

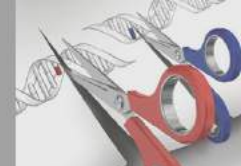
Genetika EA (BSc)

Genomszerkesztés

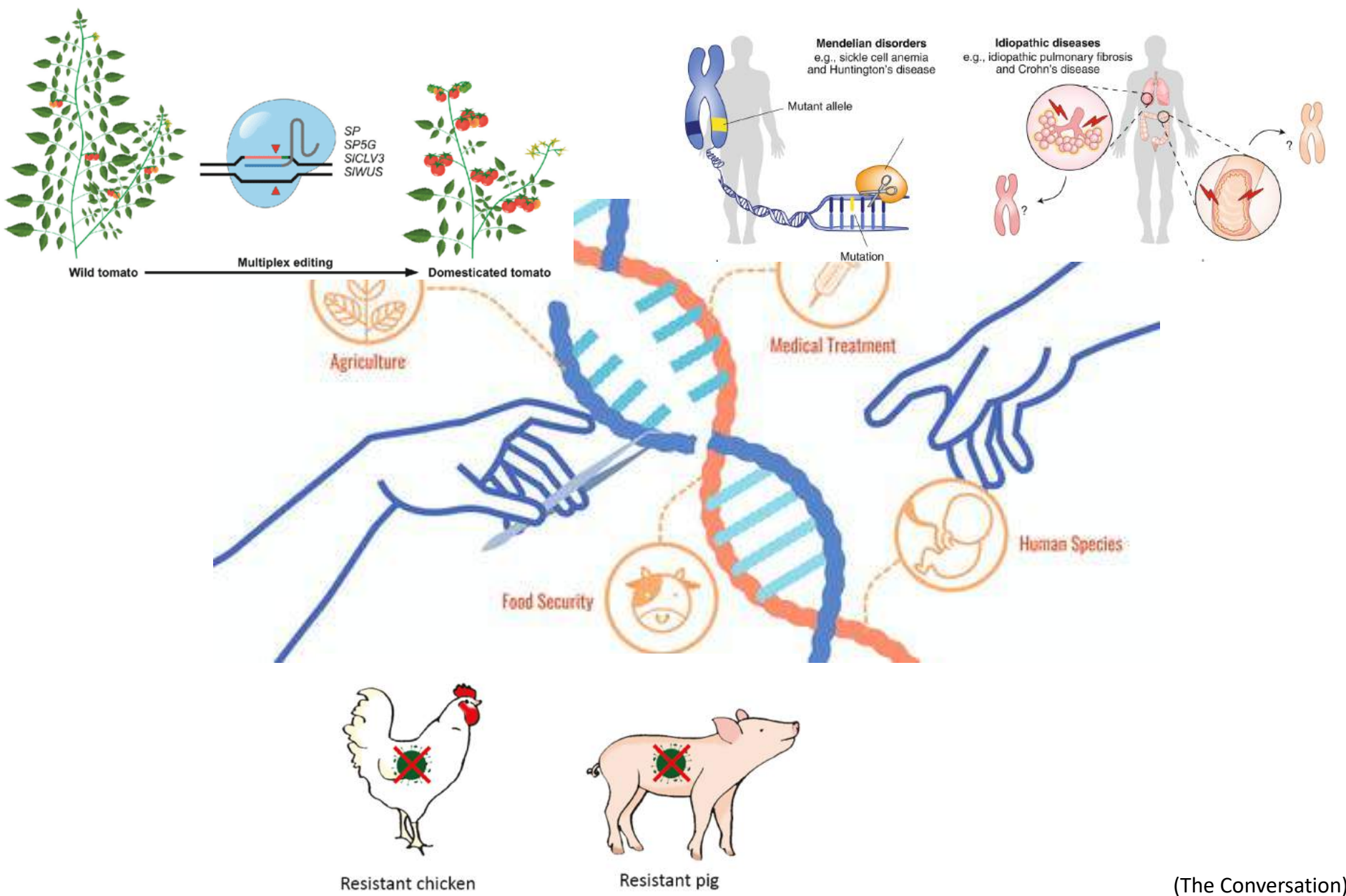


Varga Máté (Genetikai Tanszék)
mvarga@ttk.elte.hu

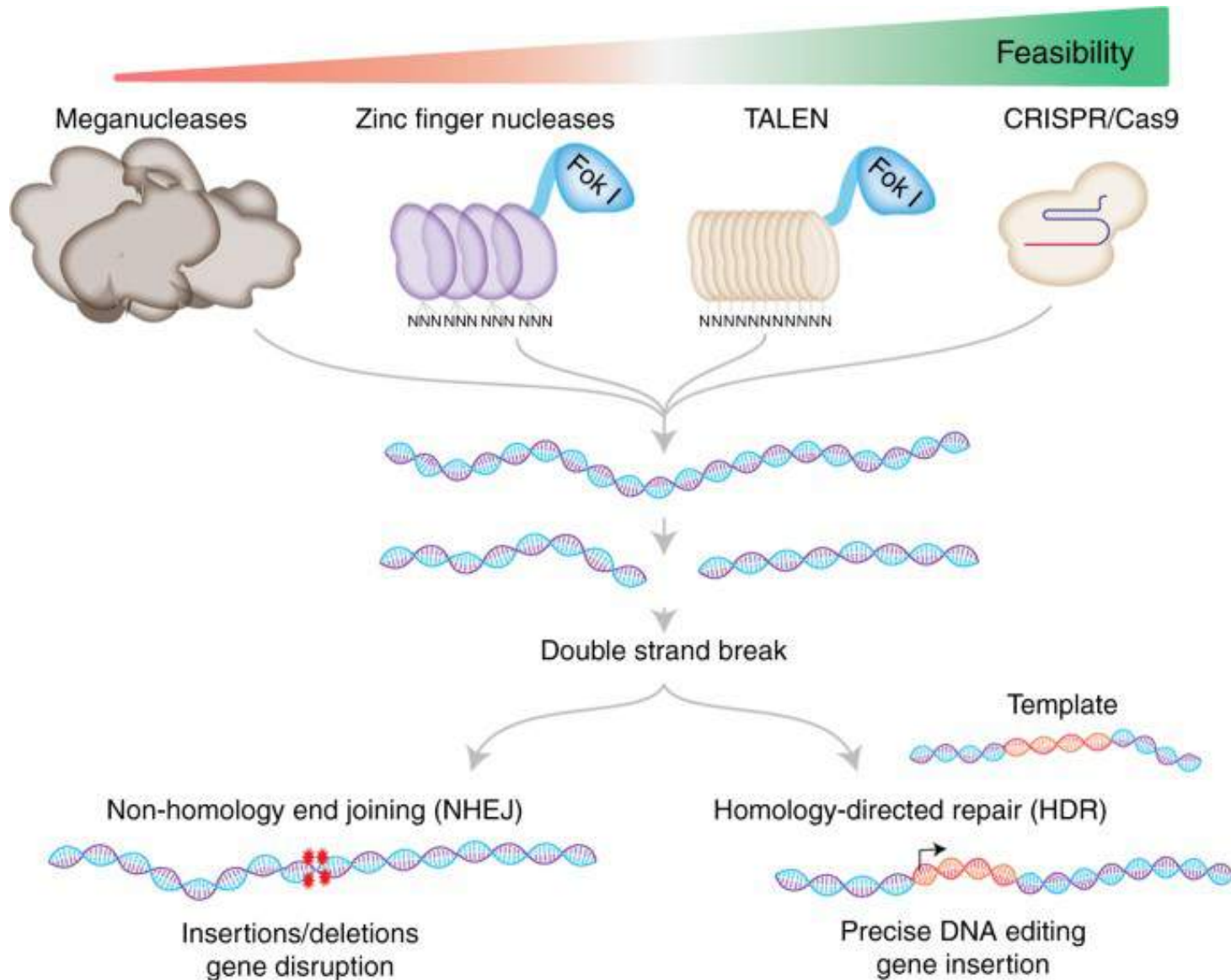
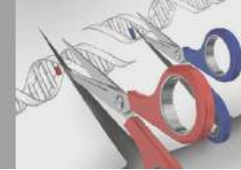
2020.10.21.



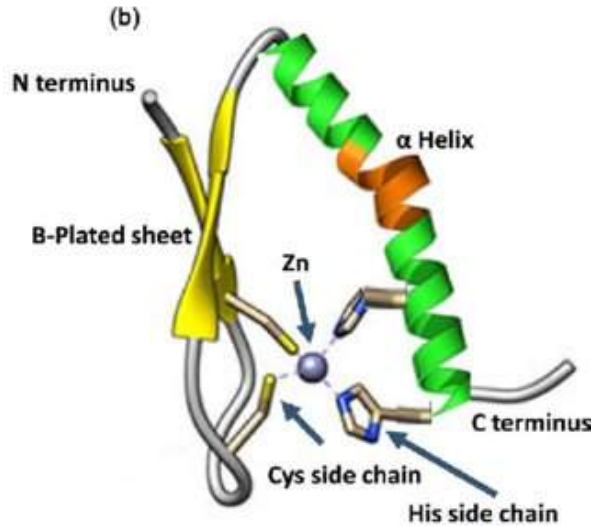
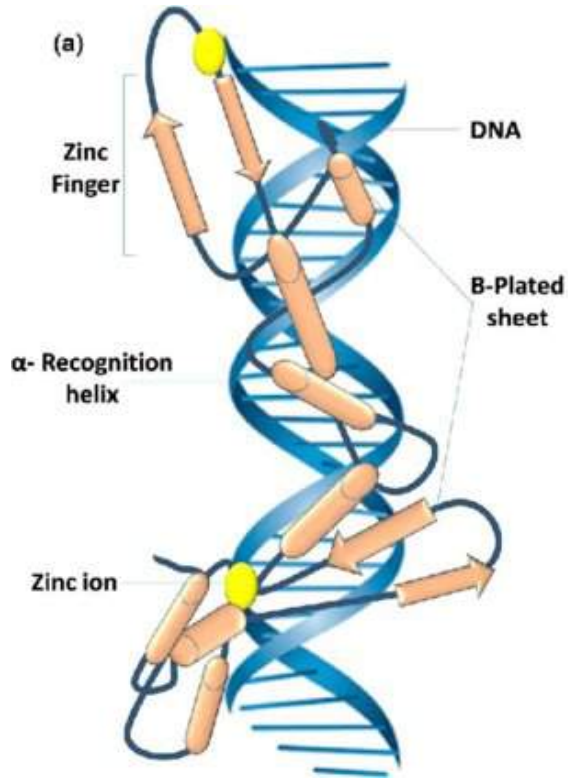
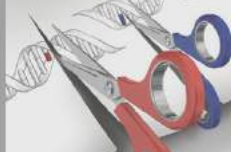
Miért akarunk „genomszerkeszteni“?



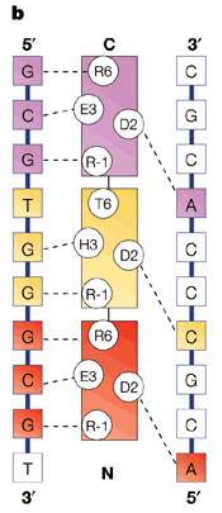
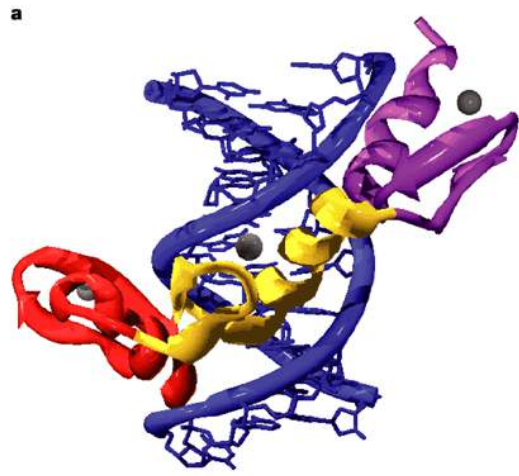
Genomszerkesztés a sejt szemszögéből: DNS hibajavító mechanizmusok



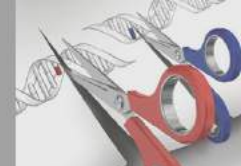
DNS-specifikus kötés: ZFN-nukleázok



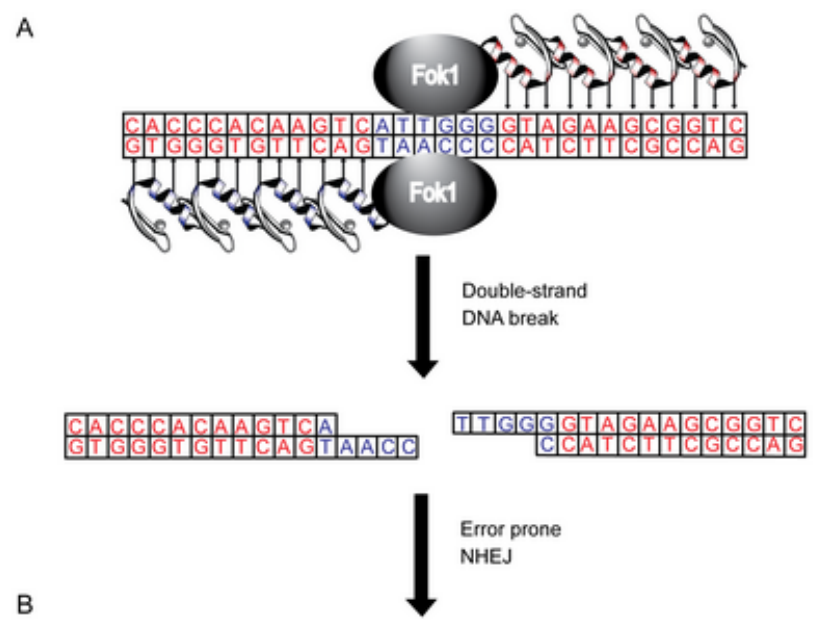
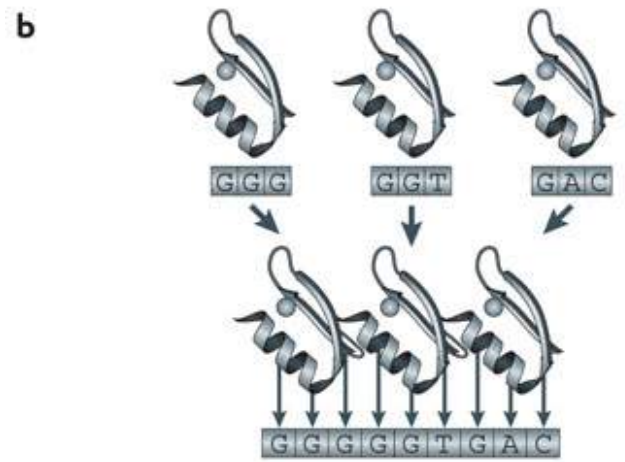
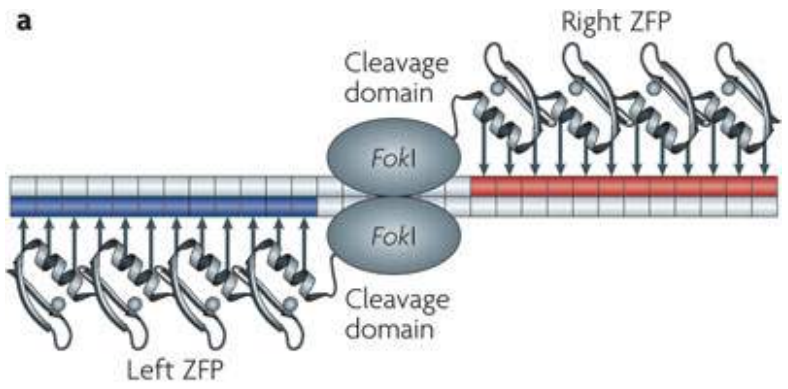
(Aslam et al. 2019 *Am J Mol Bio*)



(Jamieson et al. 2003)



DNS-specifikus kötés: ZFN-nukleázok

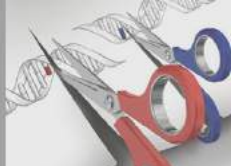


B

ACTGGAACACAACCACCCACAAGTCATTGGGGTAGAAGCGGTCACAGATATATC	Reference
ACTGGAACACAACCACCCACAAGTCATTGGTAGAAGCGGTCACAGATATATC	2bp deletion (1.4%)
ACTGGAACACAACCACCCACAA GTAGAAGCGGTCACAGATATATC	9bp deletion (4.5%)
ACTGGAACACAACCACCCACAA GAAGCGGTCACAGATATATC	12bp deletion (2.6%)
ACTGGAACACAACCACCCACAA GCGGTCACAGATATATC	15bp deletion (5.2%)
ACTGGAACACAACCACCCACAA GTCACAGATATATC	18bp deletion (11.2%)
ACTGGAACACAACCACCCACA GATATATC	25bp deletion (1.6%)
ACTGGAACACAACCACCCACAAGTCATTGGTTGGGGTAGAAGCGGTCACAGATATATC	4bp insertion (2.5%)

Nature Reviews | **Genetics**
 (Urnov et al. 2010)

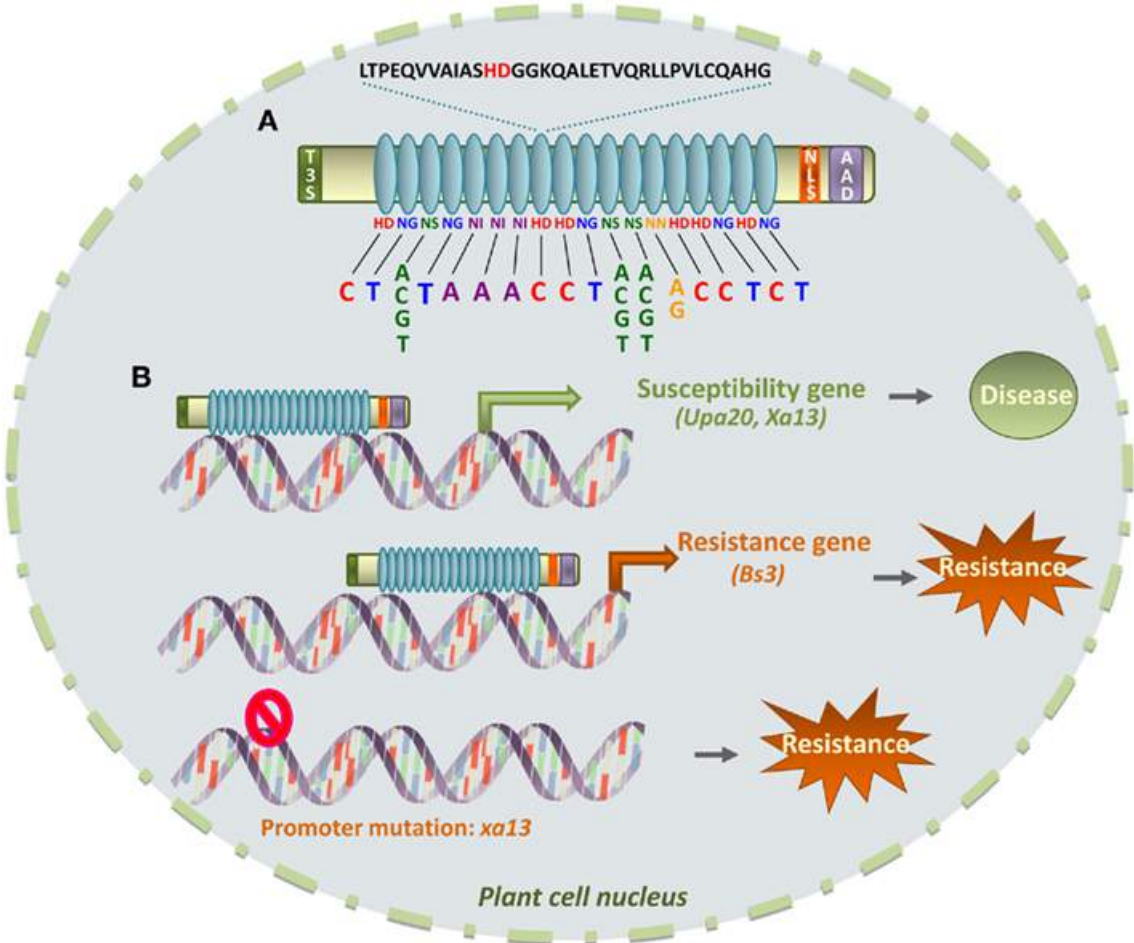
DNS-specifikus kötés: TALE nukleázok



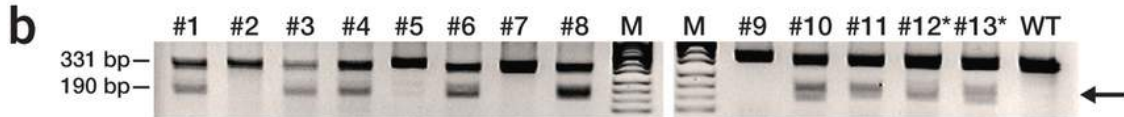
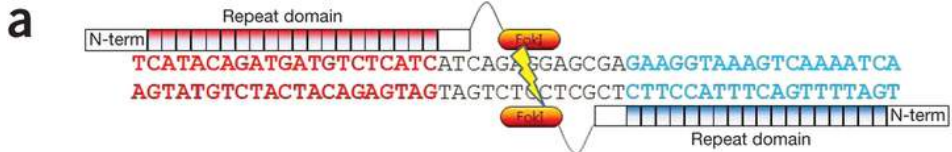
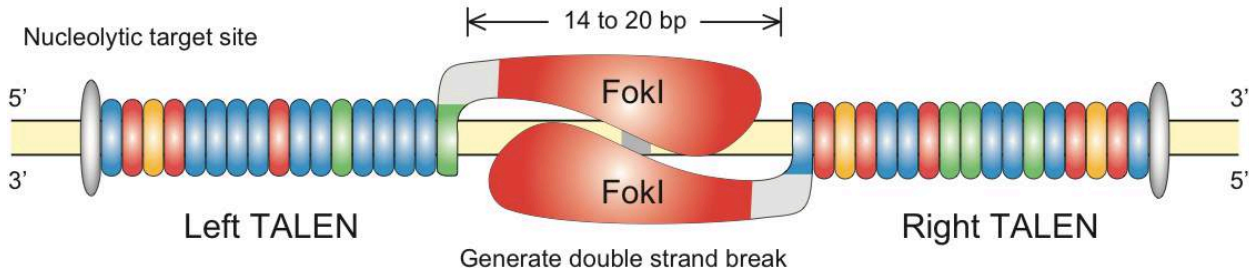
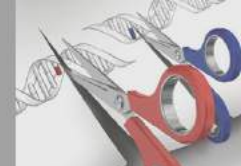
TALE = Transcription Activator Like Effectors



- *Xanthomonas* nevű növényi patogénből izolálták



DNS-specifikus kötés: TALE nukleázok

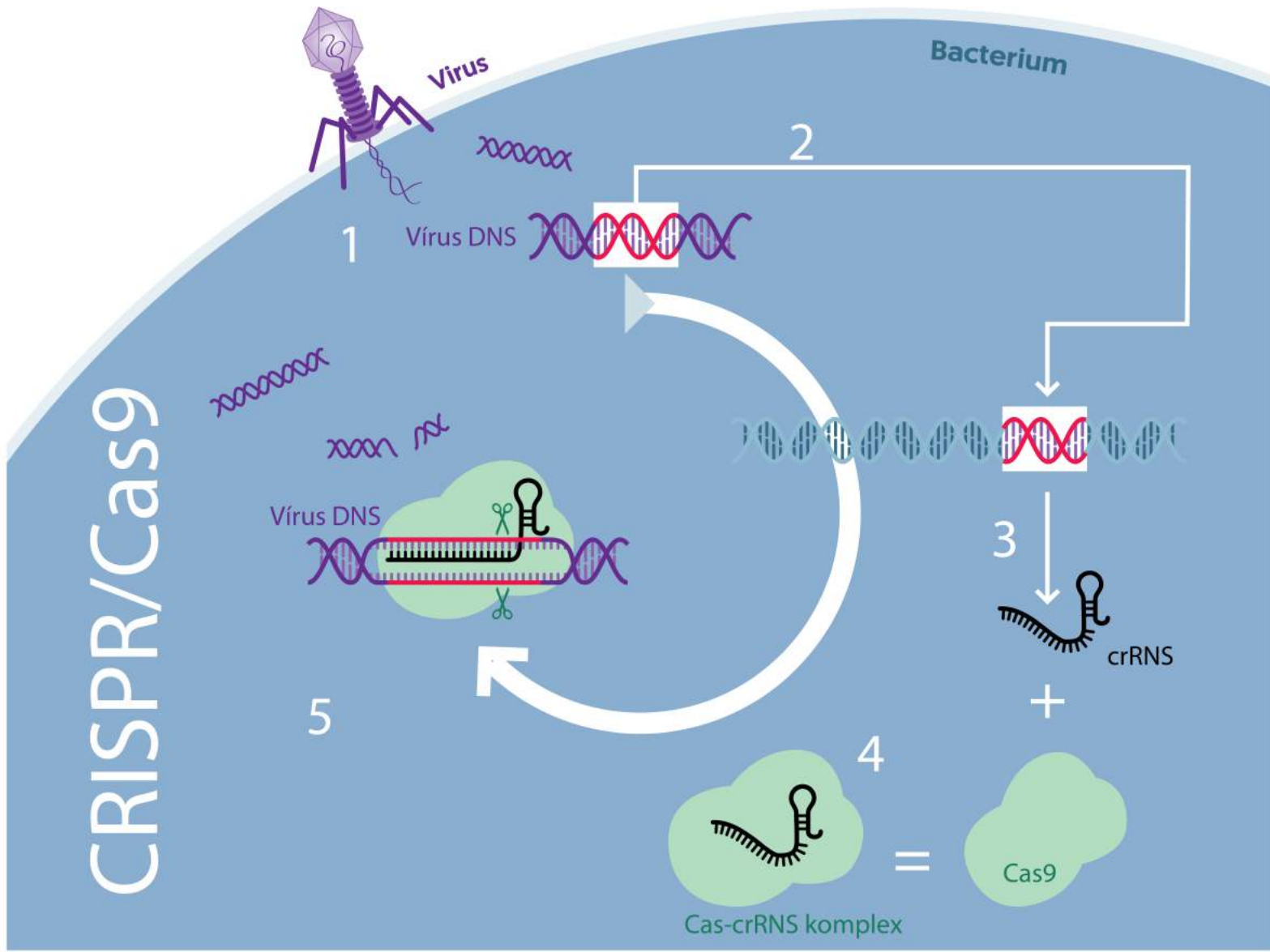
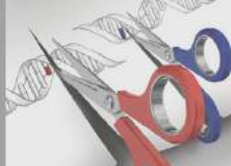


c

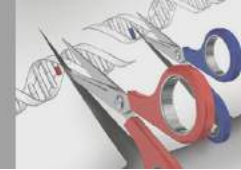
Founders	H	T	D	D	V	S	S	S	E	E	R	E	G	K	V	K	I	Amino acid sequence			
	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TCA	GAG	GAG	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	Wild-type sequence	
#1	T	CAT	ACA	GAT	GAT	<u>ac</u> -	---	---	---	---	---	---	--A	GGT	AAA	GTC	AAA	ATC	A	Δ21 bp + aa change	
	T	CAT	ACA	GgT	GAT	G--	---	---	---	---	---	---	--AA	GGT	AAA	GTC	AAA	ATC	A	Δ21 bp + aa change	
#3	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TCA	GAG	---	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	Δ3 bp	
	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TCA	G--	---	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	Δ5 bp (frameshift)	
#4	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TCA	G-G	GAG	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	Δ1 bp (frameshift)	
	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TCA	GAG	GAG	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	WT	
#5	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TC-	---	-AG	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	Δ5 bp (frameshift)	
	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TCA	GAG	GAG	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	WT	
#6	T	CAT	ACA	GAT	GAT	GTC	TC-	---	---	---	---	---	--G	CGA	GAA	GGT	AAA	GTC	AAA	ATC	Δ12 bp
	T	CAT	ACA	GAT	GAT	G-	---	---	---	---	---	---	--AA	GGT	AAA	GTC	AAA	ATC	A	Δ21 bp	
#8	T	CAT	ACA	GAT	GA-	---	---	---	---	---	---	---	--A	GGT	AAA	GTC	AAA	ATC	A	Δ24 bp	
	T	CAT	ACA	GAT	GAT	GTC	Tac	aga	t--	---	---	---	GAA	GGT	AAA	GTC	AAA	ATC	A	Δ11 bp (frameshift)	
#10	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TCA	G-G	GAG	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	Δ1 bp (frameshift)	
	T	CAT	ACA	GAT	GAa	---	---	---	---	---	---	---	---	---	AAA	GTC	AAA	ATC	A	Δ27 bp + aa change	
#11	T	CAT	ACA	GAT	GAT	GTC	TCA	TCA	TCA	G--	GAG	CGA	GAA	GGT	AAA	GTC	AAA	ATC	A	Δ2 bp (frameshift)	
	T	CAT	ACA	GAT	GA-	---	---	---	---	---	---	---	--A	GGT	AAA	GTC	AAA	ATC	A	Δ24 bp	

(Sung et al., 2013, Nat Biotech)

DNS-specifikus kötés: Cas nukleázok



CRISPR = clustered regularly interspaced palindromic repeats



NOBELPRISET I KEMI 2020 THE NOBEL PRIZE IN CHEMISTRY 2020



KUNGL.
VETENSKAPS-
AKADEMIEN
THE ROYAL SWEDISH ACADEMY OF SCIENCES

Photo: Hallbauer&Floerth



Emmanuelle Charpentier

Born in France, 1968

Max Planck Unit for the Science of
Pathogens, Germany

Photo: UC Berkeley/Doudna Lab

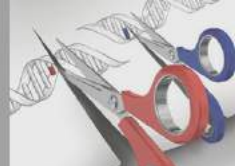


Jennifer A. Doudna

Born in the USA, 1964

University of California, Berkeley, USA
Howard Hughes Medical Institute

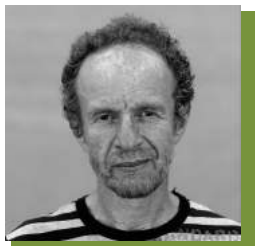
CRISPR alapú gén-editálás – a felfedezők



Yoshizumi Ishino



Francisco Mojica



Eugene Koonin



Rodolphe Barrangou



Virginijus Siksnys

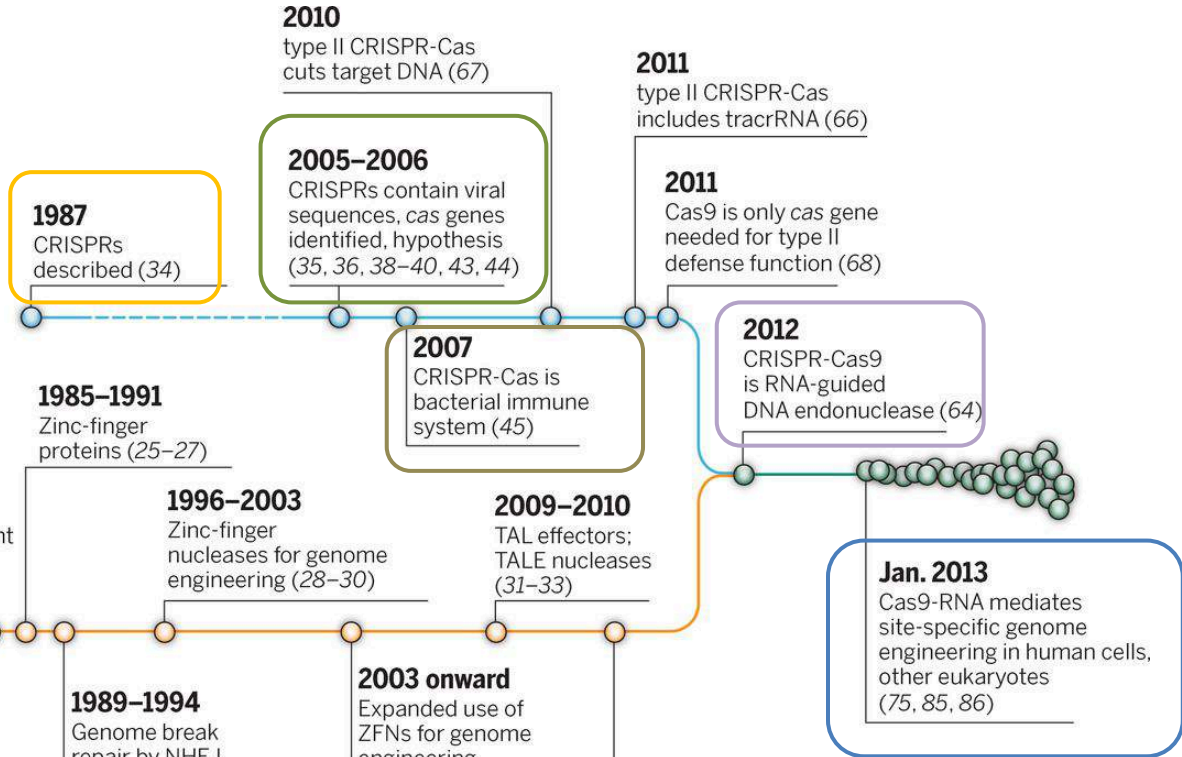


Feng Zhang



George Church

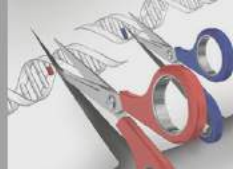
CRISPR biology



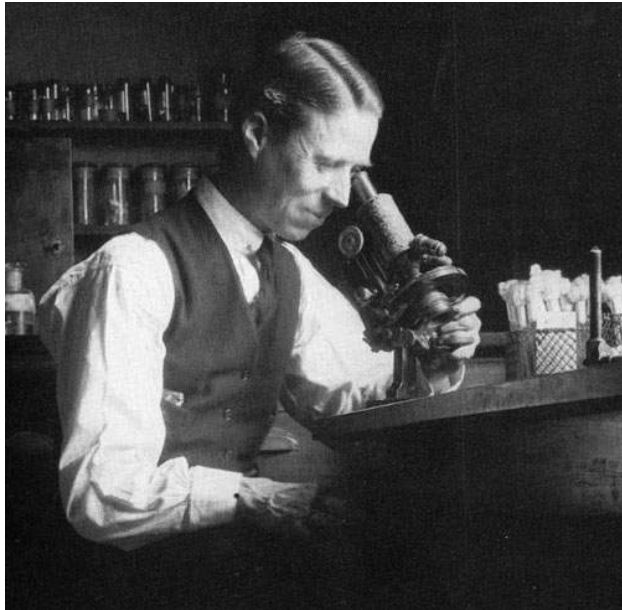
Genome editing

(Doudna and Charpentier, 2014, *Science*)

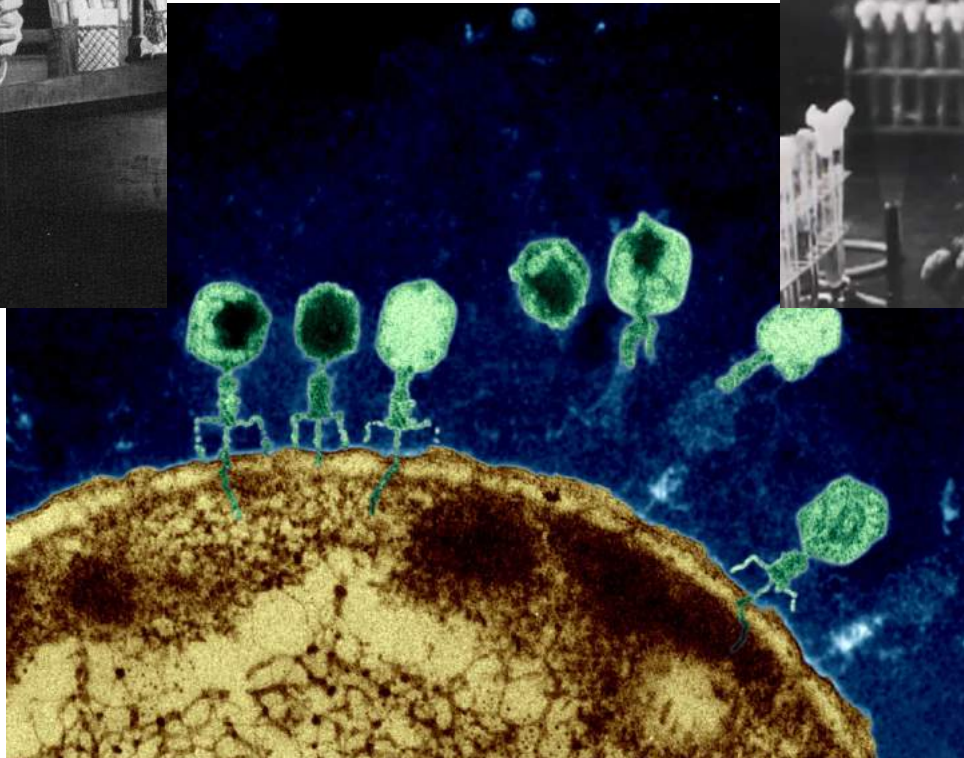
Egy évmilliárdos fegyverkezési verseny



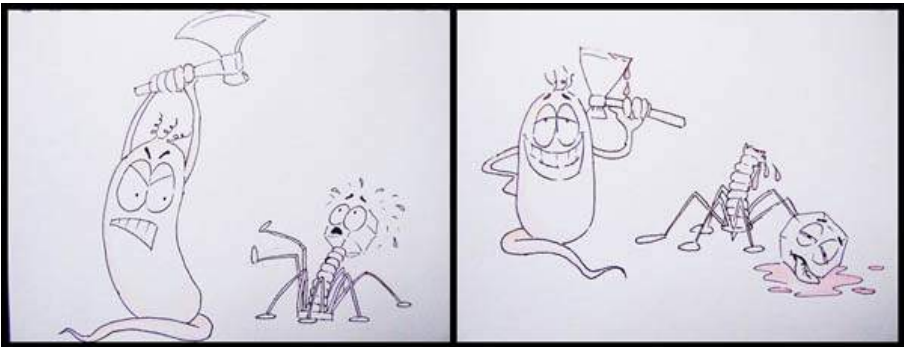
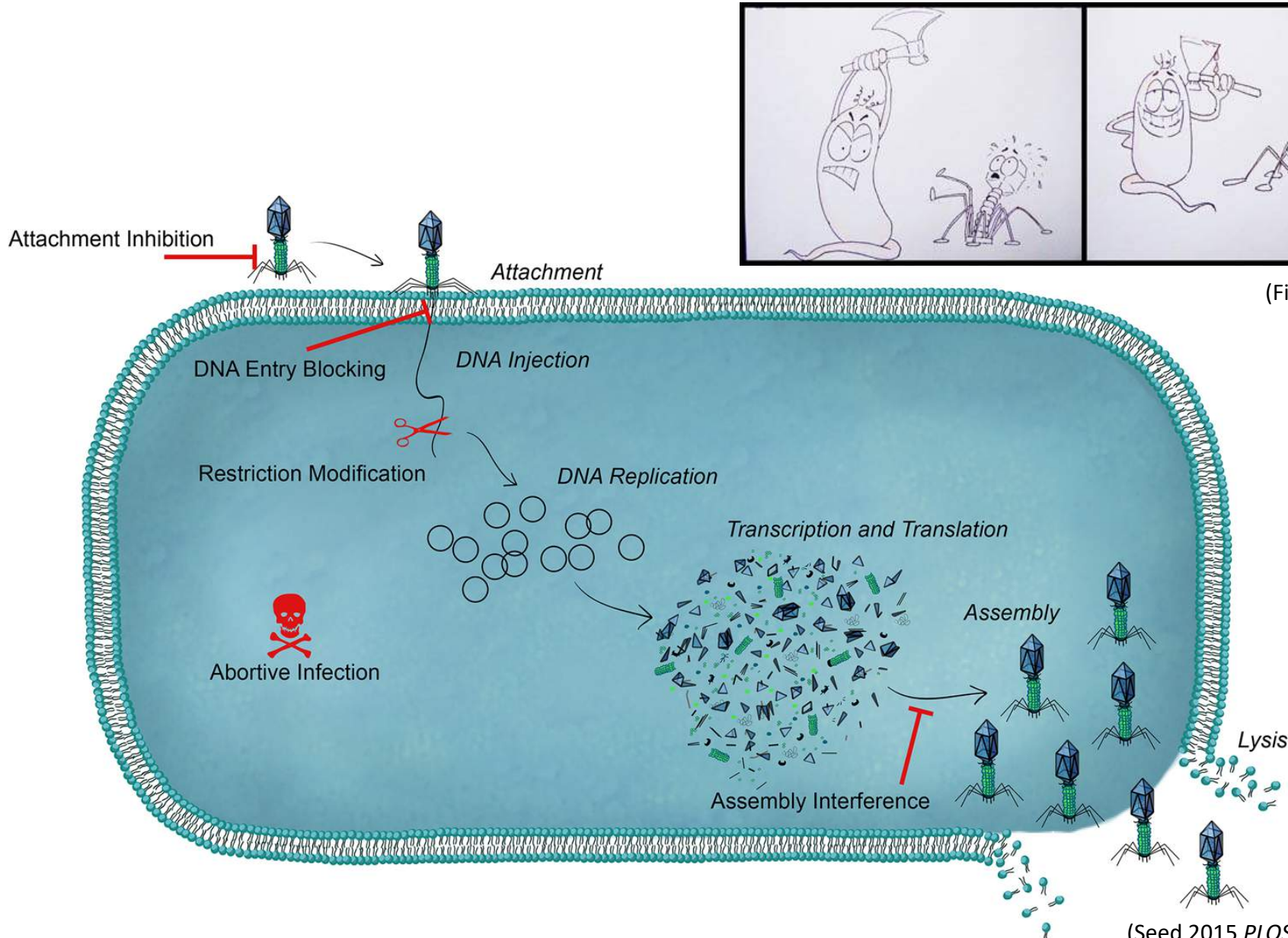
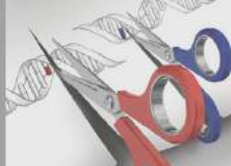
1915 – Frederick Twort:
“Egy tenyésztető enzim”



1917 – Felix d’Herelle:
“Bakteriofágok – baktériumok elfogyasztói”

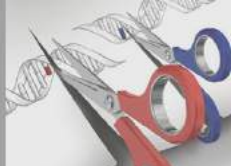


Fág-ellenes stratégiák baktériumokban

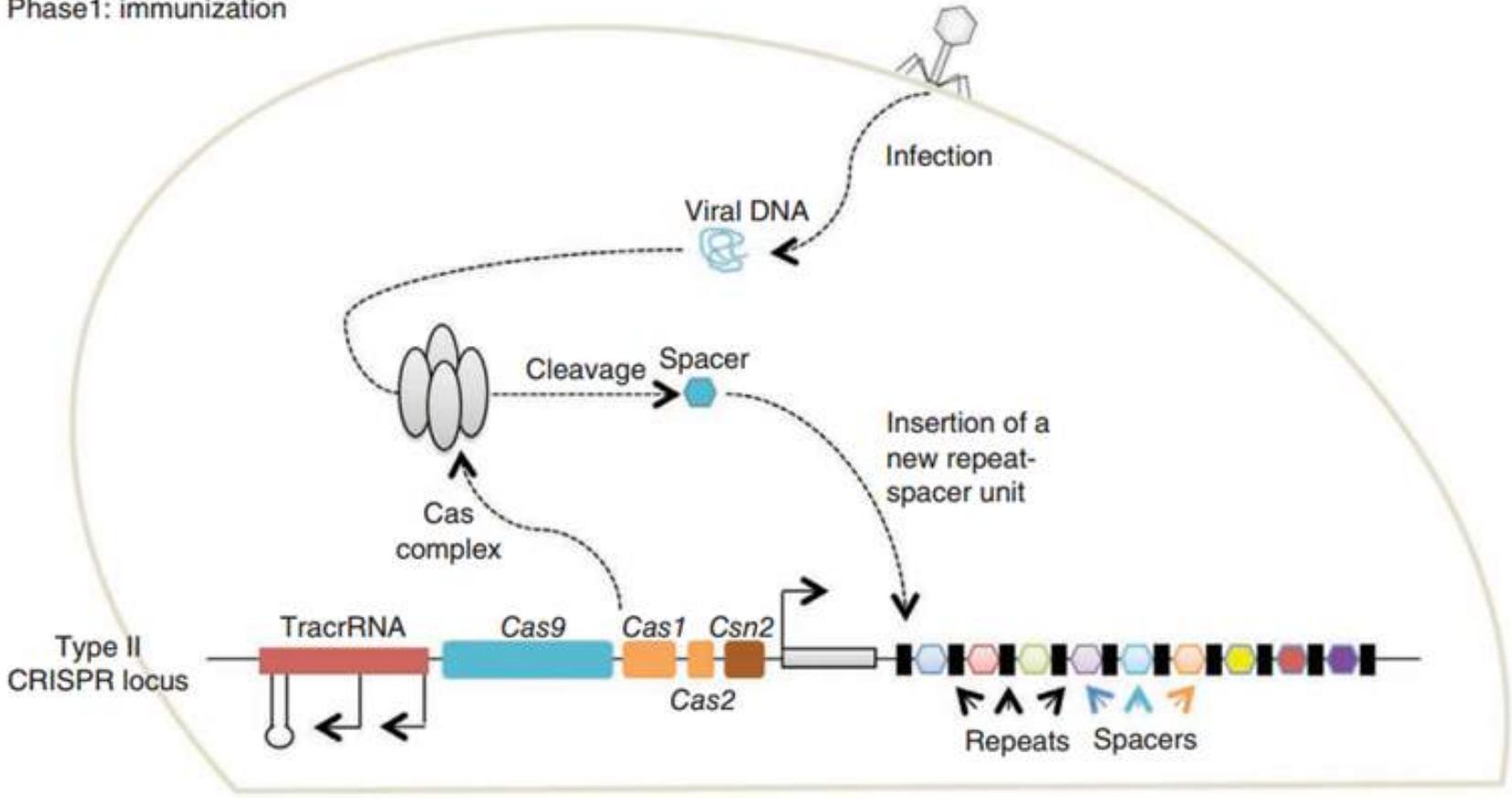


(Fineran labor)

Fág-ellenes stratégiák: prokarióta “immunrendszer”?

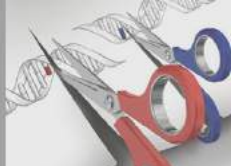


Phase1: immunization

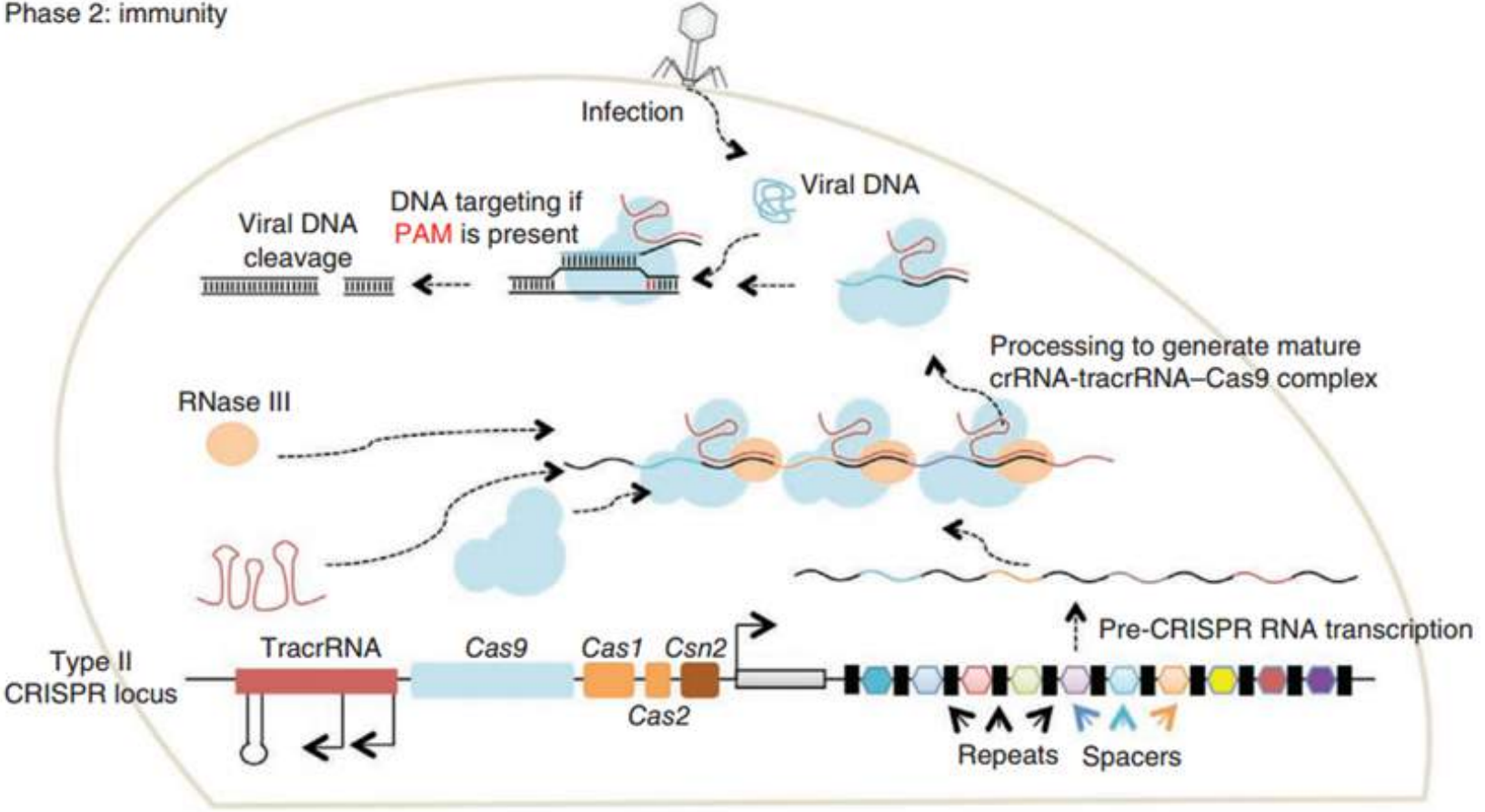


CRISPR = clustered regularly interspaced palindromic repeats

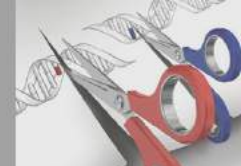
Fág-ellenes stratégiák: prokarióta “immunrendszer”?



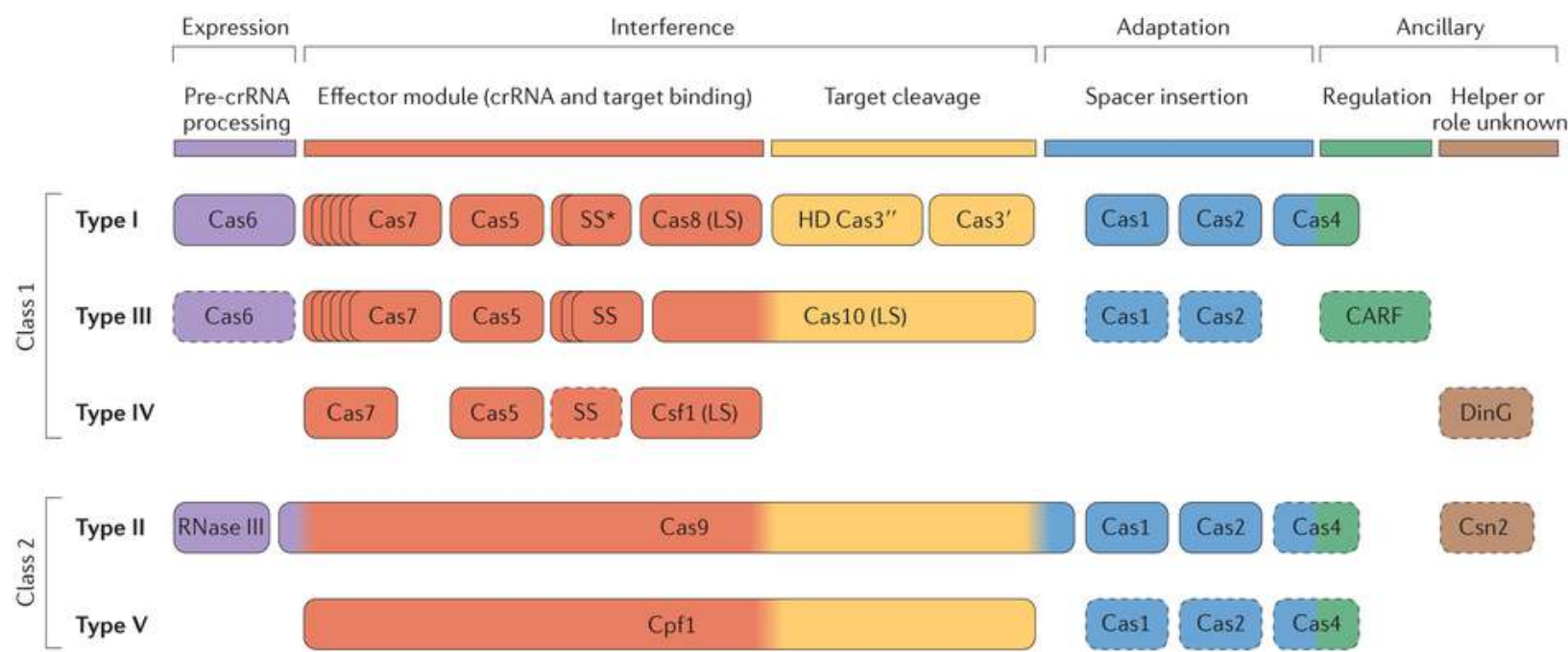
Phase 2: immunity



CRISPR = clustered regularly interspaced palindromic repeats

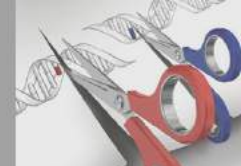


A főbb ismert CRISPR rendszerek

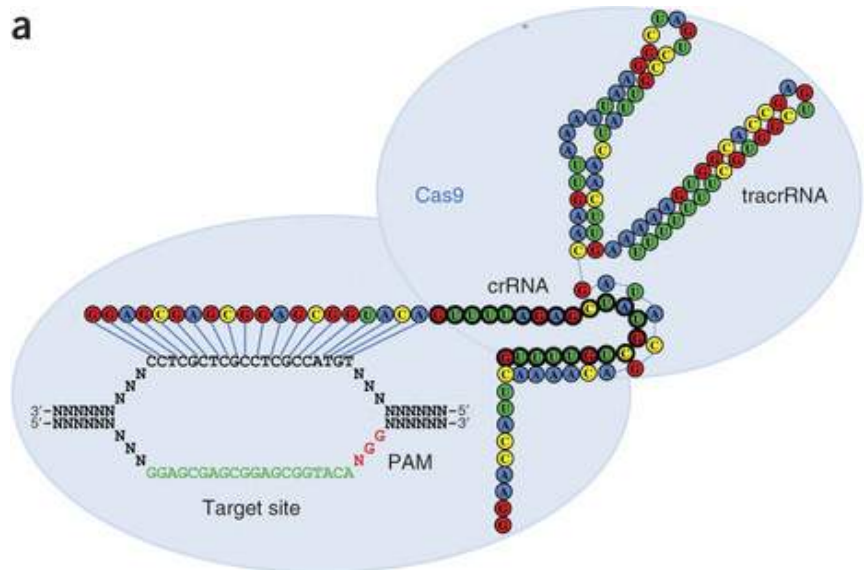


Nature Reviews | Microbiology

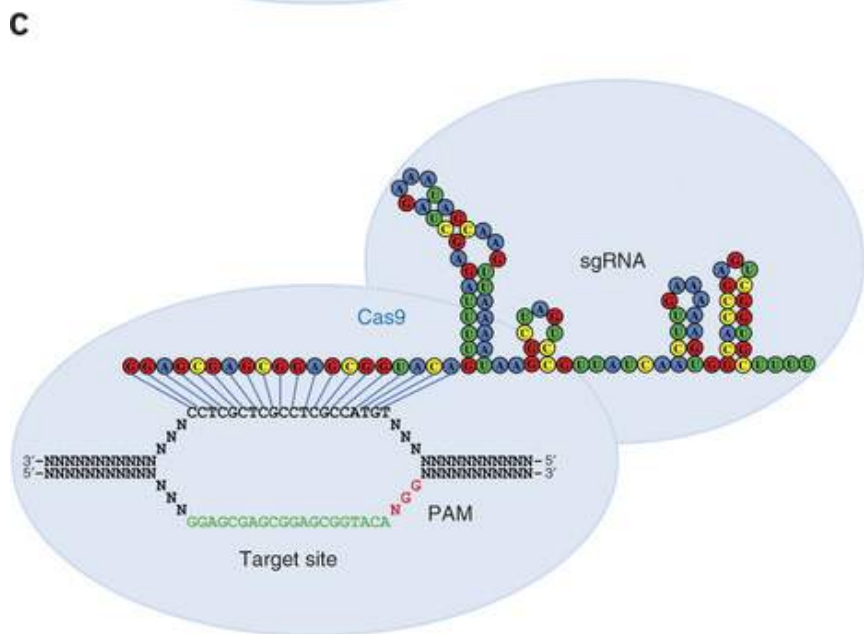
(Makarova et al. 2015 *Nat Rev Micro*)



CRISPR alapú gén-editálás



crRNA = CRISPR targeting RNA
 tracrRNA = trans activating crRNA
 PAM = protospacer adjacent motif
 sgRNA = single guide RNA



d

tia11
 Mutations in 17 out of 44 sequenced alleles

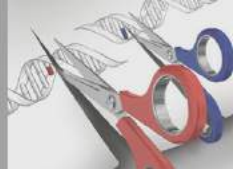
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CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCTCCAGGGATGTTACGGAGGCCCT		Wild-type
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTGGGGATGTCGGGAATCCAGGGAT	+14	(-1, +15)
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTAGGGATGTTACGGAGG	+4	(-7, +11)
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTACCACTCCAGGGATGTTACGGAGGC	+3	(-3, +6)
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCAGGGATGTTACGGAGGCCCT	-2	
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCAGGGATGTTACGGAGGCCCT	-3	(-4, +1)
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCAGGGATGTTACGGAGGCCCT	-4	
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCAGGGATGTTACGGAGGCCCT	-5	
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCAGGGATGTTACGGAGGCCCT	-6	
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCAGGGATGTTACGGAGGCCCT	-11	
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCAGGGATGTTACGGAGGCCCT	-13	
CCTGTGCTCTCCTGTTTTTAGGATATGTCGGGAACCTCAGGGATGTTACGGAGGCCCT	-23	

gsk3b
 Mutations in 8 out of 16 sequenced alleles

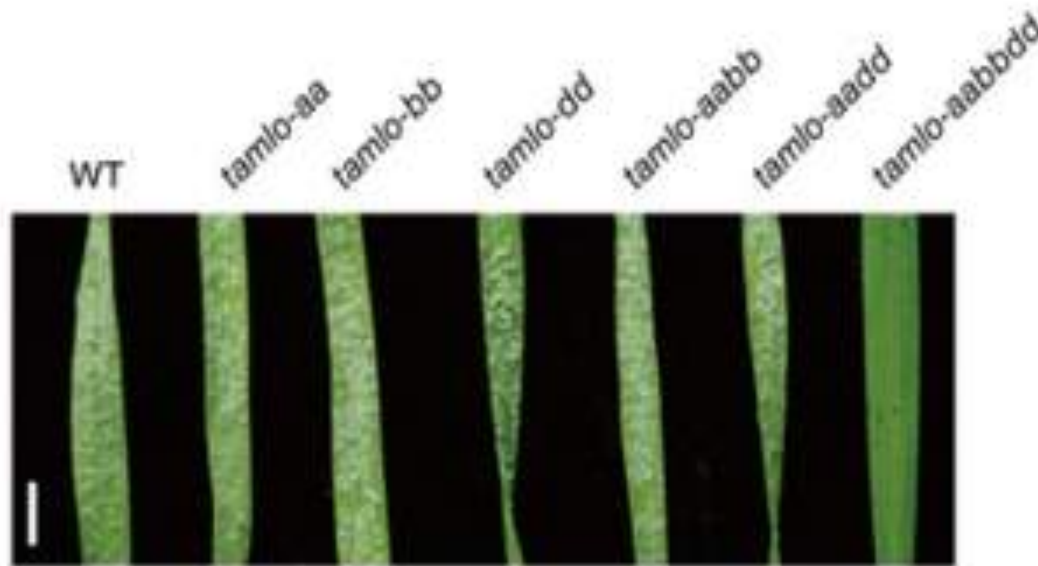
Sequence	Position	Wild-type
GTGGTGGCGACTCCTGGACAGGGACCTGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC		Wild-type
GTGGTGGCGACTCCTGGACAGGGACCTGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC	+17	(-8, +25)
GTGGTGGCGACTCCTGGACAGGGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC	+2	(-4, +6)
GTGGTGGCGACTCCTGGACAGGGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC	+1	(-2, +3)
GTGGTGGCGACTCCTGGACAGGGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC	+1	(-8, +9)
GTGGTGGCGACTCCTGGACAGGGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC	-7	
GTGGTGGCGACTCCTGGACAGGGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC	-10	
GTGGTGGCGACTCCTGGACAGGGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC	-11	
GTGGTGGCGACTCCTGGACAGGGACCTGACCGCCCGCAGGAGGTACAGTACACTGACACC	-13	

(Hwang et al., 2013, Nat Biotech)

Genomszerkesztés növényekben

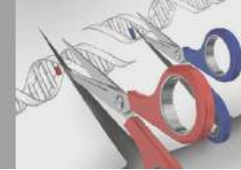


- a lisztharmatfertőzés feltételezi a működőképes MLO fehérje jelenlétét
- búzában (hexaploid) három pár *MLO* homoeoallél is van, ez nehezíti a védekezést



MLO homoeoallélok mutagenézise hexaploid búzában (rezisztencia érhető el így)

Természetes vs. mesterséges funkcióvesztés: mi a különbég...?



“Texel birka”

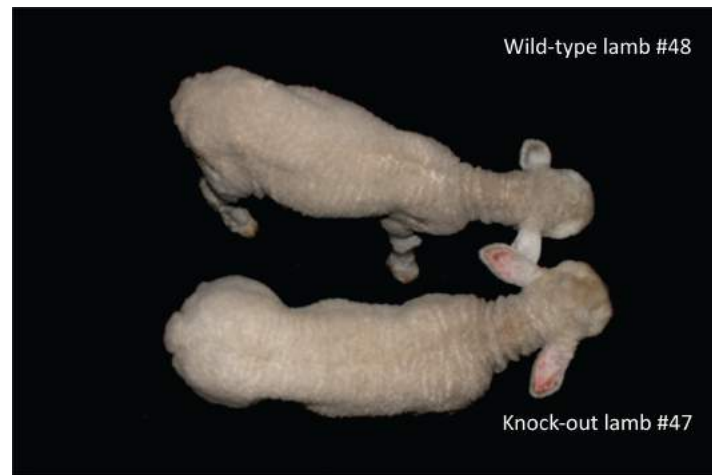


“Bully” agár



“Belgian blue”

“Természetes” *MSTN* mutánsok



Wild-type lamb #48

Knock-out lamb #47

(Crispo et al 2015 *PLOS One*)



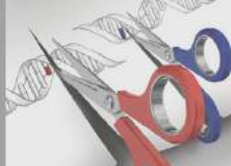
(Zou et al. 2015 *J Mol Cell Biol*)



(Wang et al. 2015 *SciRep*)

“Mesterséges” *MSTN* mutánsok

Disznó-eredetű xenotranszplantáció



CHOICE CUTS

Researchers are looking to source an increasing variety of living tissues, including solid organs, from pigs. Many are attempting to genetically engineer the animals to reduce the risk of rejection and infection in humans.

CORNEA

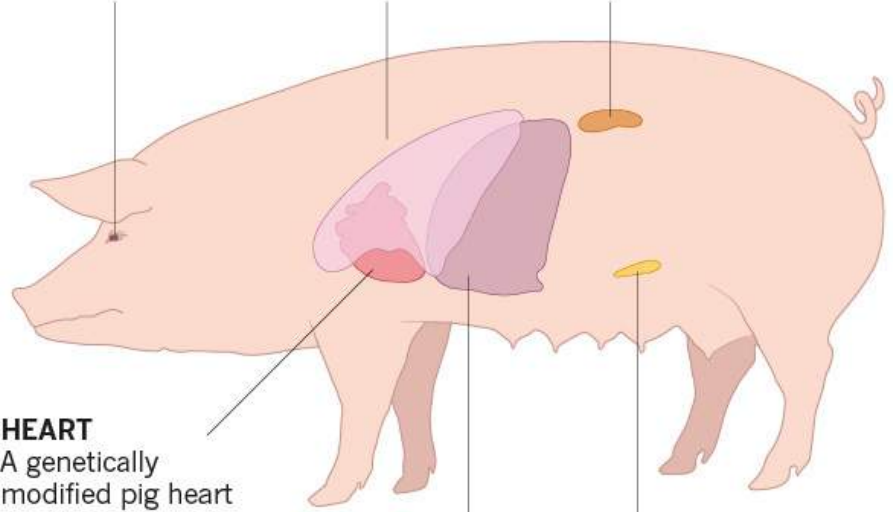
Pig corneas were approved for marketing in China in April.

LUNG

A factory farm is being designed to produce 1,000 pig lungs per year.

KIDNEY

A kidney with six genetic modifications supported a baboon's life for 4 months.



HEART

A genetically modified pig heart implanted in a baboon's abdomen survived for 2.5 years.

LIVER

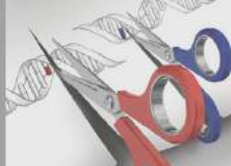
Livers could be engineered to produce their own antibodies against primate immune cells.

PANCREAS

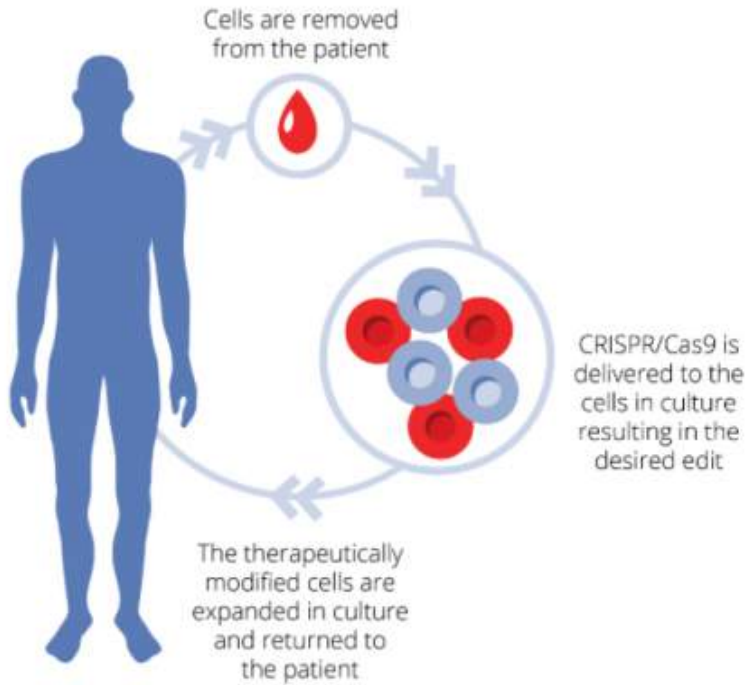
Phase III clinical trials of insulin-producing islet cells are under way.

- George Church csoportja képes volt az összes sertés endogén retrovírus (PERV) kópiát inaktiválni – számszerint 62 db-t
- további 20, sejtfelszíni immunreakciót kiváltó fehérjét kódoló gént is elrontottak

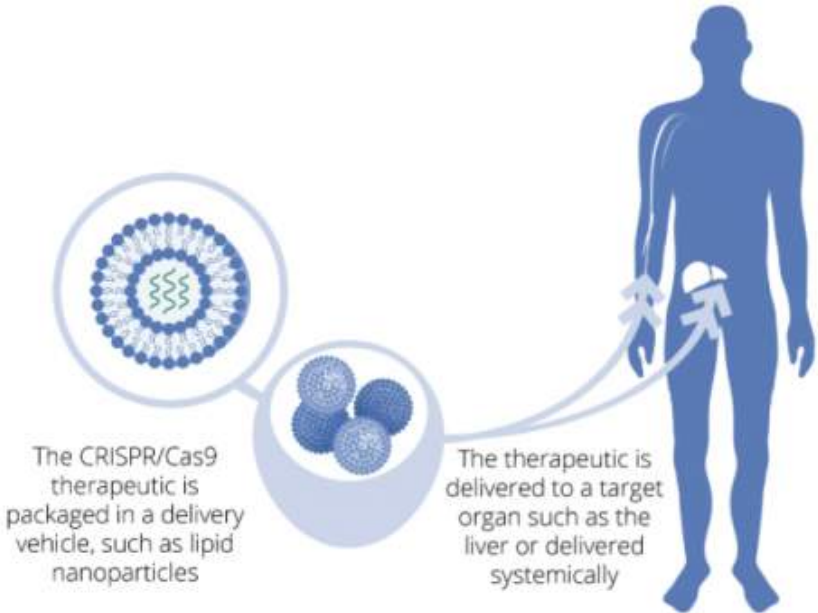
Genomeditálás-alapú terápiák



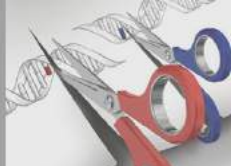
Ex vivo



In vivo



Genomedítálás-alapú terápiák




EDITING INSIDE THE BODY IN VIVO CRISPR MEDICINES


EDITING OUTSIDE THE BODY ENGINEERED CELL MEDICINES

OCULAR DISEASES

CANCER

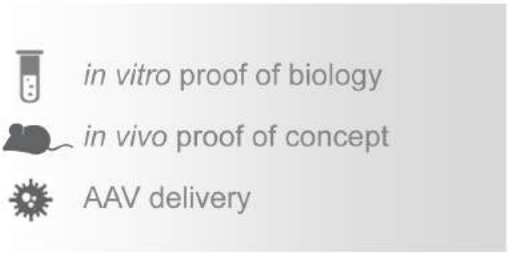
Autologous T cell medicines** 
Allogeneic cell medicines




BLOOD DISEASES

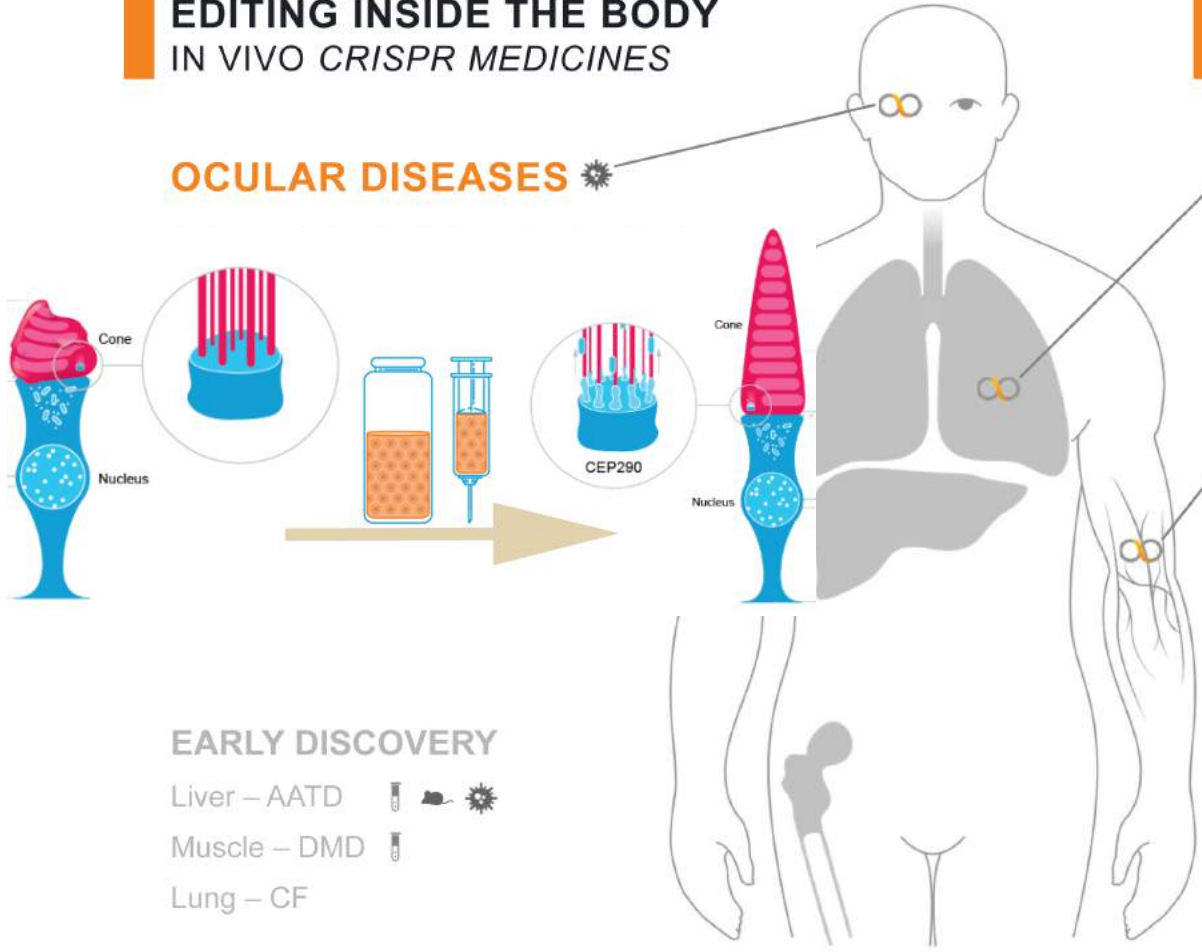
Sickle cell disease  
Beta-thalassemia  

EARLY DISCOVERY

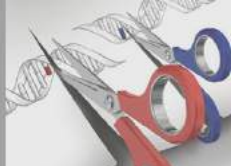
Liver – AATD   
Muscle – DMD 
Lung – CF



 *in vitro* proof of biology
 *in vivo* proof of concept
 AAV delivery



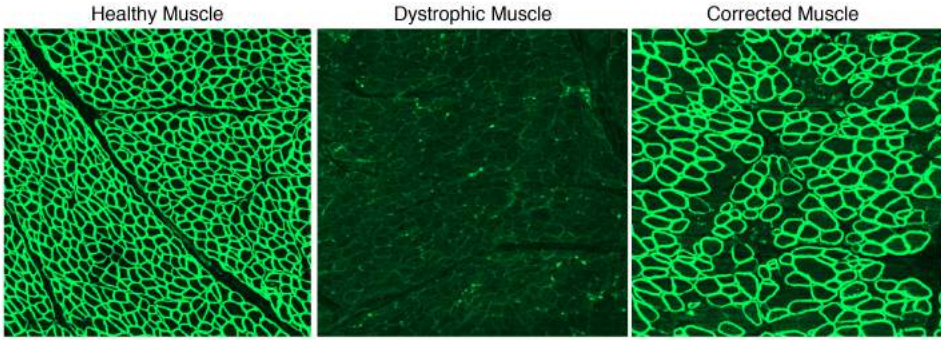
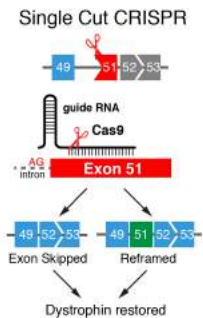
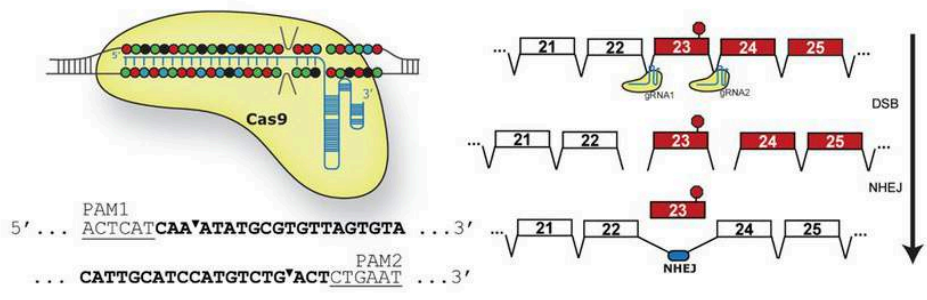
Genomedítálás-alapú terápiák (in vivo)



Duchenne's Impact on the Body

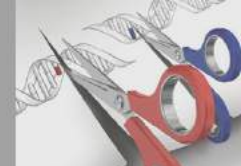
- DYSTROPHIN ABNORMALITIES**
- POSSIBLE LEARNING AND COGNITIVE DIFFICULTIES**
- DECREASED HEART FUNCTION**
- CARDIOMYOPATHY**
- LEADS TO HEART FAILURE**
- WEAKENS DIAPHRAGM**
- REQUIRES VENTILATOR**
- LEADS TO PNEUMONIA**
- LOSS OF MUSCLE MASS**
- WEAKNESS**
- INFLAMMATION**
- FIBROSIS**
- BRITTLE AND WEAK**

Source: CureDuchenne™



Kutya – DMD model

(Amoasii et al. 2018 Science)



Szép új világot építünk... (már megint)?

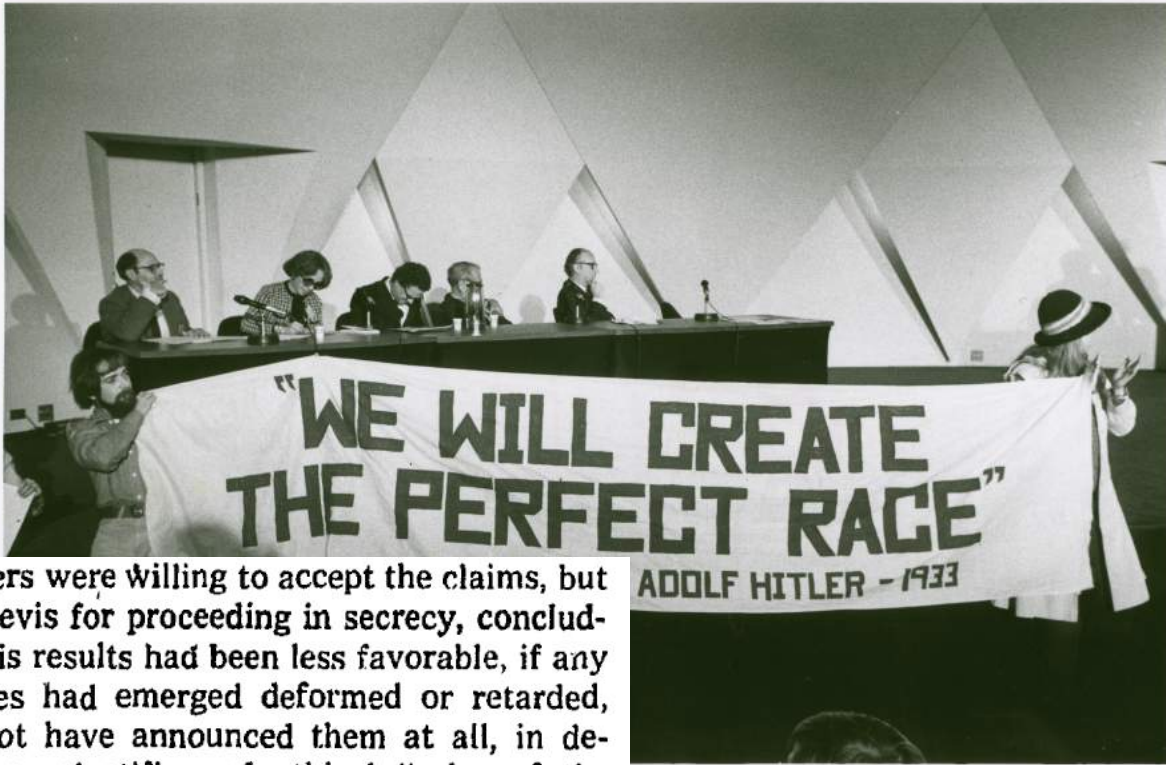
LETTERS

Potential Biohazards of Recombinant DNA Molecules

Recent advances in techniques for the isolation and rejoining of segments of DNA now permit construction of biologically active recombinant DNA molecules in vitro. For example, DNA restriction endonucleases, which generate DNA fragments containing co-

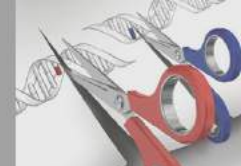
Several groups of scientists are now planning to use this technology to create recombinant DNA's from a variety of other viral, animal, and bacterial sources. Although such experiments are likely to facilitate the solution of important theoretical biological problems, also result in the creation of infectious DNA whose biological properties are completely predicted in

(Berg et al. 1974 PNAS)

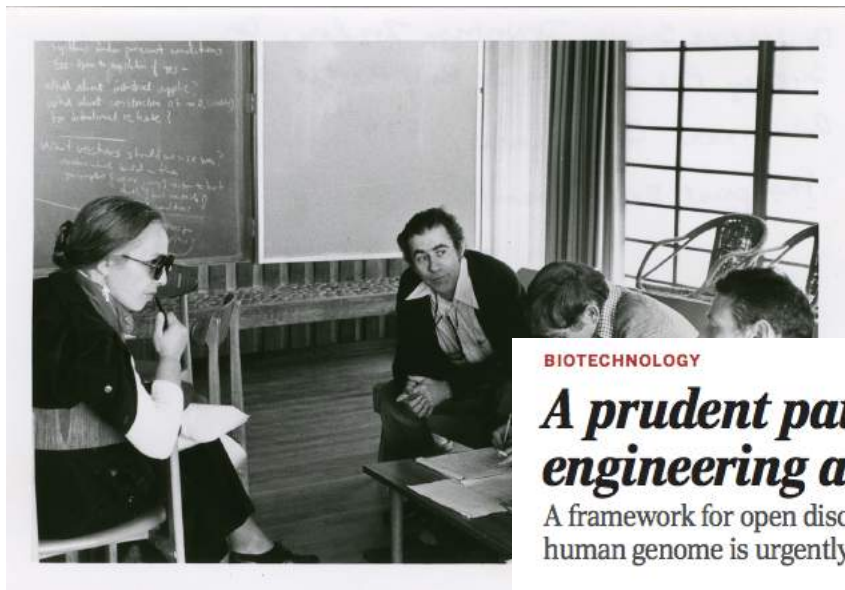


Some others were willing to accept the claims, but castigated Bevis for proceeding in secrecy, concluding that if his results had been less favorable, if any of the babies had emerged deformed or retarded, he would not have announced them at all, in defiance of the scientific and ethical "rules of the game." Still others overlooked the issue of credibility and, instead, shuddered at the implications of the claims. Not long ago, Dr. James Watson, the Nobel biologist, told a Congressional subcommittee that a successful embryo transplant would soon occur. Watson foresaw "all sorts of bad scenarios" from the achievement. "All hell will break loose, politically and morally, all over the world," he concluded, a bit rashly, some felt.

(15 Sept 1974 NYTimes)



Asilomar árnyékában



- 1975 februárjában a kaliforniai Asilomar-ban gyűltek össze kutatók, hogy egy saját szabályrendszert dolgozzanak ki a rekombináns DNS-el való munkára

BIOTECHNOLOGY

A prudent path forward for genomic engineering and germline gene modification

A framework for open discourse on the use of CRISPR-Cas9 technology to manipulate the human genome is urgently needed

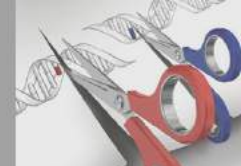
By David Baltimore,¹ Paul Berg,² Michael Botchan,^{3,4} Dana Carroll,⁵ R. Alta Charo,⁶ George Church,⁷ Jacob E. Corn,⁴ George Q. Daley,^{8,9} Jennifer A. Doudna,^{4,10,11} Marsha Fenner,⁴ Henry T. Greely,¹² Martin Jinek,¹³ G. Steven Martin,¹³ Edward Penhoet,¹⁴ Jennifer Puck,¹⁵ Samuel H. Sternberg,¹⁶ Jonathan S. Weissman,^{4,17} Keith R. Yamamoto^{4,18}

Genome engineering technology offers unparalleled potential for modifying human and nonhuman genomes. In humans, it holds the promise of curing genetic disease, while in other organisms it provides methods to reshape the biosphere for the benefit of the environment and human societies. However,



- 2015 január: a kaliforniai Napa-ban Asilomar-redux, ezúttal a CRISPR apropóján

#geneeditsummit



A dizájner bábik hajnala?

RESEARCH ARTICLE

CRISPR/Cas9-mediated gene editing in human tripronuclear zygotes

Puping Liang, Yanwen Xu, Xiya Zhang, Chenhui Ding, Rui Huang, Zhen Zhang, Jie Lv, Xiaowei Xie, Yuxi Chen, Yujing Li, Ying Sun, Yaofu Bai, Zhou Songyang, Wenbin Ma, Canquan Zhou[✉], Junjiu Huang[✉]

Guangdong Province Key Laboratory of Reproductive Medicine, the First Affiliated Hospital, and Key Laboratory of Gene Engineering of the Ministry of Education, School of Life Sciences, Sun Yat-sen University, Guangzhou 510275, China

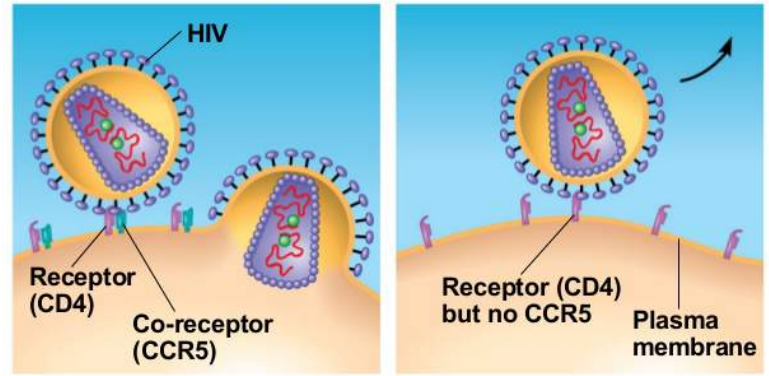
J Assist Reprod Genet
DOI 10.1007/s10815-016-0710-8

TECHNOLOGICAL INNOVATIONS



Introducing precise genetic modifications into human 3PN embryos by CRISPR/Cas-mediated genome editing

Xiangjin Kang¹ · Wenyin He¹ · Yuling Huang¹ · Qian Yu¹ · Yaoyong Chen¹ · Xingcheng Gao¹ · Xiaofang Sun¹ · Yong Fan¹

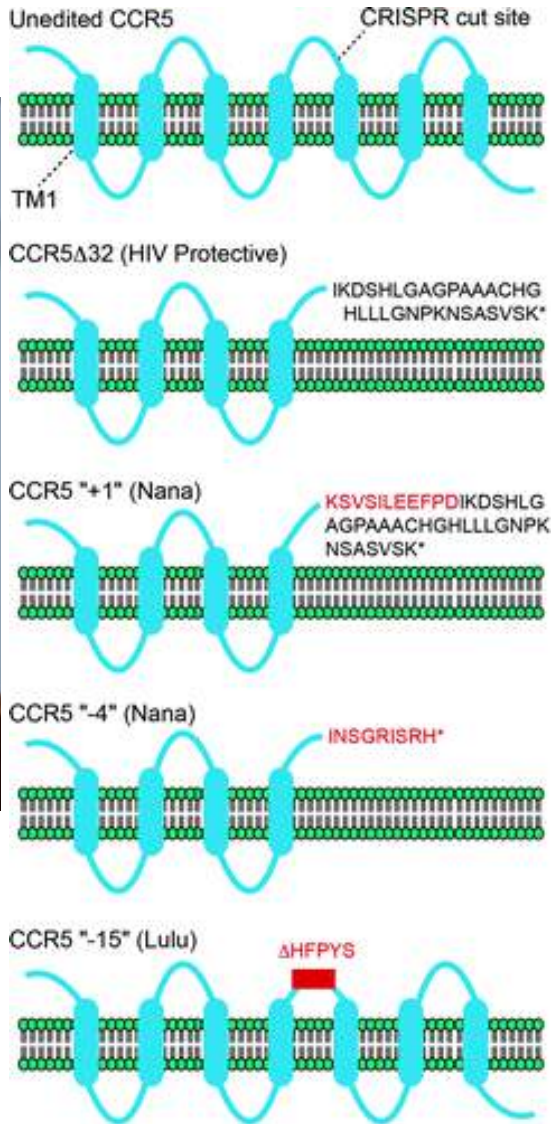
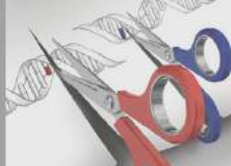


HIV can infect a cell that has CCR5 on its surface, as in most people.

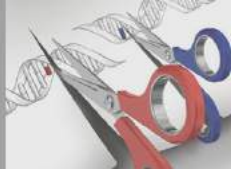
HIV cannot infect a cell lacking CCR5 on its surface, as in resistant individuals.

(Liang et al. 2015 *Protein&Cell*, Kang et al. 2016 *J Assist Reprod Genet*)

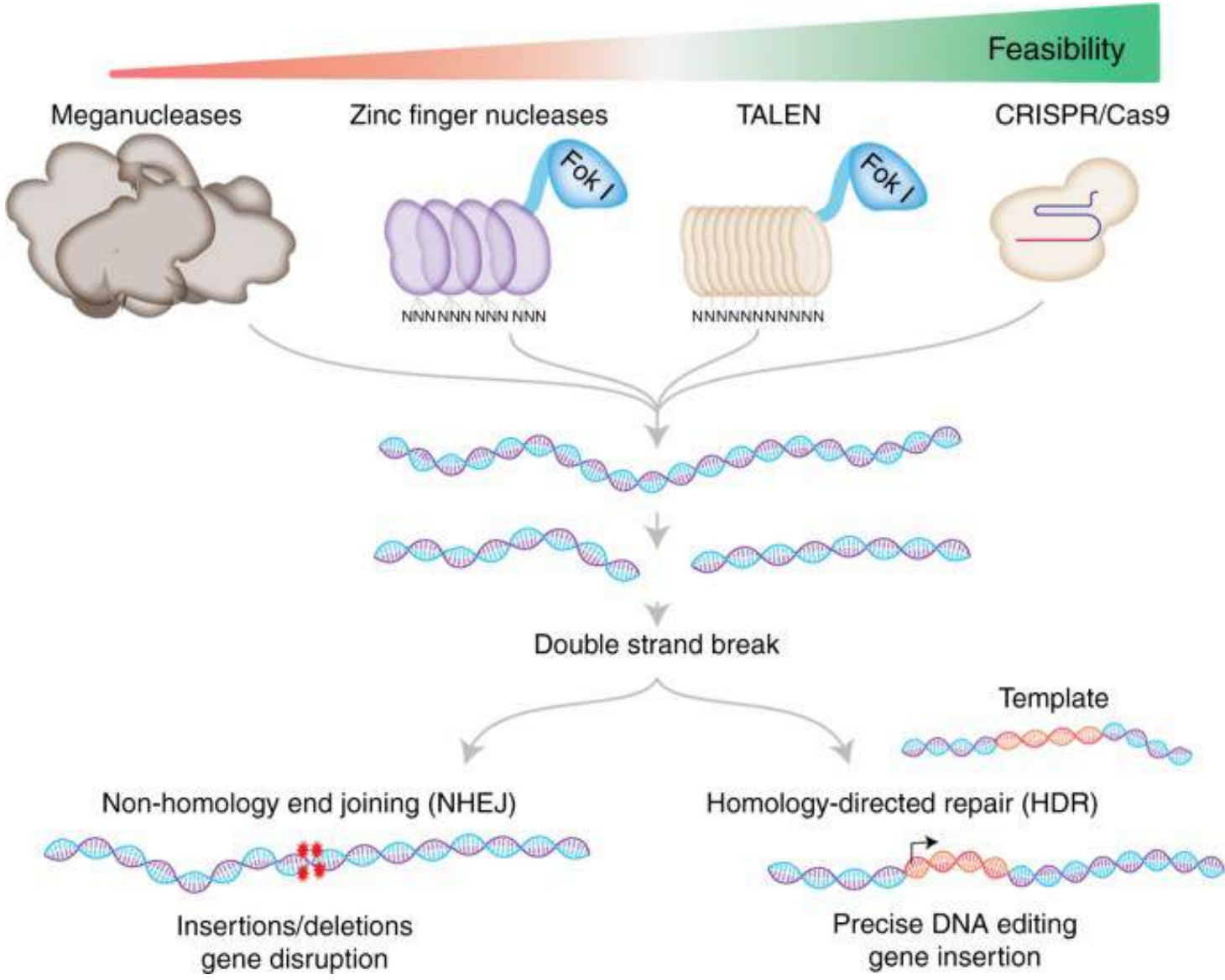
A dizájner bábik hajnala?

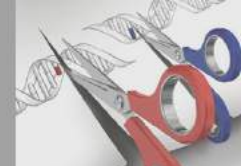


(Csiankuj Ho – 2018 november)

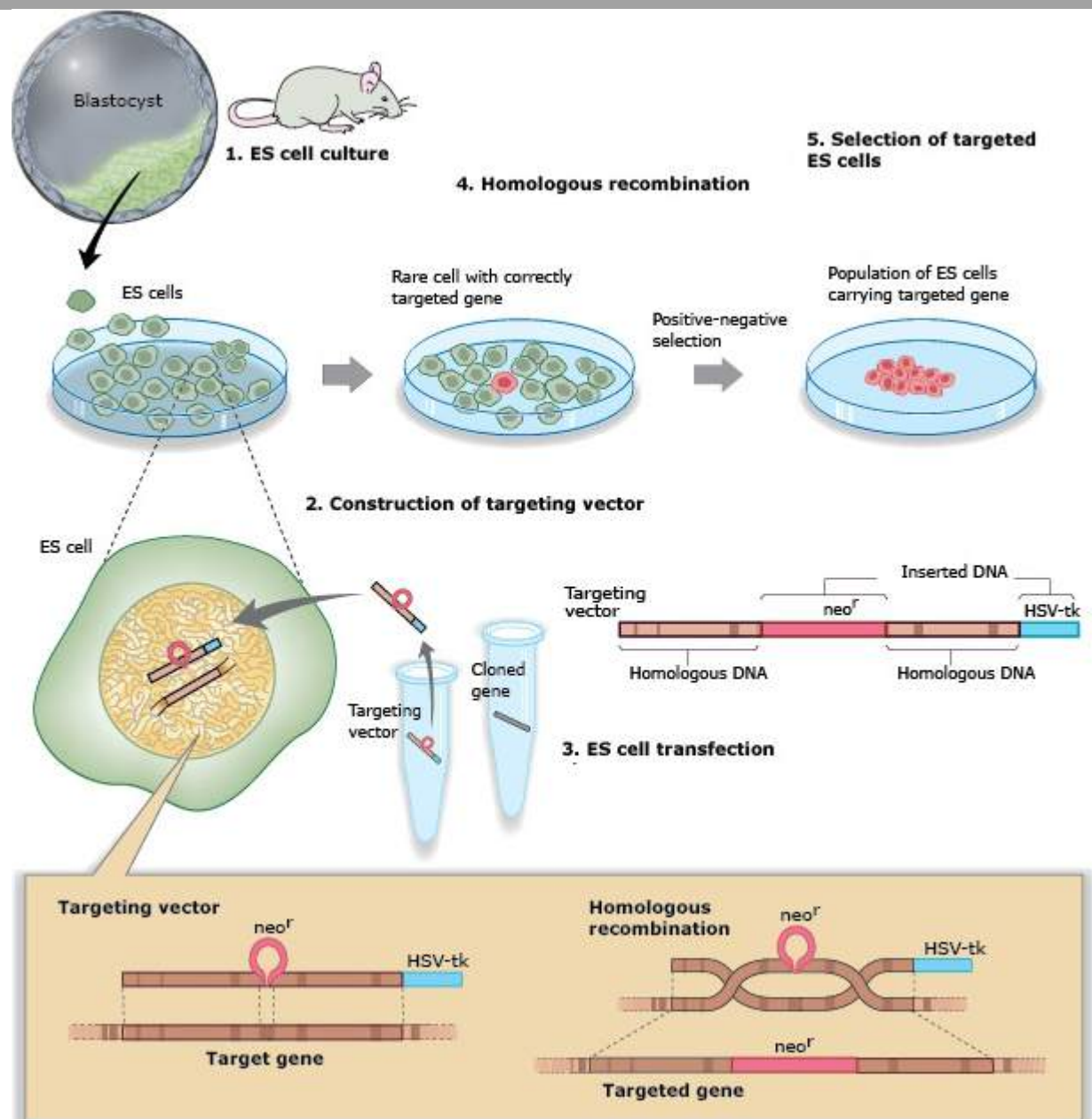


Az igazi „szerkesztés”: homológ rekombináció

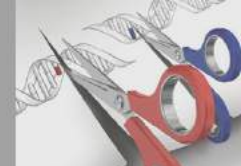




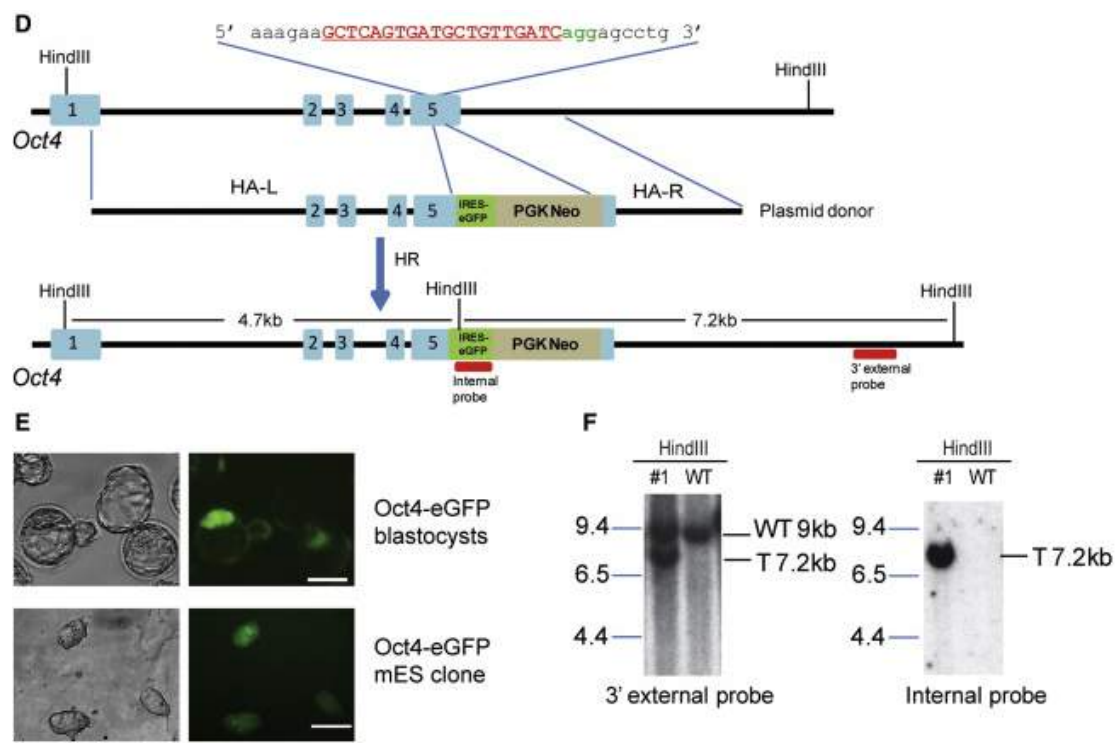
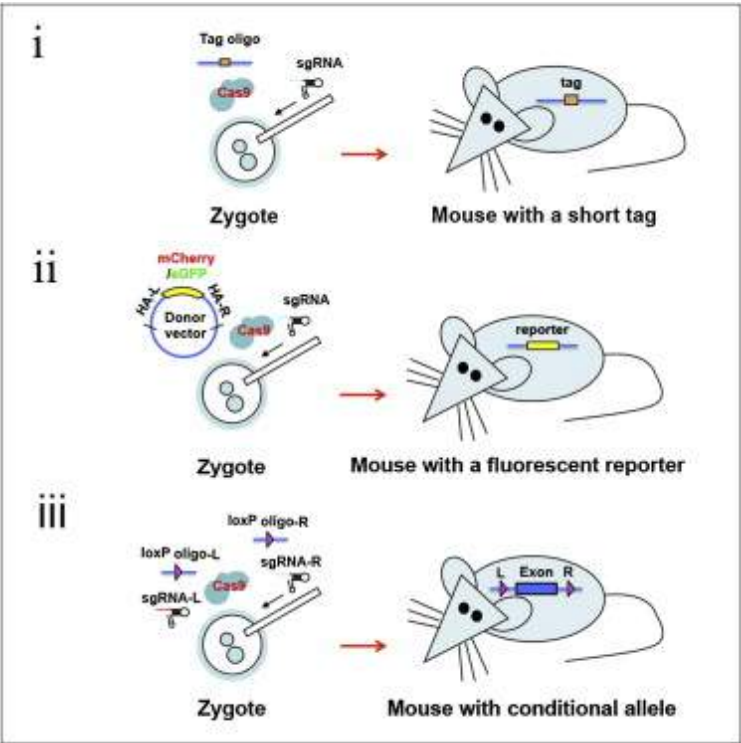
Géeditálás homológ rekombinációval (HR)



- klasszikus HR: 1:10 000 embrió (!!)
- I-SceI emésztés: 50x hatékonyabb
- "klasszikus" Cas9 (mRNS inj): 1:8, Cas9RNP: 1:2



Knock-in egerek készítése CRISPR segítségével

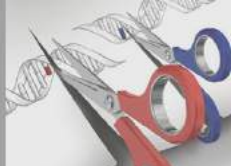


(Yang et al., 2013, *Cell*)

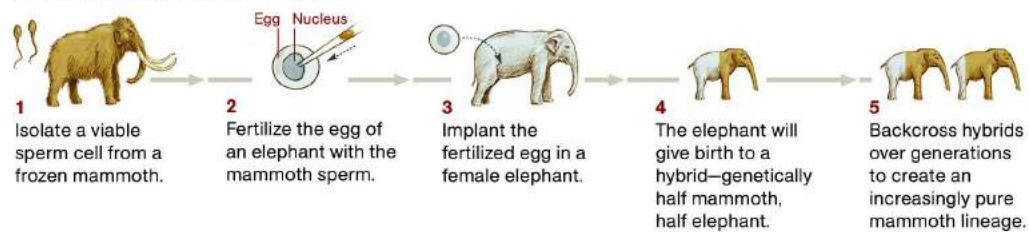
Re-Evaluating One-step Generation of Mice Carrying Conditional Alleles by CRISPR-Cas9-Mediated Genome Editing Technology

The dataset constituted 17,887 microinjected or electroporated zygotes and 1,718 live born mice, of which only 15 (0.87%) mice harbored 2 correct LoxP insertions in cis configuration indicating a very low efficiency of the method.

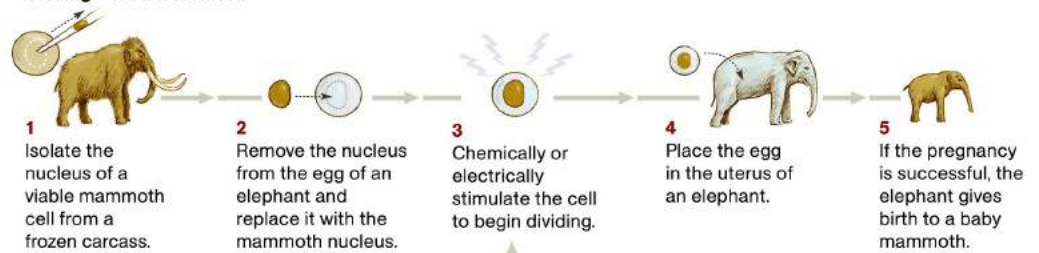
De-extinkció: kihalt fajok “újraélesztése”



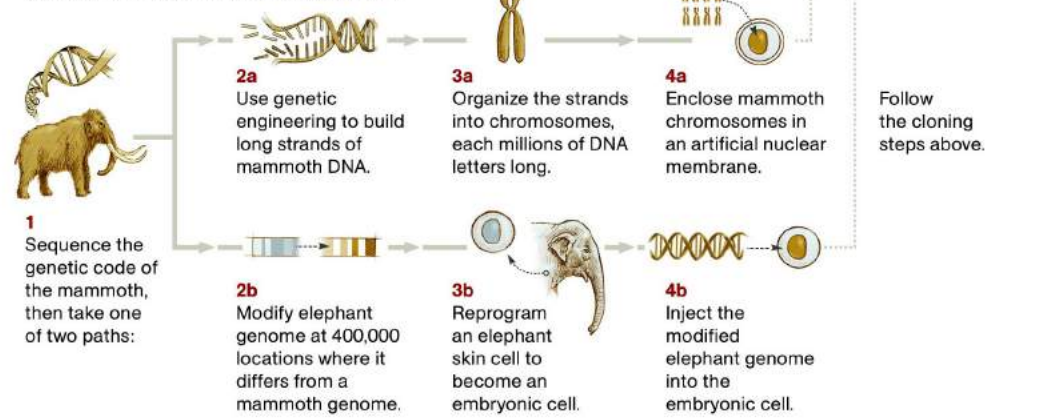
In vitro fertilization from frozen sperm

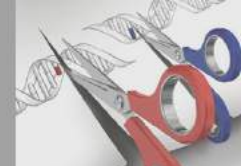


Cloning from a frozen cell

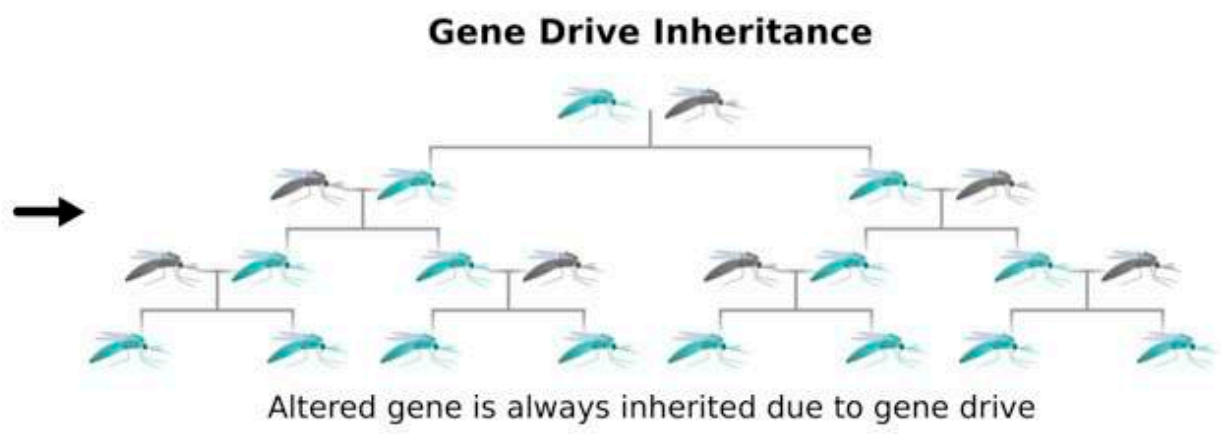
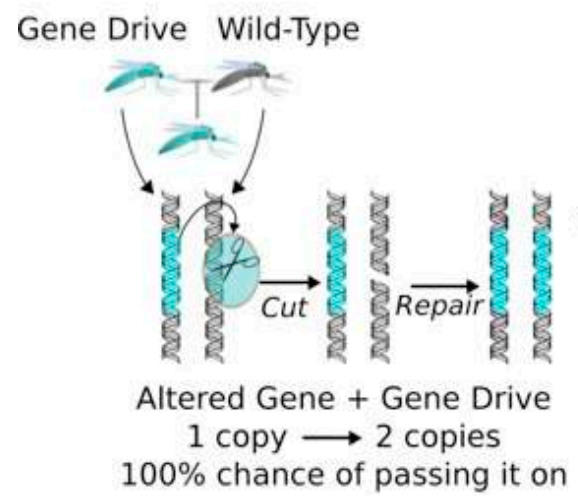
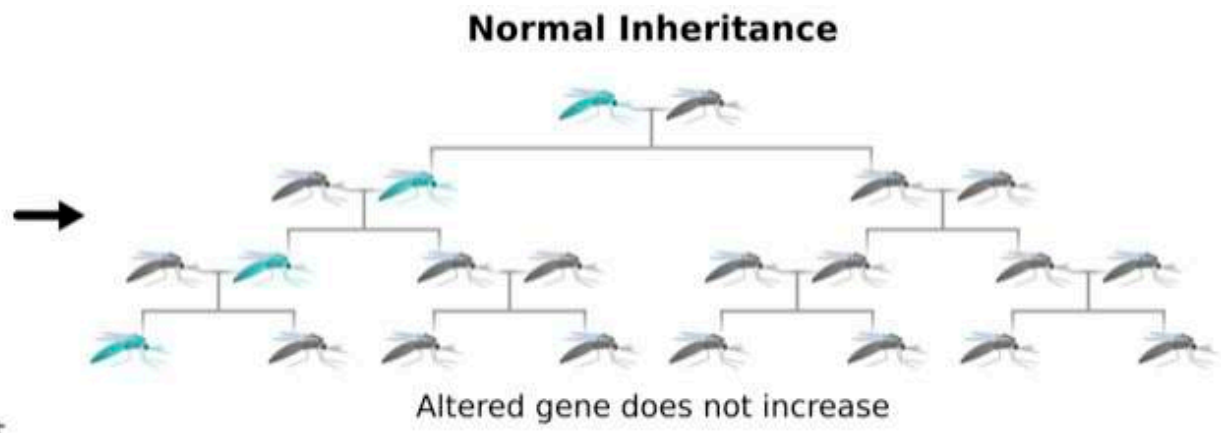
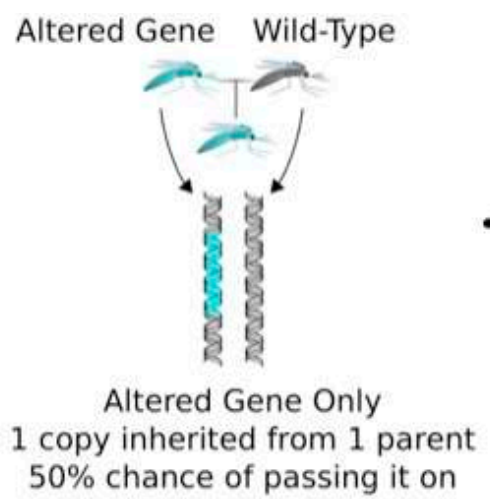


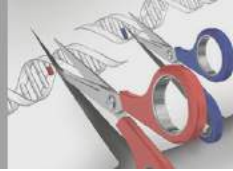
Cloning from sequenced mammoth genome





“Gene drive” – ökológiai mumus, vagy csodaszer





WIRED

The Battle Over Genome Editing Gets Science All Wrong

<http://www.wired.com/2015/10/battle-genome-editing-gets-science-wrong/>

 **THE NEW YORKER**

The Gene Hackers

<http://www.newyorker.com/magazine/2015/11/16/the-gene-hackers>

nature

CRISPR, the disruptor

<http://www.nature.com/news/crispr-the-disruptor-1.17673>

EMBO
reports

No time to waste—the ethical challenges created by CRISPR

<http://embor.embopress.org/content/16/11/1421.long>

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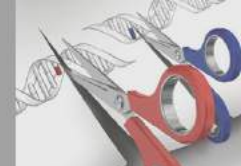
"Gene Drives" And CRISPR Could Revolutionize Ecosystem Management

<http://blogs.scientificamerican.com/guest-blog/gene-drives-and-crispr-could-revolutionize-ecosystem-management/>

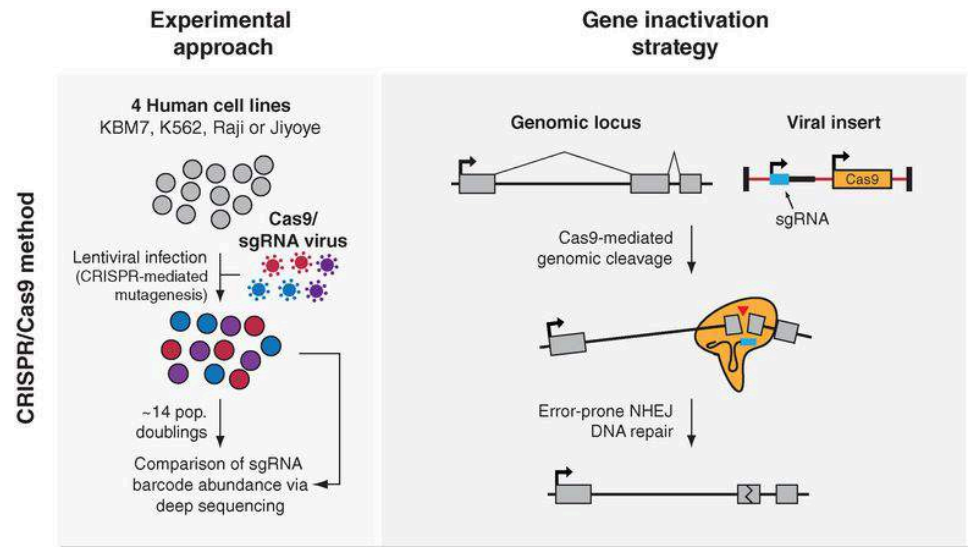
STAT

A debate: Should we edit the human genome?

<http://www.statnews.com/2015/11/30/gene-editing-crispr-germline/>

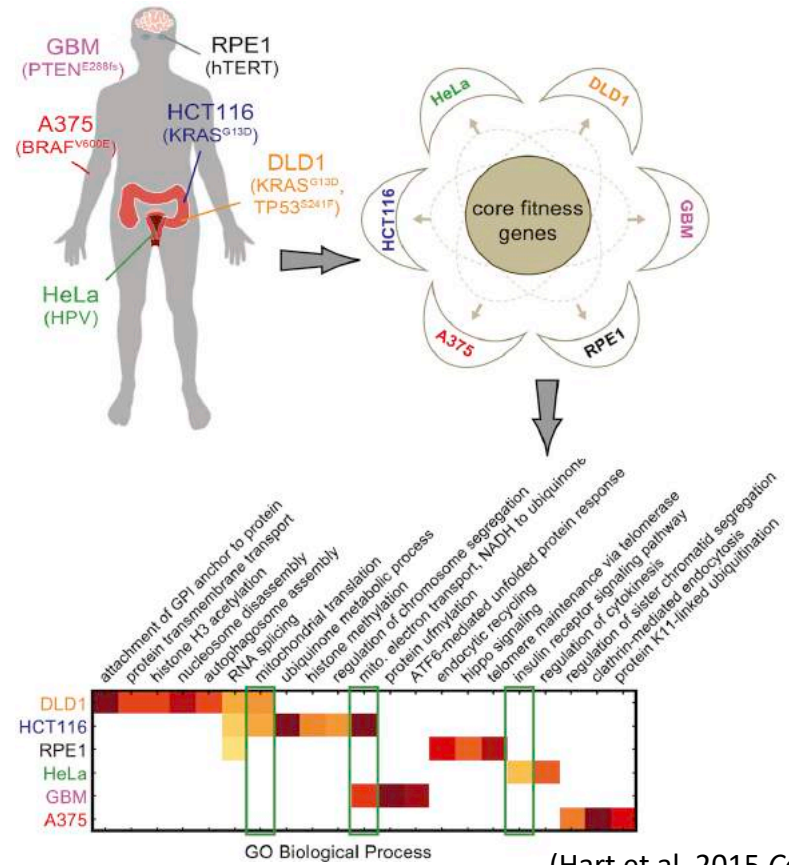


Esszenciális gének normál vs. rákos sejtekben



(Wang et al. 2015 *Science*)

- a fehérje kódoló gének kb. 10%-a esszenciális (~2000 gén) az emberi sejtvonalak in vitro fennmaradásához



(Hart et al. 2015 *Cell*)

- különböző rákos sejtvonalak vizsgálata megmutathatja, hol van az egyes betegségek Achilles-ina, hol lehet olyan célzott terápiát alkalmazni, ami specifikusan a daganatos sejteket támadja