to do a series of fermentation reactions in order to produce ALA and scale up in various volumes to assess whether the strain produced has potential for industrial applications and use.

ALA is widely utilised in medical sciences and its uses include:

- Photodynamic therapy
- Drug delivery
- Bioherbicides, biopesticides and bioinsecticides
- Tumour localiser
- Drug delivery
- Plant growth hormones

Thus, this project is promising as it holds the potential to revolutionise the current chemical methods to synthesise ALA, in a cheaper and sustainable approach.



Human Practice

We have been quite busy with human practices aspect. We participated in the Science for 4 fun fair where 200 pupils attended from 13 schools across London. We have been conducting a survey on controversial topics in synthetic biology (you can fill in our survey here: https://docs.google.com/forms/d/e/1FAIpQLSdbf_r-n4xl7OoAedAP8aVTaa1vb-mvOyzKIkRgc7VelglqEw/viewform). We have also been designing an app: Plasmid World, which educates young students on how to build a plasmid in an interactive and fun way. We will test our app in schools where they will give us feedback. Plasmid World will be available on Apple Store.

We were also delighted to host the UK meet up for two years in a row.

The #iGEMUKmeetup2016 consisted of a 2 day conference-like event; where there were talks delivered by academics: Professor Jane Lewis (University of Westminster), Dr Anatoliy Markiv (University of Westminster), Dr Robert Smiths (King's College London), Dr Tom Ellis (Imperial College London) and Dr

Vitor Pinheiro (UCL/Birkbeck). 100 students from 18 UK universities attended this event with the aim to practice their presentation skills before going to the Giant Jamboree (Boston, USA) in October. Each team had 15 minutes to present their project and their report their current progress, each team also had an extra 5 minutes for Q&A. During the Q&A section, other teams raised interesting points about their project, giving constructive feedback and suggesting new ideas – making room for the teams to improve. The teams were provided with tea, coffee, biscuits, sandwiches, crisps, pizza and soft drinks throughout the conference with no cost. Furthermore, the event also offered a pub crawl around London's iconic places as a social event where the teams could socialise.

We conducted a survey after the event where 51 people answered. We got back very positive feedback. For instance, all the 51 people liked the event and agreed that it was a good platform to meet other teams and socialise, as well as to discuss projects and future collaborations. We also asked them to describe the

Xiamen University

XMU-CHINA



@Amoy_igem



IGEM XMU-China



igemxmu@gmail.com

Our team, XMU-China, consists of 18 undergraduates. We major in different disciplines, including chemistry, chemical engineering, material, life science, pharmacy, energy and public health.

Team



Our team, XMU-China, consists of 18 undergraduates. We major in different disciplines, including chemistry, chemical engineering, material, life science, pharmacy, energy and public health. This year, we are making effort to deal with some realistic problems like antibiotic resistance, which will be talked later. Besides experiment, we also designed various activities for human practice. Newsletter is a part of it and we'll show you more details in the next part.

Project

Background

Antibiotics made a vast medical advancements over the past 70 years. The annual

annual consuming rate of antibiotics appears to be rising. It is reported that in 2010, 70 billion individual doses of top seven antibiotic classes were consumed, which equates to about 10 pills, capsules, or teaspoons for everyone on earth .

People and animals get antibiotics and develop drug resistant bacteria. When the antibiotics kill pathogenic bacteria, some of them mutating and developing the "drug-resistance". Drug resistant bacteria flourish in the absence of diminished competition, and the resistance can be transferred among the bacteria, which made their infections considerable mortality and morbidity . Furthermore, the prevalence of the infections is still increasing by abused antibiotics and resistance spreading. It is predicted that 10 million people would be killed for the drug-resistant pathogen infections each year by 2050 . Multidrug-resistant (MDR) pathogens has been identified as one of the top three threats to human health by the World Health Organization (WHO).

Bacteria develop clinically significant resistance in a period of just months to years. There have been several common drug resistant microorganisms such as methicillin-resistant *Staphylococcus aureus* (MRSA) , vancomycin-resistant *Enterococcus*, fluoroquinolone-resistant *Escherichia coli* and multidrug-resistant tuberculosis mycobacteria. The new antimicrobials or modifications of current arsenal may not address the trends in resistance effectively . Combination therapies for the treatment are needed such as quorum sensing inhibitors, bio surfactants, bacteriophage, enzymes, etc. .

Ideas

We aimed to combat the drug resistance bacteria based on the engineering of gene regulatory networks. During our brain storm, the gene circuit "plug and play" designed by Kobayashi et al. helped us to find the design strategy for constructing the programmable bacteria. The circuits of detecting and killing MRSA were from iGEM Team TU-Delft 2013 and LMU 2014. And we modified the Synthetically Synchronized Lysis Circuit (SLC) design by M. Omar Din et al. for our gram-negative bacteria detecting and self-lysing synchronously.

The programmable bacteria would be comprised of three modules.

A. Biosensor Module

- A1. Gram-positive transgenic sensing
- A2. Gram-negative transgenic sensing

B. Switch Module

B1. Self-destruction switch

B2. Toggle switches

C. Killer Module

C1. SiRNA for knocking down gene expression

C2. Toxins

Limitation

For the time limitation, we didn't test every combination of the modules. In the future work, more combinations need to be tested and more gene regulatory modules need to be constructed to enhance the capabilities of modular genetic control circuits. In this study, we tried to protect the natural host microbiome through the self-destruction switch. However, biologics have pitfalls such as *in vitro* stability, restricted delivery options and limited high-throughput, etc. It is a challenge for us to overcome.

Collaboration

***Need your help**

We need this enzyme which can destroy the biofilm. Previous study showed that the combination of triclosan and DspB effectively inhibited the formation of biofilms on both the internal and external surfaces of urinary catheters . BBa_K1659200 may be a useful part but we can not get it. If you have any advice, please help us.

An interesting web game regarding to reducing antibiotics abuse is requested to educate children. We would like to upload it to a public gaming site. If you can make it, please contact us.

*What we can do

We are making the Newsletter on which different teams can exchange their ideas and introduce their project. We issued 7 edition last year and it did help many teams with their projects. Would you like to join in this year? Please give us your email address if you are interested in it, and then we will send you an email for more details. Our email address is igemxmu@gmail.com and please notice that the deadline is 21st August.

We have skillful and kind team members who come from College of Chemistry and Chemical engineering, College of Materials, School of public health, School of life sciences, Medical College, College of energy and School of pharmaceutical Sciences. Any collaborations such as software testing and survey translating are welcome and do not hesitate to contact us.

Human Practice

We conducted online and offline bilingual surveys regarding to the understanding and usage of antibiotics. In order to get a deep understanding of people's thoughts, we did some interviews and during the interview, we showed them the appropriate usage of antibiotics. Moreover, through interviews with experts, we got a lot of expertise. We went to undeveloped rural areas to distribute leaflets about detriments of antibiotics abuse. Taking the opportunity of the start of the a new term, we showed the freshmen brochures concerning synthetic biology, our project and iGEM competition. The opinions of project-related issues on ethics, social justice and intellectual property were also spread to them. We visited primary schools and taught

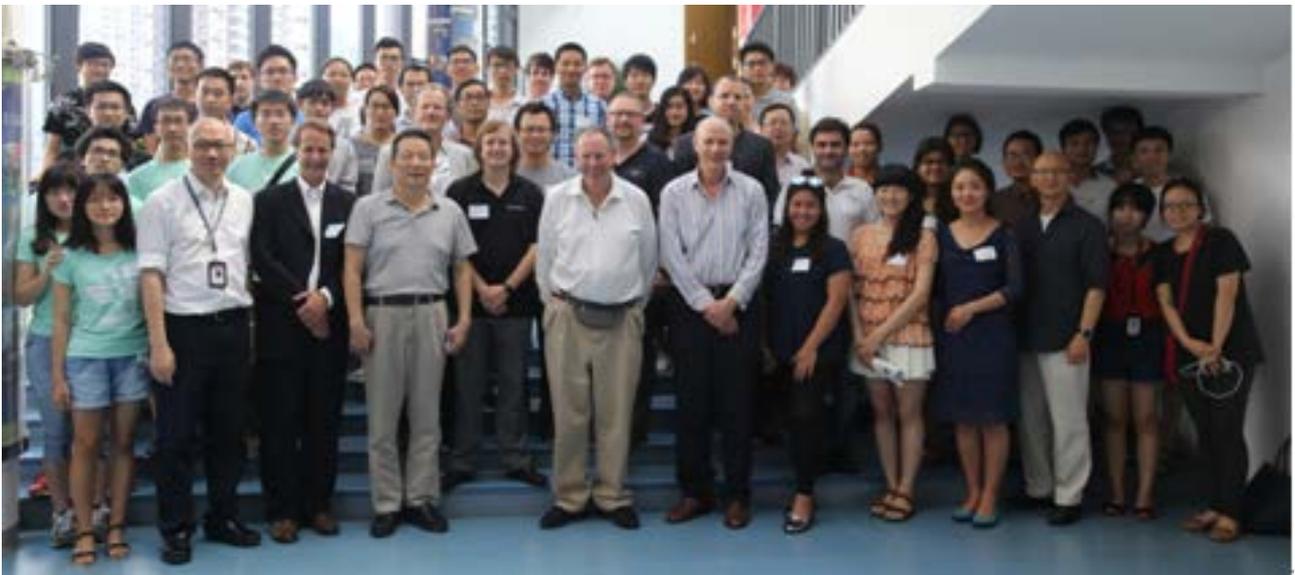
children the proper usage of antibiotics in vivid methods.

We did many efficient communications with other teams. Not only on the Internet but also by holding conference in universities. Some of us attended the Synbiobeta conference and learned



a lot of knowledge about synthetic biology.

Newsletter which is an electronic journal as an information exchange platform is a shining point. We have founded it since 2014 and it do help many teams with collaboration.



Discussion

Safety and security

In 1928, Alexander Fleming found the first antibiotic, penicillin, which led us to a new epoch of fighting against bacteria. However, the antibiotics bring great convenience as well as antibiotic-resistance to us. Sensitive strains which are killed because of long-term using of antibiotics are replaced by drug-resistance bacteria which survive. According to the First global report on antimicrobial resistance released by the WHO (1), antibiotic resistance is not prediction any more and we have entered post-antibiotic era. Antibiotic resistance has affected our daily life greatly.

The generation of antibiotic resistance is the inevitable consequence of natural evolution, while the human intervention accelerates the pace of this process. In terms of animal husbandry and aquaculture, a serial problems, including the low education level of the practitioner, cause the overuse of antibiotics. Beta-lactams, aminoglycosides, macrolides and lincosamides are major antibiotics used. What is more, the antibiotic resistance can spread with the market circulation, and finally harms the stability of this industry. If the hidden danger once breaks out, it will be a humorous disaster.

The National report on bacterial drug resistance surveillance in 2015(2) showed that 2,400,786 drug-resistance bacteria were detected in 1,143 hospitals. The detection rate of Methicillin-resistant Staphylococcus aureus(MRSA) whose fatality rate is 64 % higher than normal Staphylococcus aureus, has reached 35.8 % . The drug resistance will lead to a higher fatality rate of pathogens, threatening the cure rate of patients. The danger of some normal disease could increase highly, which could also make some clinical treatment invalid. At the same time, the increasement of medical industry risk could be against with medical industry and social stability.

Ethics

Some kind of medical injuries are unavoidable, but the intentional injuries are illegal and violate the ethical principles. The ethical principle "No Harm" means that the physician should not intentionally harm the patient, the patient should not deliberately injure himself either. Both of them would be violated by the overuse of antibiotics. In medical treatments about antibiotics, physicians are obliged to act more logically and perform the right of intervening in the treatments when

necessary.

The lack of internationally accepted principles of antibiotics usage in the food industry lead to a inadequate supervision system in this field. WHO, FAO and OIE should established a food processing monitoring platform.

Human society and bacteria world are in balance. The dosage and usage of antibiotics regulate the balance to some extent. If antibiotics are used properly, disease infected by bacteria can be controlled by mankind. So antibiotics have the satisfactory effect. But the rate of bacteria resistance would increase because of the overuse of antibiotics, which lead to the fact that scientific development can not keep up with the pace of bacteria resistance. The side effects of antibiotics are reflected by degrees.

Modern technology and industrial rationality are creating a growing gap between human and nature, and shadowing the symbiotic relationship between the two. People have the moral obligation for nature. Abusing of antibiotics is something greedy and out of control. It broke the natural balance, and under the natural forces, antibiotic resistant bacteria were created. The fact is that people will naturally be put on a firmer shackles for their greedy. If we want to meet demand, obey the rule of nature first.

Social justice

Although the academia is appealing to the restricted using of antibiotics, many developing countries do not have valid control measures. This phenomenon is not only related to the economics, but also a representation of some social problems.

According to reports in the People's Daily Online, the amount of antibiotics used in Chinese Mainland in 2003 had reached to 162,000 tons, which was about half of the amount in the world. In addition, more than 50,000 tons were discharged into soil and water. Although the Chinese Government has made laws to limit the use of antibiotics, the problem persists. In most hospitals, the use of antibiotics has been controlled suitably, while in clinics and meadows, supervision practices are insufficient. Why do some medical institutions overly rely on antibiotics? There're three main reasons: high efficiency, the lack of cognition of antibiotics and the high profit of selling antibiotics. The rate of abuse depends on the economic condition, the gap between the wealthy and poor, the educational level and other social factors, which could not only be controlled by the policy. To solve this problem, we need to pay more attention to the social justice.

Sustainability

The bacterial disease needs some sustainable treatments. As an emerging research

field, synthetic biology uses physics, chemistry, computer science, math, etc. to design and redesign the biological systems. Maybe synthetic biology can bring a new and practicable idea to a sustainable treatment. The traditional bacteria detection method may take 2 or 3 days to get the test result. During the waiting time, the doctors may use broad spectrum antibiotics to treat the patient. Using Synthetic biology methods, we can explore and design a new circuit which can detect the pathogenic bacteria quickly so that the more specific medicine can be timely used. In addition, new engineering bacteria which can degrade the antibiotics in the river or sewage can reduce the environmental pollution. What's more, the engineering bacterium killers may kill the Pathogenic bacteria instead of antibiotics.

Synthetic biology shows great potential on the sustainable development of the antibiotics treatment. But Synthetic biology is in the initial stage and needs more scientists to promote it.

Intellectual property rights

With the development of the society, intellectual wealth become more and more important and intellectual property which is protected by law gets more attention. In our project, bacteriophage which infects the Escherichia coli. specifically and non-lethally is planned to be used. We find that the resource of phage is all mentioned in the wiki in other iGEM team's project. In the academic world, everyone takes intellectual property seriously. However, there are some infringements which indicate that the law needs to be improved and public awareness of intellectual property rights needs to increase.

[1] Nischal PM. First global report on antimicrobial resistance released by the WHO. National Medical Journal of India. 2014;27(4):241.

[2] National report on bacterial drug resistance surveillance in 2015. China Licensed Pharmacist. 2016(03):3-8

Survey

- | | | |
|----|---------------|---|
| 01 | Aachen | Synthetic Biology Survey |
| 02 | AHUT | DNA 计算及最短路径规划 |
| 03 | BIT-China | Questionnaire about person health, a medical condition, breast cancer cognitive.
关于个人健康、体检情况、乳腺癌认知以及产品调研 |
| 04 | EPFL | Your opinion about your iGEM participation |
| 05 | Evry | Let's PLAY: Polylactic Acid Plastic Revolution |
| 06 | HUST | 基因表达问题的问卷调查 |
| 07 | IIT_Kharagpur | Survey 1 |
| 08 | Jilin_China | 实体瘤的认识及其治疗方式 |
| 09 | Manchester | Alcohol consumption questionnaire |
| 10 | OUC-China | 合成生物学及定量化与定性化认知情况 |
| 11 | Peking | 2016 国际基因工程大赛 (iGEM) 北京大学 Peking 团队项目调查问卷 |
| 12 | Peshawar | iGEM Peshawar |
| 13 | UCAS | Antibiotic residues/resistance |
| 14 | UCL | Synthetic Biology survey |
| 15 | USTC | 朊病毒的认识及科研应用 |
| 16 | Valencia_UPV | goo.gl/forms/offWSLR5 |
| 17 | XMU-China | Antibiotics and Antibiotic-Resistance Bacteria
抗生素调查问卷 |

17 **BGU_ISRAEL**

We are the BGU iGEM Team and our goal is to devise several approaches using synthetic biology tools for efficient plastic biodegradation using bacteria. In addition, we plan to utilize the high energy stored in polyethylene terephthalate (PET) molecules, for electricity production. We have created a game which illustrates the basic idea behind our project, our bacteria secretes LC-cutinase an enzyme found to degrade PET and uses it's biocatalysis products to create energy.

a link to our game: <http://bgu-plastikiller.netne.net/#> (PC use only)

18 **Hannover**

This year, we are organizing an online survey about our project TALEbots and its relevance for other scientists. It would help us a lot, if you could spend a few minutes to answer our questions on the following website: <https://www.umfrageonline.com/s/TALEbots>
Thank you very much!

19 **NCTU_Formosa**

We let the public understand our PANTIDE and device by doing a public survey. In this survey, we use easy words to introduce our project and current agricultural overview. And how we solve the problem. Here is our survey's link.
Link: <https://tony367.typeform.com/to/ZtmVG6>

20 **Pretoria_UP**

We have surveyed experts in the field of photobio-electrochemical cells and if more want to fill in the survey the link is <https://www.surveymonkey.com/r/XGR8RYX>.

21 **Tec-Chihuahua**

Our Human Practices team created a survey for people familiar with both synthetic biology and agriculture. If any iGEM team can answer it or share to people who can, it'd be very helpful for our investigation. Here's the link to our survey: <https://docs.google.com/forms/d/e/1FAIpQLSf8mKaPEfKZd9DgnzRuXSvBWqkiVDgTZsgzD4P7UhtKRucPCQ/viewform>

22 **UPO-Sevilla**

OPINION POLL ON GENETICALLY MODIFIED ORGANISMS

We are a group of bachelor students who belong to the iGEM competition, which tries to develop an open community for the advancement of synthetic biology.

[Link_encuesta](#)

23 **Vilnius-Lithuania**

We have created a survey for previous iGEM teams. The survey is aimed at finding out what makes a successful iGEM team. We are analyzing several criteria – the size and the composition of the team, work distribution, number of mentors etc. – and its correlation with the achievements of the team. We are planning to announce the results of the survey on our wiki page. If you are interested, pass the survey to the previous teams from your institution. The [link: https://goo.gl/forms/BSKLq4lZeHnvXGD01](https://goo.gl/forms/BSKLq4lZeHnvXGD01)

Address book

Aachen



@iGEMAachen



iGEMAachen



igem@rwth-aachen.de



igem.Aachen

AHUT-China



Igem Ahut



ahutigem@gmail.com



AHUT-iGEM

BIT-China



@iGEM_BIT



igem_bit@outlook.com



iGEM_BIT



iGEM_BIT

Aalto-Helsinki



@AaltoHelsinki



Aalto-Helsinki iGEM



team@aaltohelsinki.com



aaltohelsinki_igem



blog.aaltohelsinki.com

BGU_ISRAEL



@igembgu2016



iGEM BGU



igembgu2016@gmail.com

Cardiff_Wales



@iGEM_Cardiff



golshaiea@cardiff.ac.uk

CGU_Taiwan



@iGEMCGU_Taiwan



CGU iGEM Taiwan



igemcgu@gmail.com

Duesseldorf



@iGemHHU16



iGEM HHU



igem@hhu.de

Endinburgh_UG



@EdiGEM2016



EdiGEM2016



edinburgh.igem2016@gmail.com

EPFL



@EPFL_iGEM



igemepfl



igemmers2016@groupes.epfl.ch

Evry



@iGEM_Evry



iGEM.Evry.2016



evryigem2016@gmail.com



igem_evry

Freiburg



igemfreiburg2016



igem_2016@freigem.org

Groningen



igemgroningen2016



ilonamager@outlook.com

Hannover



@igem_hanover



iGEMTeamHannover



info@igem.uni-hannover.de

HUST

 HUST_iGEM2016@163.com

IIT_Kharagpur

 @igem_iitkgp

 iGEMIITKharagpur

 rhushikeshphadke@gmail.com

Jilin_China

 @IGEM Jilin_China

 jilin_china@outlook.com

 iGEMxHU

 +86 15754305638

NCTU_Formosa

 NCTU_Formosa-IGEM-team

 nctu5168victory@gmail.com

HokkaidoU_Japan

 @igem_hokkaidou

 iGEM HokkaidoU

 yukinosbt@eis.hokudai.ac.jp

Imperial_College

 @imperialigem

 2016imperialigem

 igem.imperial2016@gmail.com

Leiden

 @iGEM_Leiden

 igemleiden

 igem@science.leidenuniv.nl

Manchester

 @iGEMManchester

OUC-China



iGEM OUC



oucigem@163.com

Peking



pkuigem2016@126.com

Pretoria_UP



@iGEMPretoria_UP



iGEM Pretoria_UP



PretoriaUPiGEM@gmail.com

SYSU-MEDICINE



@iGEM SYSU-MEDICINE



IGEM SYSU-Medicine



SYSU_MEDICINE@163.com



SYSU_MEDICINE

Oxford



@oxfordigem



oxfordigem



oxfordigem@bioch.ox.ac.uk

Peshawar



iGEM Peshawar 2016



sidra.usman.biotech@gmail.com

Queens_Canada



@igemQueens



QGEM



director@qgemteam.com

SYSU-CHINA



@iGEM_SYSUCHina

Tec-Chihuahua



@iGEM_TecChih



iGEM ITESM Chihuahua



itesmigem@gmail.com

Tongji_Shanghai



will wang/Tongji Shanghai



+86 15821399962

UCAS



ucasigem@163.com

UCL



UCL iGEM



ucligem2016@gmail.com

UNIK_Copenhagen



@iGEM_TecChih



CosmoCrops



igem16.cph@gmail.com

Tel-Hai



iGEM Tel Hai



igemtelhai@gmail.com

Tianjin



@iGEM_Tianjin



yuanma@tju.edu.cn



iGEM2016Tianjin

UCC_Ireland



@ucc_igem



UCCIgem



corkigem@gmail.com

UPO-Sevilla



@iGEM_UPO



Igem Upo



igem.upo@gmail.com

UrbanTundra_Edmonton



UrbanTundra Edmonton - IGEM HS 2016



urbantundraedmonton@gmail.com

USTC



yukae@mail.ustc.edu.cn

Vilnius-Lithuania



@Vilnius_iGEM



VilniusiGEM



vu.igem@gmail.com

Westminster_UoW



@WestminsteriGEM



westminsterigem2016



uow.igem@gmail.com

Valencia_UPV



@UPVigem



Valencia UPV iGEM



Valencia.upv.igem@gmail.com

Washington



UW IGEM Team



uwigem@uw.edu

XMU-China



@Amoy_igem



IGEM XMU-China



igemxmu@gmail.com