

A circular graphic containing a grayscale illustration of various microorganisms, including bacteria with flagella and a DNA double helix structure, set against a light background.

ADVANCES IN MICROBIAL BIOTECHNOLOGY

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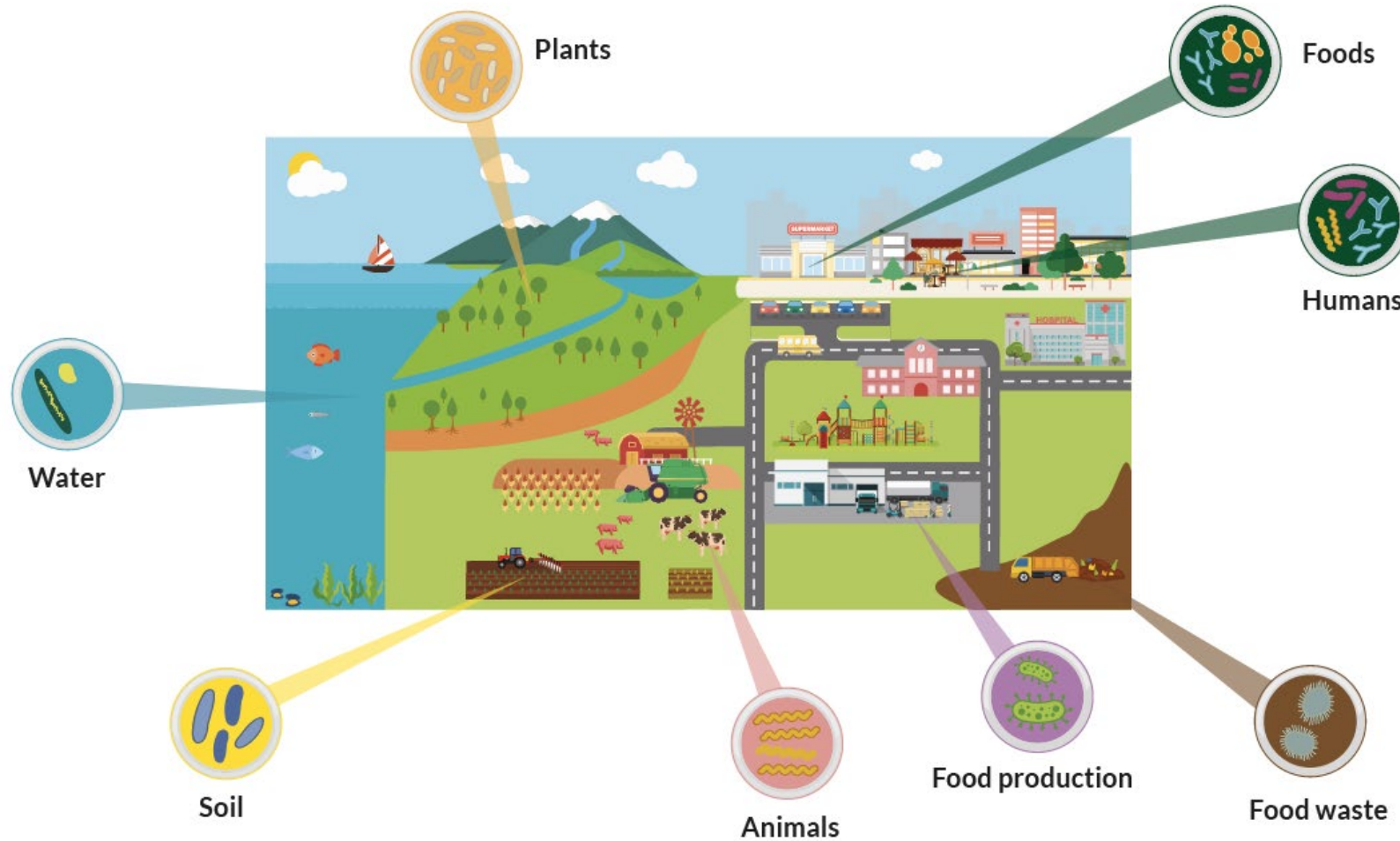
DISCLAIMER

- **Employee of Société des Produits Nestlé SA (Switzerland)**
Senior R&D specialist in Biotransformation; Nestle Culture Collection coordinator
Expertise in beneficial microbes, their metabolism and genetics (probiotics and fermentation)
- **PhD in Microbiology from Wageningen University (The Netherlands)**
Host & Microbe Interactomics group; Prof. M. Kleerebezem



Research and
Development

MICROBES ARE EVERYWHERE...



Extremophiles



Hot springs, fumaroles

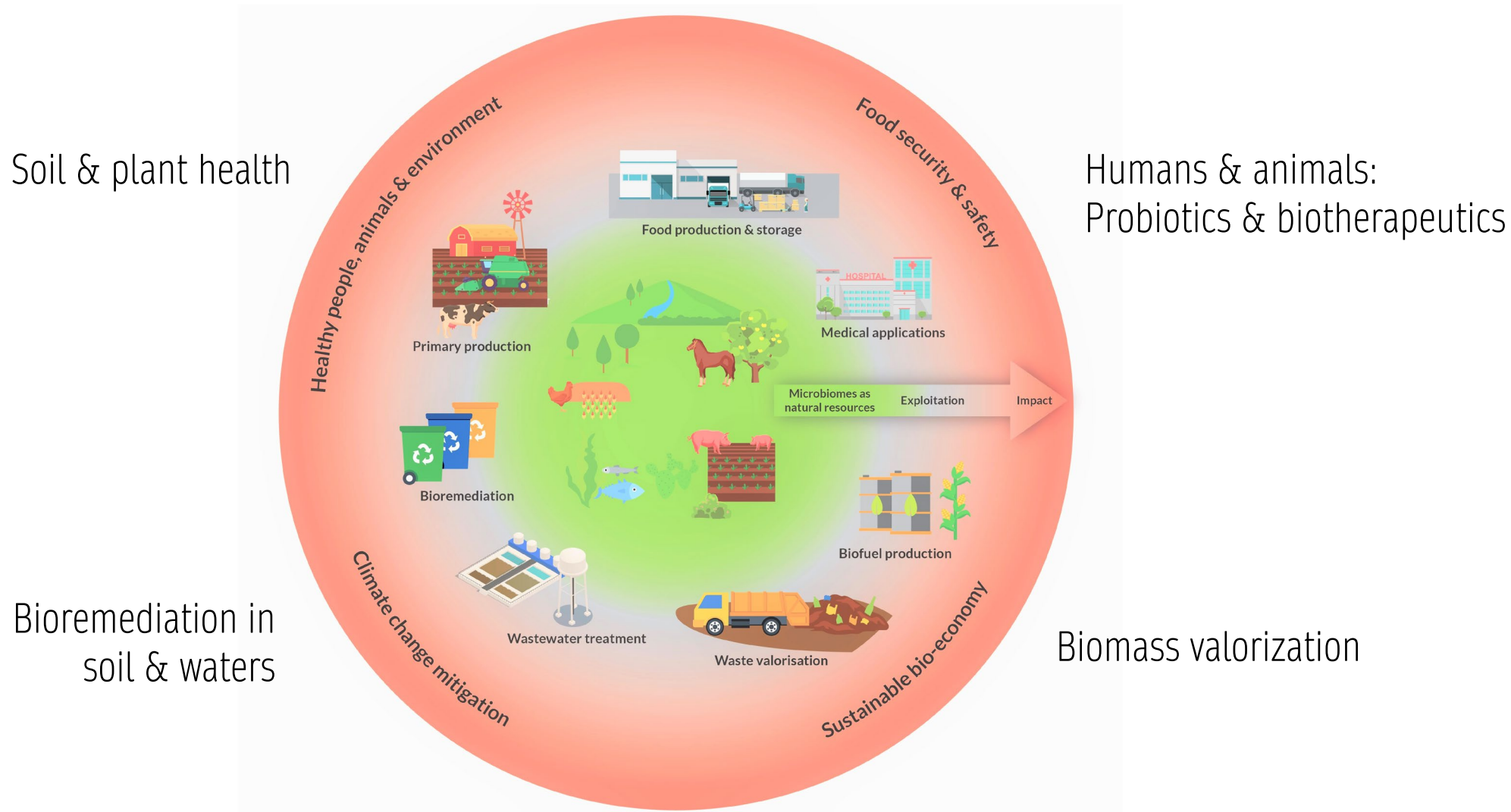


Hypersaline lakes



... AND POSSESS TREMENDOUS VALUE

Fermented foods



COMPLIANT EXPLOITATION OF MICROBES

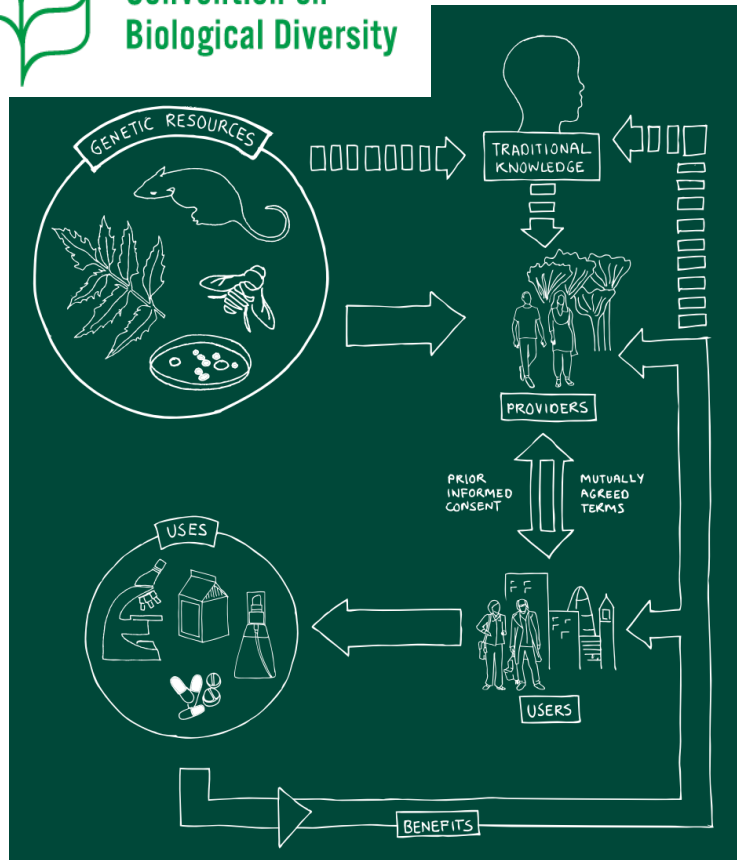
Environmental isolates
(soil, water, plants, etc..)



Human isolates



Convention on Biological Diversity



International Ethical Guidelines for Health-related Research Involving Humans



WMA Declaration Of Helsinki : Ethical Principles For Medical Research Involving Human Subjects

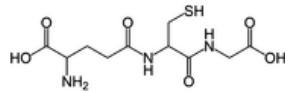


ICH -Guidelines for good clinical practices

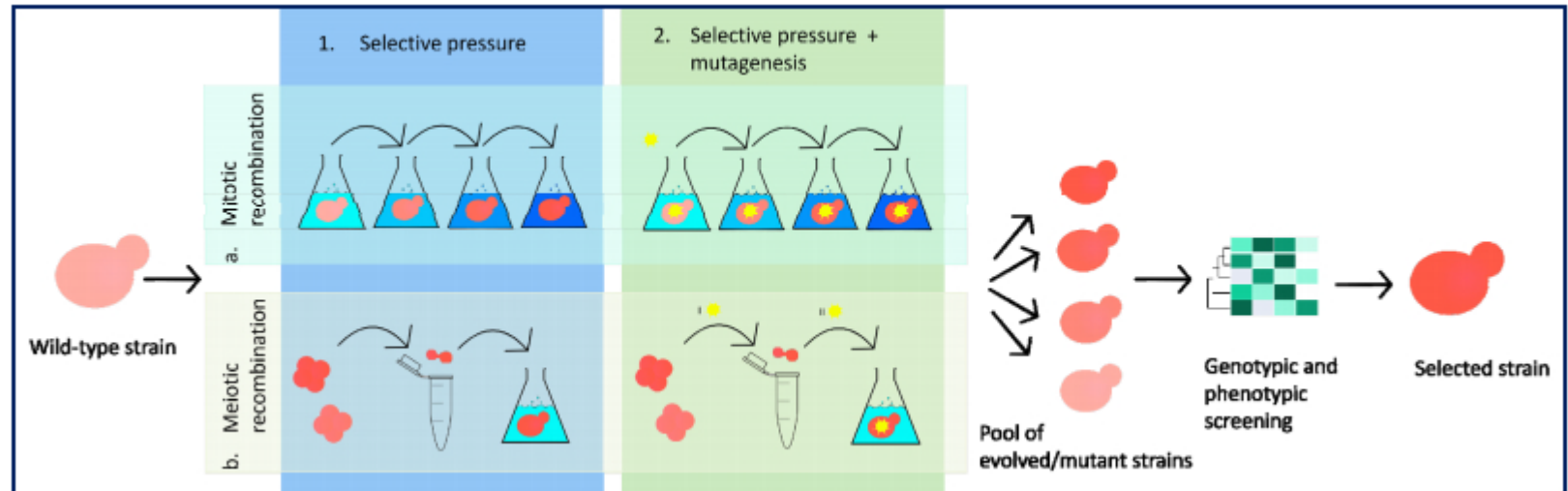


ENHANCE NATURE'S CAPACITIES ADAPTIVE EVOLUTION

Glutathione



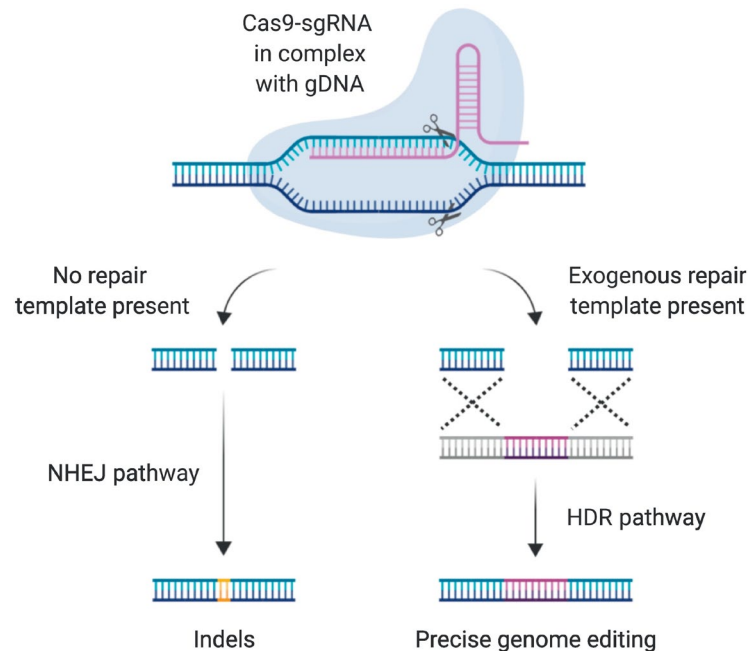
“Adaptive evolution results from the propagation of **advantageous mutations through positive selection**”



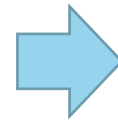
De Vero et al. AIMS Microbiology, 2017, 3(2): 155-170

ENHANCE NATURE'S CAPACITIES TARGETED GENOME EDITING

CRISPR-Cas9



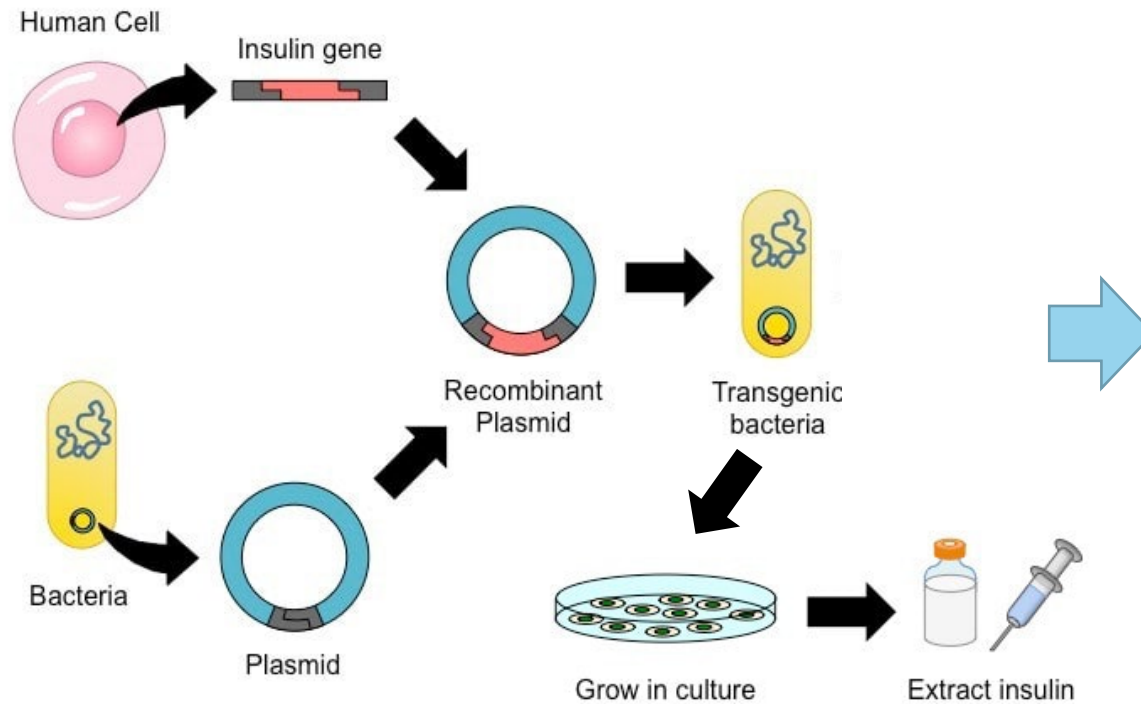
Trends in Molecular Medicine



Industrial strains	product
<i>Synechococcus elongatu</i>	succinic acid
<i>Saccharomyces cerevisiae</i>	free fatty acid
<i>E.coli</i>	n-butanol
<i>Corynebacterium glutamicum</i>	glutamic acid
<i>Saccharomyces cerevisiae</i>	β -carotene
<i>Bacillus subtilis</i>	BLA
<i>Clostridium tyrobutyricum</i>	n-butanol
<i>Corynebacterium glutamicum</i>	γ -amino-butyric acid
<i>E.coli</i>	fatty acids
<i>E.coli</i>	5-amino-levulinicacid
<i>E.coli</i>	isopropanol
<i>E.coli</i>	β -carotene
<i>Bacillus subtilis</i>	riboflavin
<i>Myceliophthora thermophile</i>	cellulase
Filamentous fungus	mucic acid
<i>Aspergillus niger</i>	galactaric acid

Zhao et al. 2020, Synthetic and Systems Biotechnology 5 (2020) 269–276

«COPYING» NATURE CAPACITY RECOMBINANT DNA TECHNOLOGY

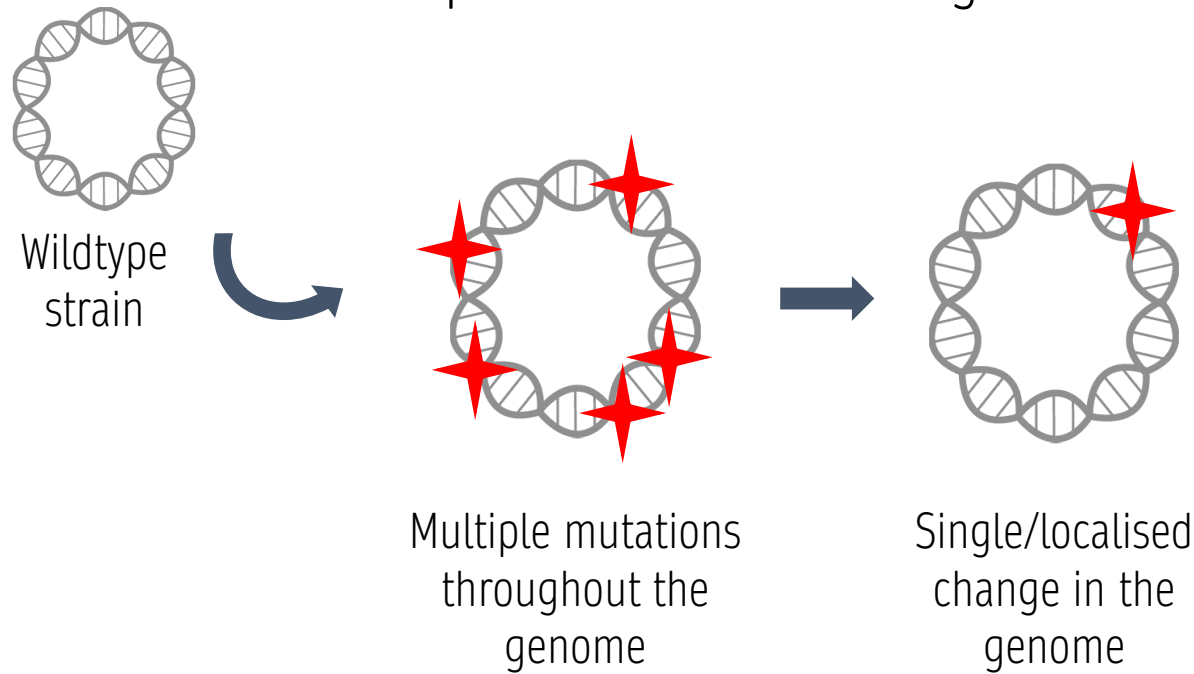


<https://mysciencesquad.weebly.com/>

Chassis cells	Therapeutic payload	Diseases	Development stages
<i>Lactobacillus</i>	IL-10	IBD	Mice
<i>Lactobacillus</i>	IL-4	IBD	Mice
<i>E. coli</i>	IL-35	IBD	Mice
EcN	Trefoil factor	IBD	Mice
<i>Bacillus thermophilus</i>	Superoxide dismutase	IBD	Mice
<i>Bifidobacterium</i>	RhMnSOD	IBD	Mice
<i>Lactobacillus</i>	Elafin	IBD	Mice
<i>Lactobacillus</i>	Recombinant mouse heme oxygenase-1	IBD	Mice
NZ9001			
<i>Lactobacillus</i>	Pancreatitis-related protein	Intestinal mucositis	Mice
EcN	Butyrate	Colon cancer HT29	Mice
<i>Salmonella</i> , <i>Typhimurium</i>	IL-1β	Colon cancer	Mice
		CT26	
<i>E. coli</i>	B-glucuronidase	Colon cancer	Mice
<i>Salmonella</i> , <i>Typhimurium</i> ,	Autoinducer	Colorectal cancer	Mice
		MC26	
EcN	Tum-5	Melanoma	Mice
<i>Salmonella</i> VNP20009	Sox2	Lung cancer	Mice
<i>Salmonella</i>	Transforming growth factor alpha- <i>pseudomonas</i> exotoxinTGfa-PE38	Colon cancer CT26 & Breast cancer 4T-1	Mice
<i>Salmonella</i> SL7207	Diaminopimelate DAP	Hepatocellular carcinoma	Mice
<i>Lactobacillus</i>	GLP-1	Diabetes	Mice
<i>Lactobacillus</i>	Heat shock protein 65HSP65, IA2P2	Diabetes	Mice
<i>Lactobacillus</i>	GLP-1	Obesity	Mice
<i>Bacillus subtilis</i> SCK6	Butyric acid	Obesity	Mice
<i>Bacillus subtilis</i> SCK6	BA	Obesity	Mice
EcN SYN1020	L-arginineI-arg	HyperammonemiaHA	Stop
EcN SYN1618	Insert phenylalanine ammonia lyase and L-amino acid deaminase gene	PhenylketonuriaPKU	Phase 1/2a
<i>Lactobacillus plantarum</i>	Angiotensin-converting enzyme inhibitory peptidesACEIPS	Hypertensive	Mice
<i>Vibrio cholerae</i> strain Haiti V	Delete CTXF, CTXA, RECA genes	Cholera	Infant rabbit
<i>Meningitis</i> MenB YH102, YH103	Delete rfaF, metH, siaD	Meningitis	Mice
EcN	Insert Phl p1 and Phl p5 gene, control the level of IgE	Allergic poly-sensitization	Mice

CORRESPONDING GENETIC SIGNATURES

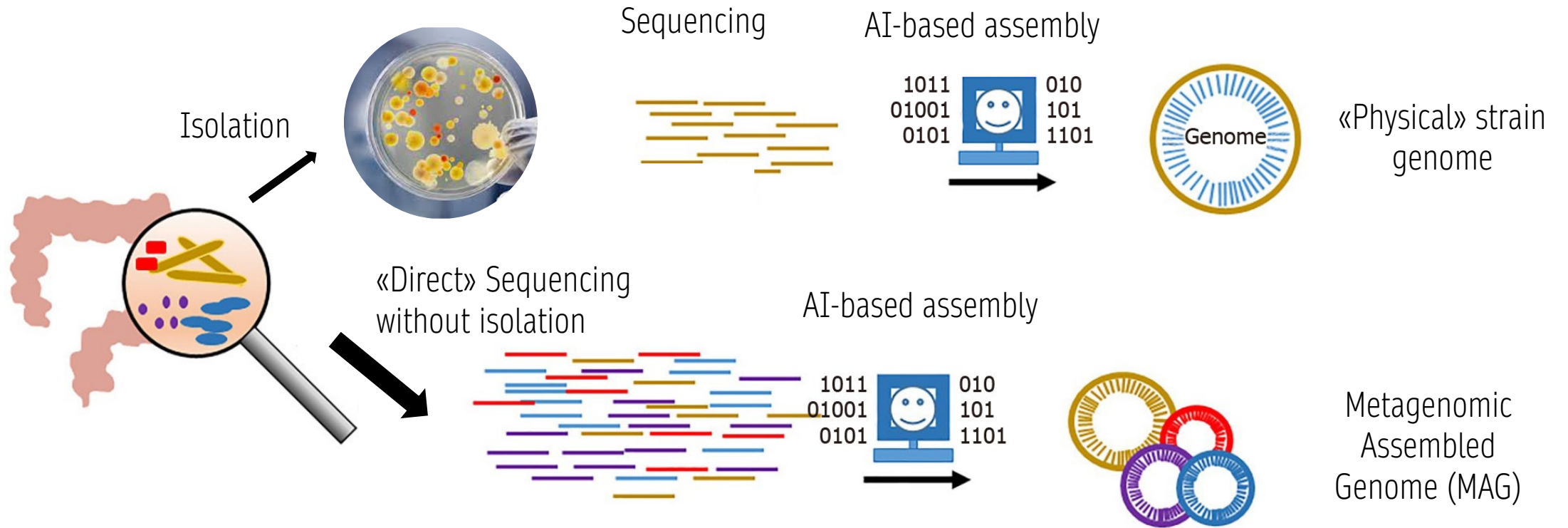
Adaptive evolution → targeted editing



Recombinant DNA technology



DO WE NEED «PHYSICAL» STRAINS ? METAGENOMIC ASSEMBLED GENOMES



DO WE NEED «PHYSICAL» STRAINS ? E.G. IN SILICO ENZYME DISCOVERY

«Physical» strain genome



Metagenomic Assembled Genome (MAG)

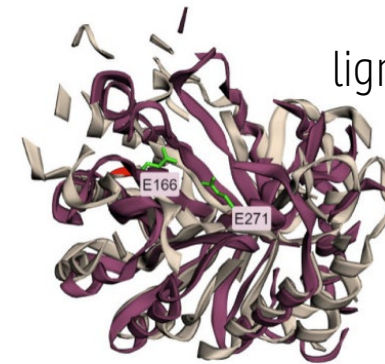


AI-base prediction of function



[NCBI Prokaryotic Genome Annotation Pipeline \(nih.gov\)](https://www.ncbi.nlm.nih.gov/genome/annotation_prok/pgap/)

In silico functional prediction
(e.g. enzyme with increased
lignocellulose degradation activity)



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LVLDICGKLGIIYCIIDWH-VLNEGSGN---PKNT----LN-DALPFWDYMSAKHKI
: ::::::::::::::::::::: ..... ..
IIVDLAESNGQSVRGHTLVVHNQL---PSWVDDVPAGELLGVMRDHITHEVDHFK
: ::::::::::::::::::::: ..... ..
CNEPNGFMVWRS----H---VKEYADQVIPVIRANDPKIIICGTPMWS----Q
: ::::::::::::::::::::: ..... ..
VNEAFEEDGSRQSVFQQKIGDSYIAEAFKAARAADPDVKLYNDYDIEGIGPK
: ::::::::::::::::::::: ..... ..
MPL--SYNNVMTLHF-YSG-DHTQSLRDKAQATLNNGAAIFVTEFGTTKASGNG
: ::::::::::::::::::::: ..... ..
3WUG SDAVYEMVKSFKAQGIPIDGVGMQAHLIAGQVPASLQENIRRFADLGVDAITELDIRM-TLPR
PersiCelXyn1 ---GV--FLDECDRWMMNERK-I-SWVNSF-ADK---N-----ESSAALQPGASRSKNW
3WUG : ::::::::::::::::::::: ..... ..
IAAKDAQQATDYGAVVEACLVVSRVCGITVWDYTDKYSWVPSVFPGQGAALPWE-----
PersiCelXyn1 NLVSESGQYIKOKLSQPK
3WUG : ::::::::::::::::::::: ..... ..
--DFAKK-PAYHAIAAALN
    
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Ariaeenejad et al. International Journal of Biological Macromolecules 177 (2021) 211-220

SUMMARY

- Microbes are everywhere, but their exploitation should be compliant with National/International reglementation
- Beyond recombinant DNA technology, new «non-GMO» technics can be used to improve/inhibit specific strain capacities
 - Depending on the technology used, the extend of genetic changes can vary a lot
- With the advancement of sequencing and artificial intelligence, it is today possible to discover new strains / functions, without going through «physical» isolation



THANK YOU



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