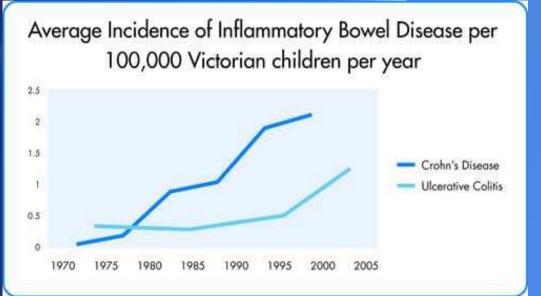
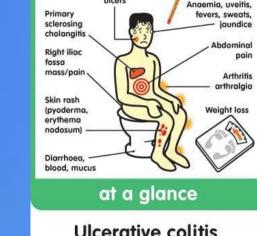
Why Crohn's Disease?

wel disease that has symptoms which Crohn's disease is a type of inflammatory local. Nowadays it spreads more mostly among the youth (see th is in constant inflammation and ap below). A part of the gastro-intesting that the disease can cause ulcers, vo diarrhea and anemia (see the figur Over 600 000 people in the US suffe is disease.

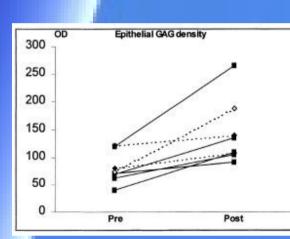


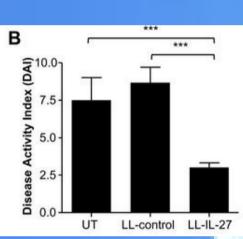


Ulcerative colitis and Crohn's disease

Why NAG and Why Probiotic Bacteria?

ed their patients (age up to 16), with NAG (N Researchers from En ly and observed that 8 out of 12 of them y biopsy and other means and published their res ional substrate for glycosaminoglycan synthesis, in paediatric chronic ished an article on making and transferring producing bacteria into the intestines of mice w

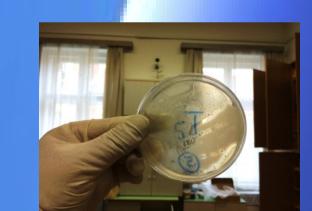




The Role of Natural Chitinases in Our Project

ne basic material of many organisms such as fu and worms. Due to nce of chitin in soil, some soil decomposers produ ected soil samples from different locations and in o examine them, we have olloidal chitin agar medium. If they break down ch e can see clearing zones d the colonies on the colloidal chitin agar as visible





In order to determine how chitinases are able to he intestine, we ordered Sigma Aldrich Chitinase Assay Kit (SACAK) to study y of natural chitinase enzymes under different environmental conditions. y is based on the enzymatic hydrolysis of chitinase substrates and it release ophenol which can be measured colorimetrically at 405 nm. We used BioRad I rk microplate reader to carry out these measurements.

Our key findings were as follows, 1. The SACAK was appropriate for measur tural chitinase activity. We could see strong color changes at the kit's own ch

2. The measurment of the natural chit of Bacillus thuringiensis liquid media showed 2.08 U/ml activity according to the g ule by the SACAK provider. 3. The chitinase activity of Bacillus f sis bacteria was the highest if it was

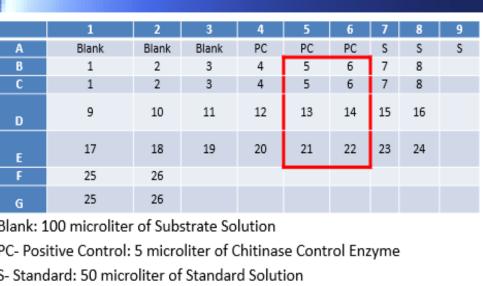
4.8-5.8. These findings strengthen the cultured in shaking plate on 37°C, idea of GMO-based chitinase acti a normal large intestine environment conditions.

4. The large intestine extract of the rat n't show any chitinase activity itself. 5. The large intestine extract didn't ha y enhancing or inhibiting effect on natural

chitinase activity. 6. We found strong exochitinase activ t no endochitinase activity was measured.



0.023 0.011 0.076 0.085 0.084 0.065 0.077 0.072 0.078 0.078 0.071 0.069 0.094 0.091 0.081 0.069 0.082 0.104 0.094 0.083 0.074 0.077 0.068



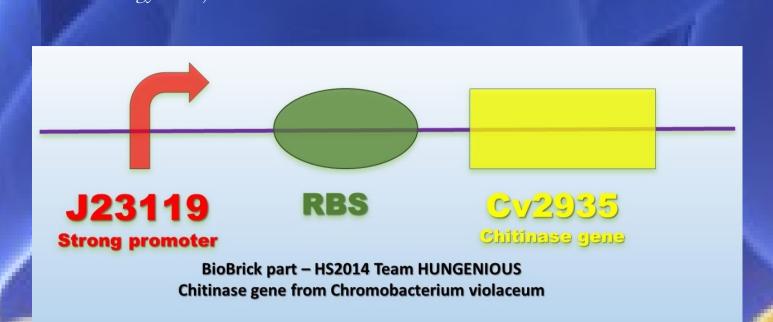
25: 5 microliter of PBS + 2,5 microliter of healthy rat intestine

26: 5 microliter of PBS + 2,5 microliter of inflamed rat intestine



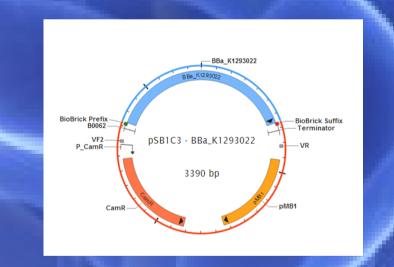
BRC

We have constructed two BioBricks based on our experimental design. One of them is BBa K1293022 which contains the Cv2935 gene of Chromobacterium violaceum. The other one is BBa K1293023 which is constructed from a strong promoter BBa 23119 and our aforementioned basic part. The article in which we found our target gene is: and efficient secretion of a functional chitinase from Chromobacterium violaceum in Escherichia coli. Lobo et al. BMC Biotechnology 2013, 13:46 doi:10.1186/1472-6750-13-46



PROBIOTIC CHITINASE PRODUCING BACTERIA ENGINEERED AGAINST CROHN'S DISEASE







Our Future Plans

Research quetsions:

- We would like to prove the tissue repairing effect of NAG coming from our GMO bacteria
- We would like to find an appropriate mean of transfer of GMO bacteria into mice/humans

Biosafety questions:

- Does our GMO bacteria cause any changes in the natural microflora of the colon?
- Can we ensure that Gmo bacteria will die after leaving the patients' digestive

Genetic Experiments

irst, we were going to get DNA from DT (Integrated DNA Technologies), but unfortunately, ole to synthesize the chitinase gene. In the lack of time r many attempts, the company was un plan was the following: we ordered enomic DNA of Chromobacterium violaceum from

we had plenty of things to do, our tean up into 2 smaller groups: someone was working at researchers, whilst the others were working at the Biological Research Centre (BRC) with 1

nstead of standard, and annealing temperatures the BRC we have used gradient PCR techn so had to be set to an appropriate value. neously, we also amplified the pZA31 plasmid, ted. To avoid 3' overhangs different polymerases to which the Cv2935 gene was going to be: nd buffers had to be used in both reactions. cide whether the PCR reactions were successful, e ran the products on gel. The results sho that, the 'Phusion polymerase' was the most ccessful with the annealing temperature of 5 is it can be seen in these pictures.

e prepared the electrocompetent cells by su , chilling and centrifuging 3 times, then we bjected the cells to electroporation in order form our plasmid, containing the chitinase ne, into Nissle bacteria (E.coli strain). Specia ttes and an electroporator was used for this urpose. Unfortunately our transformation exper nt was unsuccessful.





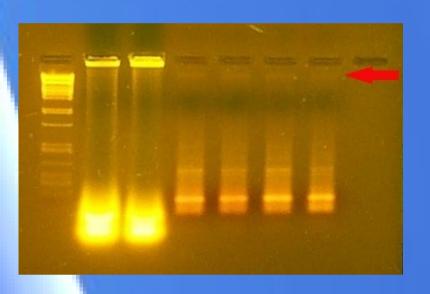
SANOFI

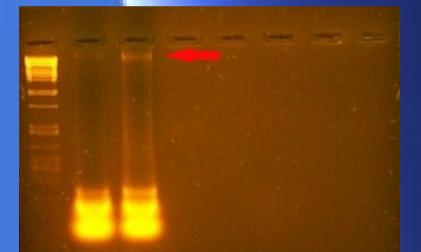
Ist well: size ladder **2nd~6th well:** Samples treated at different annealing temperatures. he red arrow indicates the slightly visible line of DNA under ne 5th well. During the PCR of the DNA of the 5th well, the realing temperature was set to 58C.



4th well: Samples treated with different polymerases: we the DNA in the second and third well with the i Taq polymerase. We added Phusion High-Fidelity ase to the DNA, that can be found in the $4^{
m th}$ well. The ow shows the line of chitinase gene at 2kbps.

e the procedures at BRC were going on, similars wei rried out at our school and at the ersity of Szeged. Though, here we wanted to construc Brick in a pSB1C3 backbone, not o transform a plasmid into bacteria. We amplified th ase gene using colonial PCR and stion. After we have purified the prefix and suffix primers. PstI & EcoRI were used for a ligation was performed according to the table abo st, the product of the ligation nsformed into <mark>DH5-alpha strain of E. coli</mark> with hea ck method. All the experiments cked by gel-electrophoresis, even from the transforme lonies we have made colonial results can be seen on the figures below.





Human Practices

project we had several meetings with people who c In order to pro problem we worke met for example:

A medical doctor Farkas M.D.), who works with patients affected by C. hn's disease.

talk to medical students. We had an oppo

We had the pr k about our project to Nobel prize winner Aaron Chi anover, who visited our school

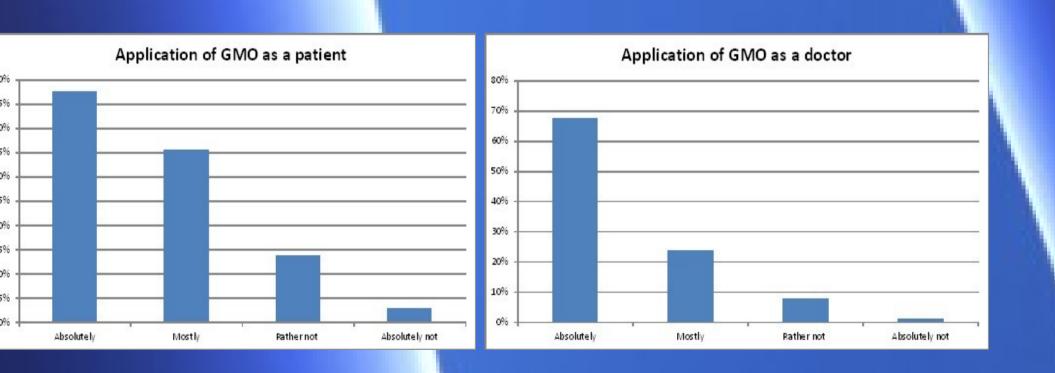
ke a survey to find out people's opinion about our idea Moreover we have de topic in which it may l. We created this survey using SuveyMonkey. It has 101 times mostly by peo Our key findings were as

disease is believed to be a moderately spread disorder 1. Considering these rea

ne survey more than 35% approves non-medical appl 2. In accordance to the of a genetically modifie

f those, who would take part in a treatment which inv 3. We could also conclude GMO: 45% would not re cured with GMO. 4. Interestingly, 70% claim they were a doctor they would absolutely use a treatme

lotic bacteria.



Team and Acknowledgments

Miklos Boldogkoi, Reka Fabian, Mark Harangozo, Our team consists of 8 pupils all from Akos Marton, Gergo Nikolenyi, Martin l etra Varga, Andras Volford.

Our mentors were: Sandor Ban (RMG-Balint Csorgo (BRC-Szeged), Akos Nyerges (BRC-

Main designer of our website: Csongor Kis Szeged) Institutions involved:

BRC: Biological Research Centre of Hu Academy of Sciences,

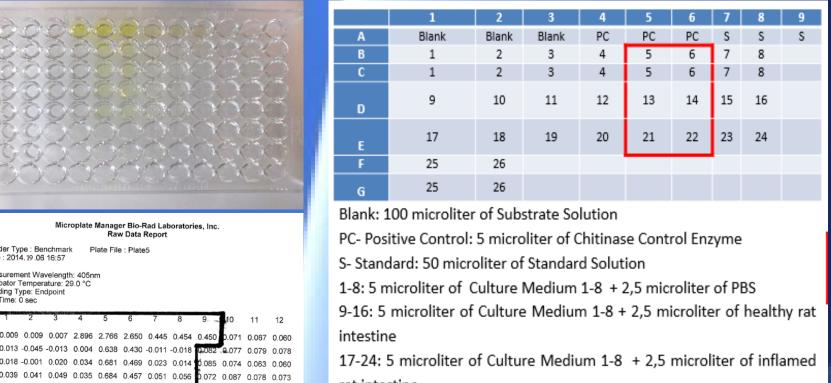
RMG: Radnoti Miklos Experimental Gra chool, Szeged

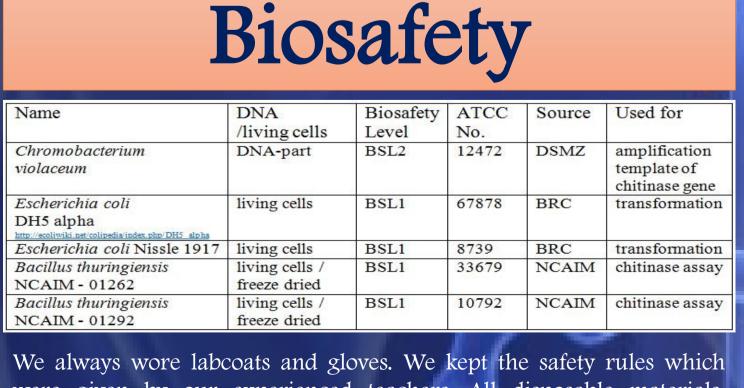
Department of Physiology, Anatomy and I cience at the University of Szeged

The sources of bacteria and genom: Soil samples

involving genetically modi

NCAIM: National Collection of Agricult ndustrial Microorganisms – Hungary DSMZ: Deutsche Sammlung von Mikroor n und Zellkulturen





were given by our experienced teachers. All disposable materials, (pipette tips, Eppendorf- and PCR-tubes, inoculating loops, etc.) connecting with microorganisms were put into a biohazard container, filled with strong oxidative disinfectant.

We have used microorganisms belonging only to BSL1 level. The only exception is Chromobacterium violaceum which is itself a BSL2 bacterium; however we have used only parts of its DNA (chitinase gene). The product of this gene is not connected with the Chromobacterium violaceum's pathogenic effect.