

GOVERNING BENEFITS AND RISKS OF EMERGING BIOTECHNOLOGIES

RETROSPECTIVE AND CURRENT CASES

10:05-10:25 AM 17 FEBRUARY 2022

Professor Kenneth A. Oye

MIT Political Science and Data Systems and Society

OUTLINE

Risk Governance Under Uncertainty: Permissive, Precautionary, Adaptive
Retrospective and Current Biotechnology Cases

- Agriculture: Genetically Engineered Plants and Animals
- Industry: Synthesis of Biofuels, Fragrances, and Drugs
- Medicine: SCGT, Regenerative Medicine, HGGT
- Environment: Gene Drives

A Closing Conundrum: Governing Risks with Information Hazards

The research, conferences and workshops on which this presentation is based have been supported by the Netherlands Rijksinstituut voor Volksgezondheid en Milieu; the US Defense Advanced Research Projects Agency, US National Science Foundation Molecular and Cellular Biology, the MIT Center for Biomedical Innovation, the International Risk Governance Council and Richard Johnson.

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APPROACHES TO RISK GOVERNANCE UNDER UNCERTAINTY

Permissive

Allow innovation unless environment, health, security are clearly compromised

After-the-fact reaction if crisis materializes; backlash may limit innovation

Examples: DDT, post-Fukushima nuclear shutdown, US stasis on gene therapy



Precautionary

Limit innovation unless environment, health and security are clearly protected

Diversion of innovation to less regulated areas may heighten risks

Examples: EU on GMOs, US on stem cell research, Y2K



11:59:59

31 DECEMBER 1999

12:00:00

01 JANUARY 2000

Planned Adaptive

- * Prepare: Fund research to inform priors on benefits and risks
- * Discriminate: Foster initial applications with most favorable priors
- * Observe: Harvest and process information from initial experience
- * Adapt: Learn from experience and update/correct practices

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WHY BIOENGINEERING IS CHANGING SO FAST

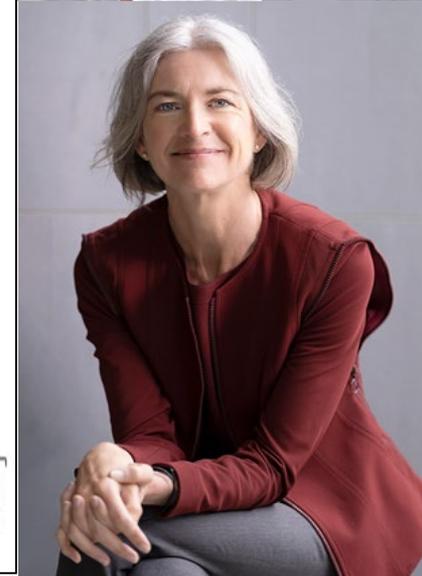
ENABLING TECHNOLOGIES DEVELOPING AT EXTRAORDINARY RATES

Making DNA: Cost and speed of DNA synthesis

Reading DNA: Cost and speed of DNA sequencing

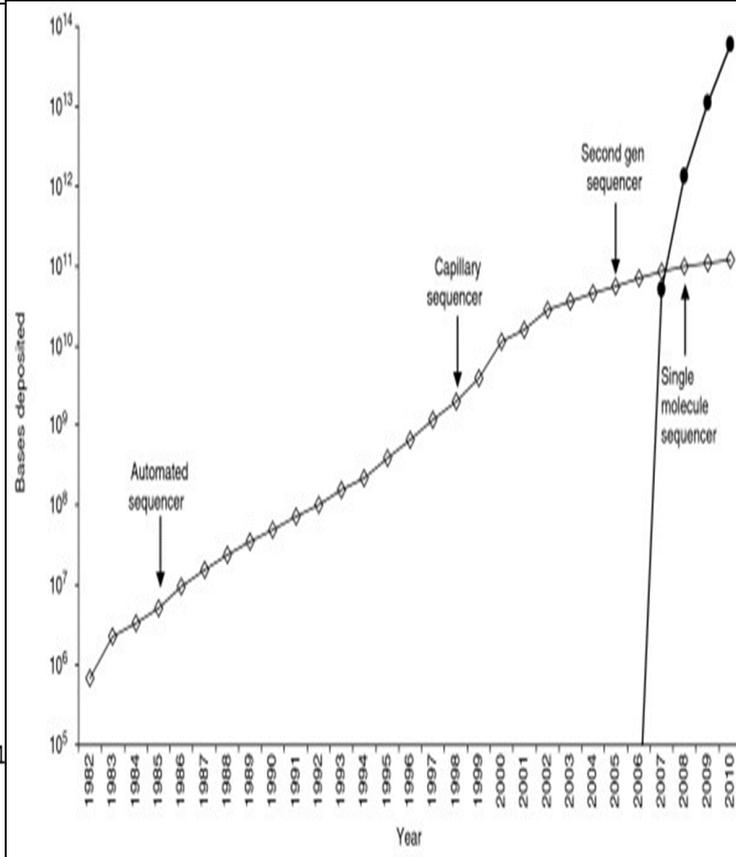
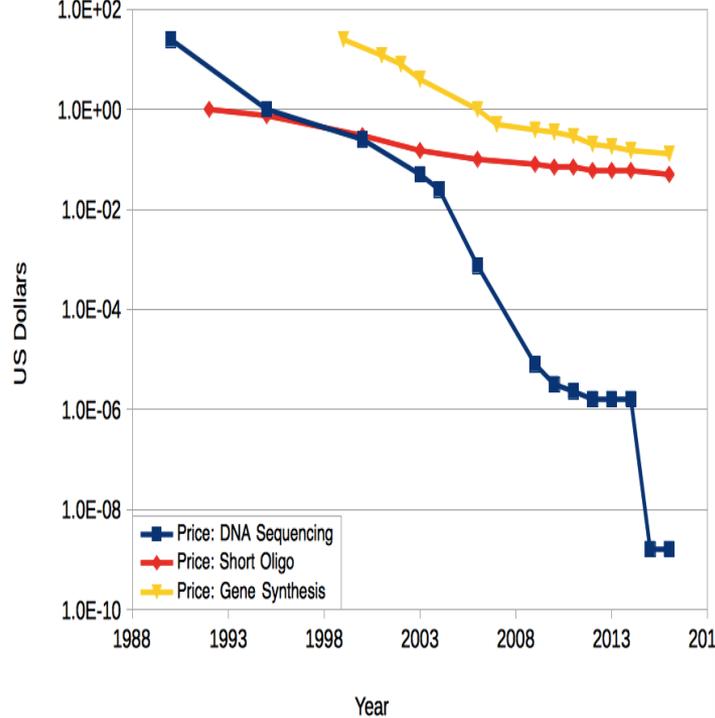
Characterizing and Designing DNA: AI and data storage

Editing DNA: From zinc fingers and talens to CRISPR and base editing



Price Per Base of DNA Sequencing and Synthesis

Rob Carlson, March 2016, www.synthesis.cc



Cost of DNA Synthesis & Sequencing

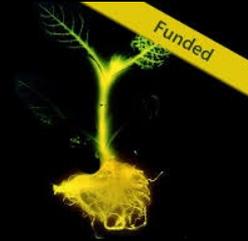
Info Deposited in Data Bases

Design/Editing Tools

EMERGING APPLICATIONS

AGRICULTURE

GM Crops and Livestock N Fixation, Glowing Plants, Aquabounty



INDUSTRY

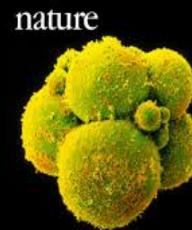
Synthesis of Organic Materials

Fuel, Flavors, Drugs



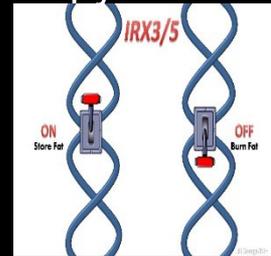
MEDICINE

Regenerative Medicine, Somatic and Germline Cell Therapy



Chinese scientists have reported genetically modifying human embryos

bit.ly/editedembryo



ENVIRONMENT

Remediation; Control Vector Borne Disease and Invasive Species





AQABOUNTY ATLANTIC SALMON WITH POUT AND CHINOOK GENES

FOOD SAFETY: FDA compared non-GE farmed and GE salmon, with focus on estradiol, testosterone, 11-ketotestosterone, T3, T4 and insulin-like growth factor 1 (IGF1)) + complete compositional analysis.

ENVIRONMENT: FDA issued “Finding of No Significant Impact” (FONSI)

Physical containment: All fish in land based tanks with safeguards

Geophysical containment: High water temp or bad water quality g

Biological containment – organism design

- Adults sterile triploid (diploid broodstock)
- Adults all female
- Fitness disadvantage relative to wild type due to limited food

EMERGING APPLICATIONS

AGRICULTURE

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Synthesis of Organic Materials

Fuel, Flavors, Drugs



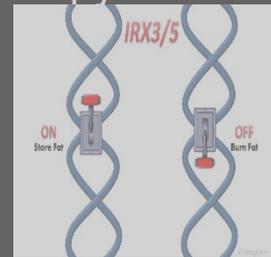
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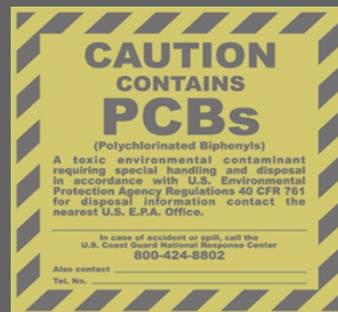
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ENVIRONMENT

Remediation; Control Vector Borne Disease and Invasive Species



INDUSTRIAL APPLICATIONS: MAKING LOW VALUE MATERIALS

SGI, Sapphire, Algenol

synthesis of biofuels

Universal Biomining

extraction and effluent treatment

Economics > low cost containment > environment release probable

Environmental effects

- Fitness / reproduction / propagation
- Horizontal gene flow
- Mutation > effects on fitness, gene flow?

Raceways and Sluices



Plastic Bags



Surface Ponds



ALGAE BIOFUELS CONTAINMENT WORKSHOP



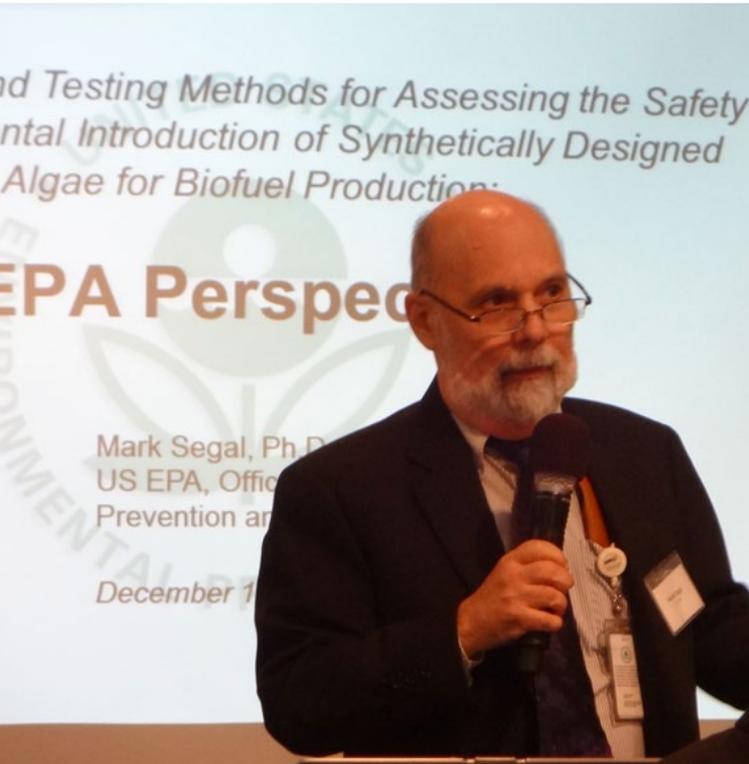
SYNTHETIC GENOMICS*

ALGENOL
B I O F U E L S

HARNESSING THE SUN TO FUEL THE WORLD™

Sapphire
Energy

Agilent Technologies



...nd Testing Methods for Assessing the Safety
...ntal Introduction of Synthetically Designed
...Algae for Biofuel Production:

EPA Perspectives

Mark Segal, Ph.D.
US EPA, Office of
Prevention and
Control

December 1



TECHNOLOGISTS DESIGN TO LIMIT RISKS OF RELEASE

Daisuke Kiga: Synthetic Amino Acids

Biocontainment Design: Church and Isaacs Labs

Unfilled gap - independent systematic demonstration and testing

ARTICLE

doi:10.1038/nature14121

LETTER

doi:10.1038

Biocontainment of genetically modified organisms by synthetic protein design

Daniel J. Mandell^{1*}, Marc J. Lajoie^{1,2*}, Michael T. Mee^{1,3}, Ryo Takeuchi⁴, Gleb Kuznetsov¹, Julie E. Norville¹, Christopher J. Gregg¹, Barry L. Stoddard⁴ & George M. Church^{1,5}

Recoded organisms engineered to depend on synthetic amino acids

Alexis J. Rovner^{1,2}, Adrian D. Haimovich^{1,2*}, Spencer R. Katz^{1,2*}, Zhe Li^{1,2}, Michael W. Grome^{1,2}, Brandon M. Gass^{1,2}, Miriam Amiram^{1,2}, Jaymin R. Patel^{1,2}, Ryan R. Gallagher^{1,2}, Jesse Rinehart^{2,3} & Farren J. Isaacs^{1,2}

US EPA FUNDS RESEARCH ON IMPROVED METHODS OF TESTING

First grants awarded July 2021



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Research Grants

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Assessment Tools for Biotechnology Products

INDUSTRIAL APPLICATIONS: MAKING HIGH VALUE MATERIALS

Keasling/Amyris/Sanofi

Prather

Evolva

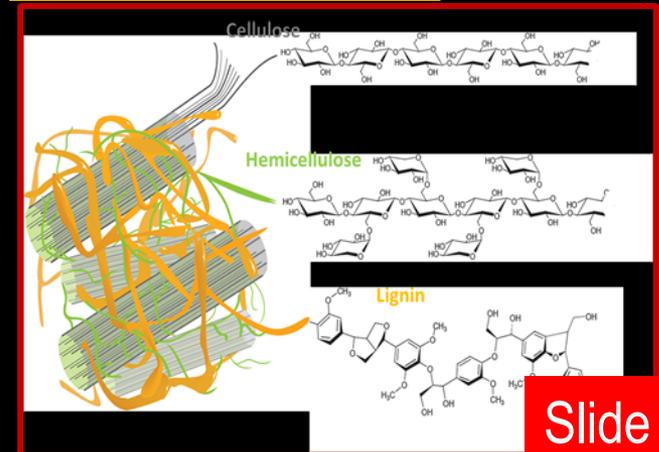
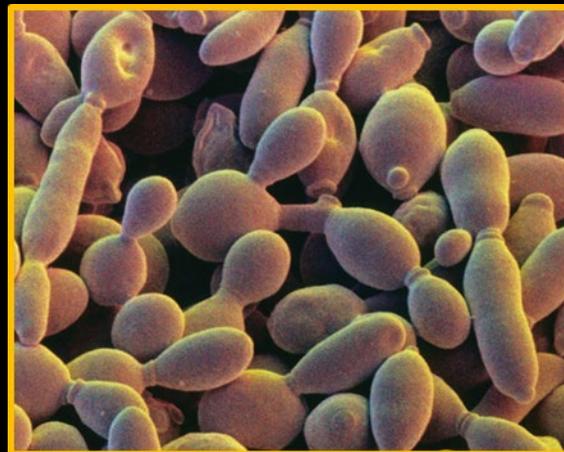
Smolke, Dueber, Martin, Facchini

amorphadien, artemisinin

glucaric acid

vanillin

reticuline, hydrocodone, morphine



OPIATE PRODUCTION IN YEAST DUEBER, MARTIN, SMOLKE
Benefits: Ability to vary scaffolds, control intermediates, create novel analgesics
Hazard: Production beyond control of cartels and law enforcement
Technical Measures: markers, finicky strains, unappealing final product
Policy Measures: lab security, licensing, synthesis screening

nature International weekly journal of science



Illegal use of opiates such as heroin and morphine affects more than 16 million people worldwide.

Regulate 'home-brew' opiates

The research community and the public require a fast, flexible response to the synthesis of morphine by engineered yeasts, urge **Kenneth Oye, Tania Bubela and J. Chappell H. Lawson.**

Every year, thousands of students from across the world compete to build biological systems from pre-existing parts in a competition organized by the International Genetically Engineered Machine (iGEM) Foundation. Last November, to spark discussion on security and health risks raised by synthetic biology,

FBI Special Agent Edward You presented an example: the production of opiates from sugar by yeast (*Saccharomyces cerevisiae*) that has been genetically modified.

Your hypothetical scenario is becoming a reality. One week after the iGEM competition, two developers of opiate-producing yeast strains approached us, specialists in

biotechnology policy. They had results in advance of publication, and requested advice on how they might maximize the benefits of their research while mitigating the risks. Now, published papers by these researchers — John Dueber at the University of California, Berkeley, and his colleagues¹, and Vincent Martin ▶

Science

SYNTHETIC BIOLOGY

Complete biosynthesis of opioids in yeast

Stephanie Galanie,¹ Kate Thodey,² Isis J. Trenchard,² Maria Filsinger Interrante,² Christina D. Smolke^{2*}

Opioids are the primary drugs used in Western medicine for pain management and palliative care. Farming of opium poppies remains the sole source of these essential medicines, despite diverse market demands and uncertainty in crop yields due to weather, climate change, and pests. We engineered yeast to produce the selected opioid compounds thebaine and hydrocodone starting from sugar. All work was conducted in a laboratory that is permitted and secured for work with controlled substances. We

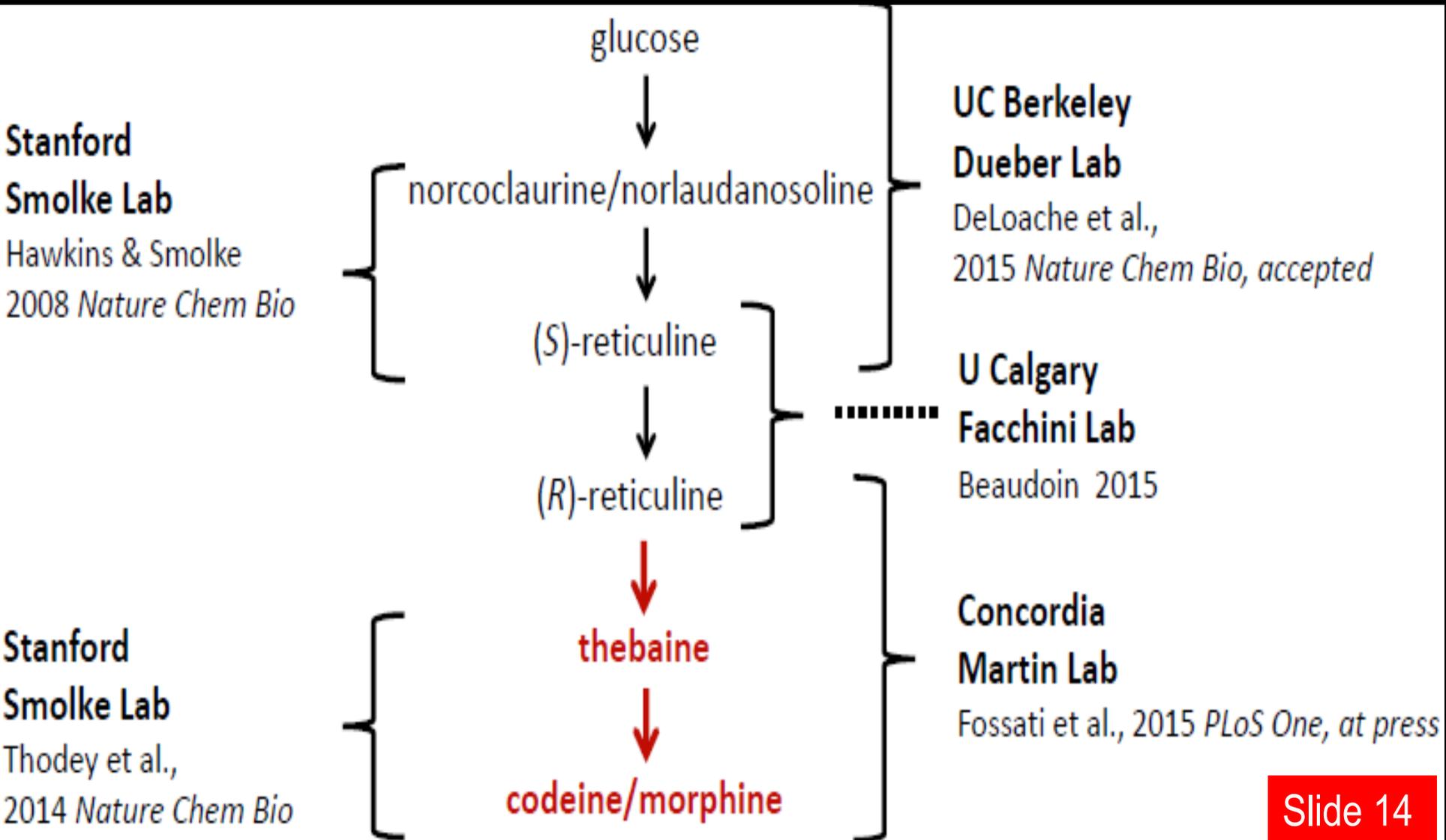


Slide 13

OPIATE PRODUCTION IN YEAST: TWO RESEARCH GROUPS

Smolke, Thodey, Hawkins (Stanford)

Dueber (UCB), Facchini (Calgary), Martin (Concordia)



(NOT) BREWING BAD

PERSONNEL SECURITY

Psychological disorders

- Psychopathy
- Borderline personality disorder
- Narcissistic personality disorder

Other Risk Factors

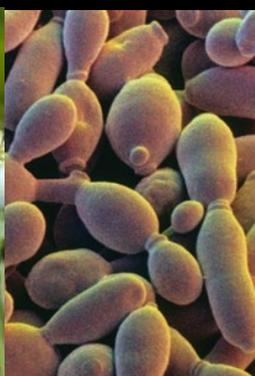
- Financial stress
- Status insecurity
- Sleep deprivation
- Perceived unfairness

LAB SECURITY

- Entry and exit control
- Materials access
- Inventory management
- Information controls

REDUCE APPEAL OF STRAINS

- Insert markers for traceability
- Make hard to cultivate
- Stop short of opiates
- Make distasteful to consume



EMERGING APPLICATIONS

AGRICULTURE

GM Crops and Livestock N Fixation, Glowing Plants, Aquabounty



INDUSTRY

Synthesis of Organic Materials

Fuel, Flavors, Drugs

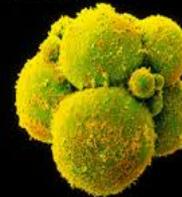


MEDICINE

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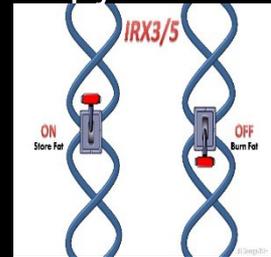


nature



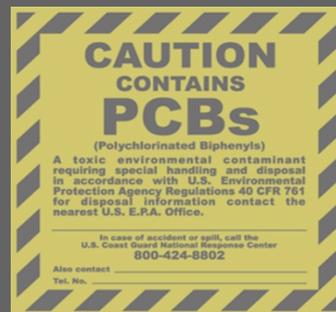
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ENVIRONMENT

Remediation; Control Vector Borne Disease and Invasive Species



VERSION 1.0 ADVERSE EFFECTS, OFF TARGET EFFECTS AND MUTATIONS

doi:10.1038/nature16526



High-fidelity CRISPR-Cas9 nucleases with no detectable genome-wide off-target effects

Benjamin P. Kleinstiver^{1,2*}, Vikram Pattanayak^{1,2*}, Michelle S. Prew¹, Shengdar Q. Tsai^{1,2}, Nhu T. Nguyen¹, Zongli Zheng³ & J. Keith Joung^{1,2}

CRISPR-Cas9 nucleases are widely used for genome editing but can induce unwanted off-target mutations. Existing strategies for reducing genome-wide off-target effects of the widely used *Streptococcus pyogenes* Cas9 (SpCas9) are imperfect, possessing only partial or unproven efficacies and other limitations that constrain their use. Here we describe SpCas9-HF1, a high-fidelity variant harbouring alterations designed to reduce non-specific DNA contacts. SpCas9-HF1 retains on-target activities comparable to wild-type SpCas9 with >85% of single-guide RNAs (sgRNAs) tested in human cells. Notably, with sgRNAs targeted to standard non-repetitive sequences, SpCas9-HF1 rendered all or nearly all off-target events undetectable by genome-wide break capture and targeted sequencing methods. Even for atypical, repetitive target sites, the vast majority of off-target mutations induced by wild-type SpCas9 were not detected with SpCas9-HF1. With its exceptional precision, SpCas9-HF1 provides an alternative to wild-type SpCas9 for research and therapeutic applications. More broadly, our results suggest a general strategy for optimizing genome-wide specificities of other CRISPR-RNA-guided nucleases.

Unexpected mutations after CRISPR-Cas9 editing *in vivo*

To the Editor: CRISPR-Cas9 editing shows promise for correcting disease-causing mutations. For example, in a recent study we used CRISPR-Cas9 for sight restoration in blind *rd1* mice by correcting a mutation in the *Pde6b* gene¹. However, concerns persist regarding secondary mutations in regions not targeted by the single guide RNA (sgRNA)². Algorithms generate likely off-target sites for a given gRNA, but these algorithms may miss mutations. Whole-genome sequencing (WGS) has been used to assess the presence of small insertions and deletions (indels)³ but not to probe for single-nucleotide variants (SNVs) in a whole organism. We performed WGS on a CRISPR-Cas9-edited mouse to identify all off-target mutations and found an unexpected number of SNVs compared with the widely accepted assumption

Slide 17

Public Interest Group Calls for Public Disclosures in Gene Therapy Death



Contact: Osagie Obasogie
510-625-0819, ext 310

Troubling new revelations have emerged this week in the death of an Illinois woman in a gene therapy trial for arthritis, prompting the Center for Genetics and Society to call on the federal government to consider firmer regulatory action.

The death of 36-year-old Jolee Mohr in a gene transfer experiment sponsored by Targeted Genetics Corporation of Seattle

Jolee Mohr, with husband Robb and daughter Torie in 2006, died after two injections of a gene therapy. The cause is not yet known. Photo courtesy of Mohr Family

raises important concerns about other participants in the 20 study locations across the country, and about participants in clinical trials of gene therapy and biomedical research more generally.

VERSION 1.5

CURRENT SOMATIC CELL GENE THERAPY (SCGT)

Single gene alterations to cure thalassemia, cystic fibrosis, hemophilia.

800+ SCGT now under development

Bluebird LentiGlobin BB305 for β -thalassaemia approved by EMA FDA



bluebirdbio™

VERSION 2.0 REGENERATIVE MEDICINE

REPLACE

Engineer differentiated tissue/organ
Insert/transplant in subject

- Tracheal implants - Macchiarrini
- Retinal Tissue Implant – Kurimoto

REGENERATE

Trigger internal healing in subject
Insert extracellular matrix, modified stem cells

- Own cord blood stem cells
- Donor stem cells, marrow
- Procymal for graft-versus-host disease

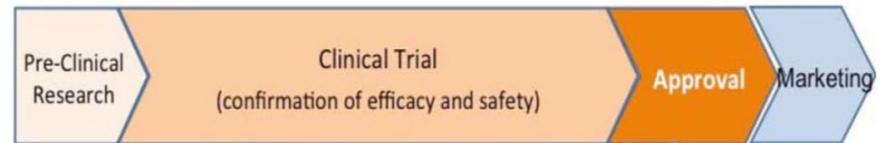
PMDA “Conditional Time Limited Authorization”

Pharmaceuticals Licensing and Reimbursement in the European Union, United States, and Japan

KA Oye¹, HG Eichler², A Hoos³, Y Mori⁴, TM Mullin⁵ and M Pearson⁶



【Traditional Approval Process】



【New Scheme for Regenerative Medicine Products】



* Probable benefit: Confirmation of efficacy with small patient population.

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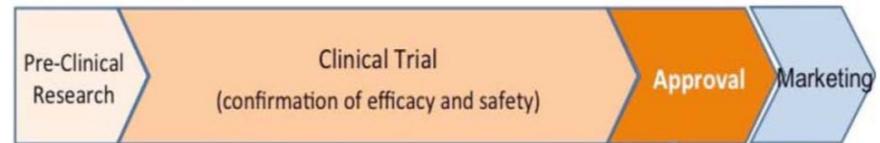


Pharmaceuticals Licensing and Reimbursement in the European Union, United States, and Japan

KA Oye¹, HG Eichler², A Hoos³, Y Mori⁴, TM Mullin⁵ and M Pearson⁶



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VERSION 3.0 GERMLINE GENE THERAPY (GGT)

SCGT works in individual, GGT changes in germline will be heritable

- Huang@Sun Yat-sen edited β -thalassaemia gene 28 embryos. Experiment failed with many off-target effects (4/2015 Protein&Cell)
- Zhang@Broad Improved Cas9 Specificity (12/2015 Science)
- Joung@MGH Hi-fi CRISPR no off-target effects (1/2016 Nature)
- Mitalipov@Oregon GGT pathogen in human embryo (8/2017 Nature)
- HGP Write – ultra-safe cells with codon knockout to limit gene flow

Protein Cell 2015, 6(5):363–372
DOI 10.1007/s13238-015-0153-5



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[✉], Junju 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
✉ Correspondence: hjunju@mail.sysu.edu.cn (J. Huang), zhoucanquan@hotmail.com (C. Zhou)

Received March 30, 2015 Accepted April 1, 2015

GENOME EDITING

Rationally engineered Cas9 nucleases with improved specificity

Ian M. Slaymaker,^{1,2,3,4*} Linyi Gao,^{1,4*} Bernd Zetsche,^{1,2,3,4} David A. Scott,^{1,2,3,4} Winston X. Yan,^{1,5,6} Feng Zhang^{1,2,3,4,†}

The RNA-guided endonuclease Cas9 is a versatile genome-editing tool with a broad range of applications from therapeutics to functional annotation of genes. Cas9 creates double-strand breaks (DSBs) at targeted genomic loci complementary to a short RNA guide. However, Cas9 can cleave off-target sites that are not fully complementary to the guide, which poses a major challenge for genome editing. Here, we use structure-guided protein engineering to

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NATURE | ARTICLE

日本語要約

Correction of a pathogenic gene mutation in human embryos

Hong Ma Nuria Marti-Gutierrez Sang-Wook Park Jun Wu Yeonmi Lee Keiichiro Suzuki Amy Koski Dongmei Ji Tomonari Hayama Riffat Ahmed Hayley Darby Crystal Van Dyken Ying Li Eunju Kang A.-Reum Park Daesik Kim Sang-Tae Kim Jianhui Gong Ying Gu Xun Xu David Battaglia Sacha A. Krieg David M. Lee Diana H. Wu Don P. Wolf Stephen B. Heitner Juan Carlos Izpisua Belmonte Paula Amato Jin-Soo Kim Sanjiv Kaul Shoukhrat Mitalipov

Affiliations | Contributions | Corresponding authors

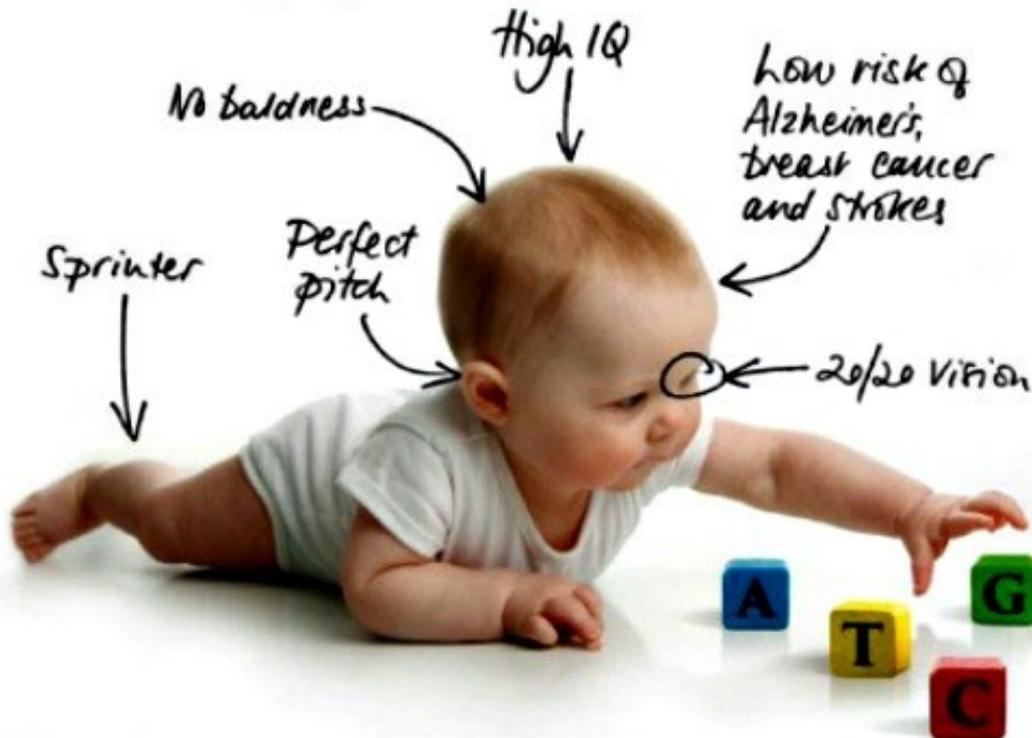
Nature 548, 413–419 (24 August 2017) | doi:10.1038/nature23305

Received 28 March 2017 | Accepted 27 June 2017 | Published online

Slide 21

Editing humanity

The prospect of genetic enhancement



BEYOND THERAPEUTICS

OLYMPIC ENHANCEMENTS

- SIZE
- SPEED
- AGILITY
- PAIN THRESHOLD
- JOINT STRENGTH
- EYE-HAND COORDINATION
- DEPTH PERCEPTION

MILSPEC ENHANCEMENT

- PAIN THRESHOLD
- TOLERANCE OF SLEEP LOSS
- VISUAL ACUITY
- PARALLEL PROCESSING
- COURAGE
- POSITIVE OWN GROUP AFFECT
- NEGATIVE OUT-GROUP AFFECT

WHICH ENHANCEMENTS ARE IN FACT FEASIBLE? ETHICAL?

EMERGING APPLICATIONS

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INDUSTRY

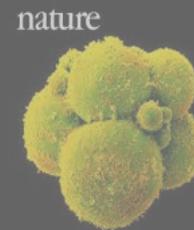
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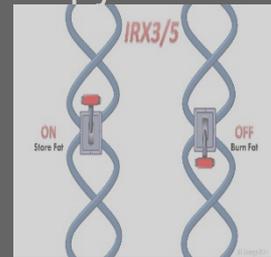
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ENVIRONMENT

Remediation; Control Vector Borne Disease and Invasive Species



Regulating gene drives

Kenneth A. Oye,^{1,2*} † Kevin Esvelt,^{3*} Evan Appleton,⁴ Flaminia Catteruccia,^{5,6} George Church,³ Todd Kuiken,⁷ Shlomiya Bar-Yam Lightfoot,² Julie McNamara,² Andrea Smidler,^{5,8} James P. Collins⁹

¹Political Science Department, Massachusetts Institute of Technology. ²Engineering Systems Division, Massachusetts Institute of Technology. ³Wyss Institute, Harvard University. ⁴Bioinformatics, Boston University. ⁵Harvard School of Public Health. ⁶University of Perugia, Italy. ⁷Woodrow Wilson International Center for Scholars. ⁸Harvard Medical School. ⁹School of Life Sciences, Arizona State University.

*Principal contributors to this piece.

†Corresponding author. oye@mit.edu

Regulatory gaps must be filled before gene drives could be used in the wild

Genes in sexually reproducing organisms normally have, on average, a 50% chance of being inherited, but some genes have a higher chance of being inherited. These genes can increase in relative frequency in a pop-

ulation. However, ecological effects would not necessarily be re-

verse. Genome engineering that uses the CRISPR-Cas9 nuclease Cas9 to cut sequences specified by guide RNA molecules (5, 6). This technique is in widespread use and has already engineered the genomes of more than a dozen species. Cas9 may enable "RNA-guided gene drives" to edit nearly any gene in sexually reproducing populations (1).

To reduce potential negative effects in advance of construction and testing, Esvelt et al. have proposed several novel types of drives (1). Precision drives could exclusively affect particular species or subpopulations by targeting sequences unique to those groups. Immunizing drives could block the spread of unwanted gene drives by preemptively altering target sequences. Reversal drives could overwrite unwanted changes introduced by an initial drive or by conventional genome engineering, even restoring the original sequence. However, ecological effects would not necessarily be re-



WHAT IS A GENE DRIVE ?

Mendelian: 50% odds genetic alteration will pass to next generation
IFF fitness or reproductive edge, THEN propagate

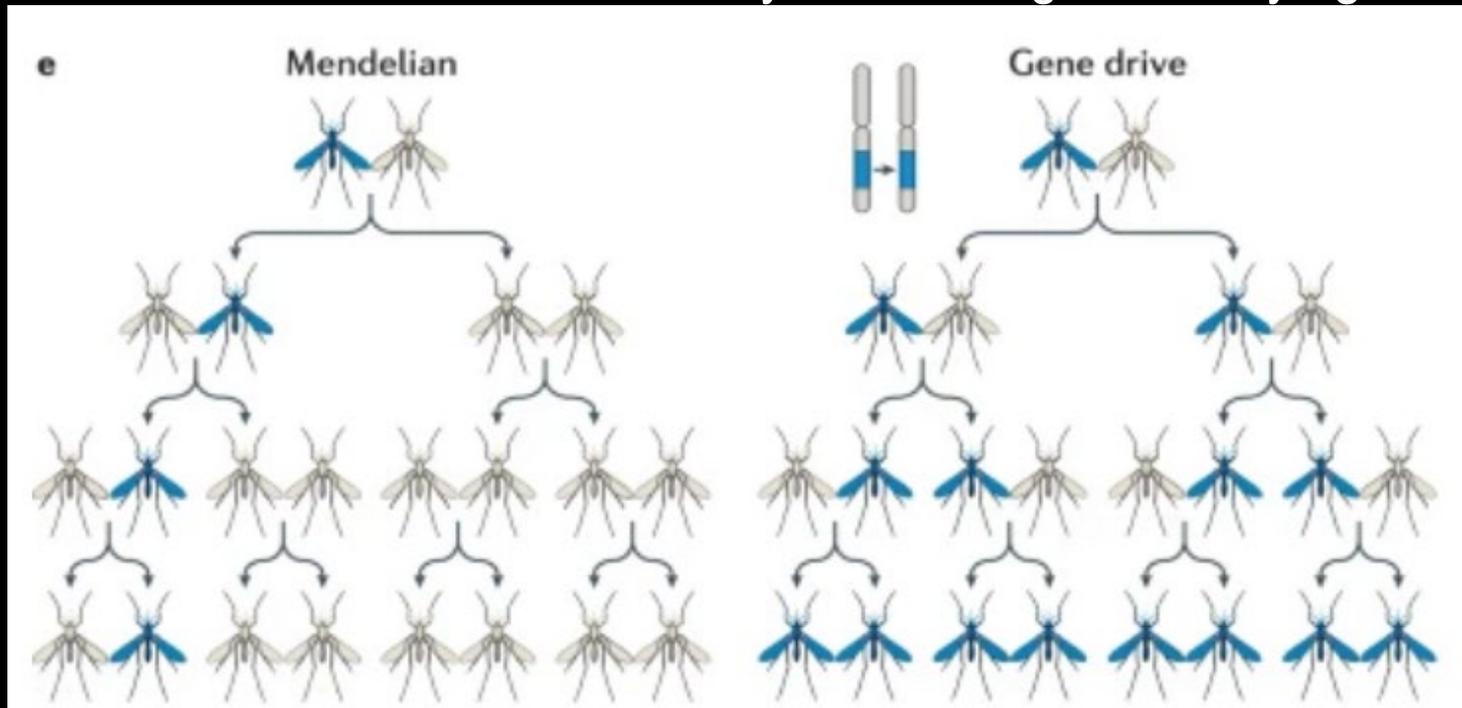
Gene drive: 99.5% odds alteration will pass to next generation (caveat)
Edit whole population without fitness or reproductive edge

FOR WHAT APPLICATIONS?

Environment: Control invasive species

Agriculture: Cut herbicide/pesticide resistance, eradicate pests

Health: Control vector borne diseases by eradicating or modifying vector



ENVIRONMENTAL ISSUES

Mutation of gene drives inevitable, will alter effects

Lateral gene transfer may reduce discrimination

Immunization and reversal may not be effective

Diseases borne by vectors will evolve

Environmental effects will vary by species and alteration

SECURITY ISSUES

Gain-of-function enabling ability to host diseases

Suppression of crops and livestock in traditional agriculture

Suppression of pollinators and other keystone species

Immunization drives may protect self and allies from effects

Reversal drives may be withheld for economic or political gain

Security implications uncertain - note ingenuity and creativity

(MOST) WIZARDS ARE ACTING RESPONSIBLY

Code of Conduct, Use IP to Enforce, Open Access to Data



RESEARCHERS CODE OF CONDUCT SCIENCE AUGUST 2015

BIOSAFETY

Safeguarding gene drive experiments in the laboratory

Multiple stringent confinement strategies should be used whenever possible

By Omar S. Akbari^{1,2}, Hugo J. Bellen^{3,4}, Ethan Bier^{5,*}, Simon L. Bullock⁶, Austin Burt⁷, George M. Church^{8,9}, Kevin R. Cook¹⁰, Peter Duchek¹¹, Owain R. Edwards¹², Kevin M. Esvelt^{8,*}, Valentino M. Gantz⁵, Kent G. Golic¹³, Scott J. Gratz¹⁴, Melissa M. Harrison¹⁵, Keith R. Hayes¹⁶, Anthony A. James¹⁷, Thomas C. Kaufman¹⁰, Juergen Knoblich¹¹, Harmit S. Malik^{18,19}, Kathy A. Matthews¹⁰, Kate M. O'Connor-Giles^{14,20}, Annette L. Parks¹⁰, Norbert Perrimon^{9,21}, Phillip Port⁶, Steven Russell²², Ryu Ueda^{23,24}, Jill Wildonger²⁵

Gene drive systems promote the spread of genetic elements through popula-

fore used institutionally approved stringent barrier methods. Only one experimenter

research involving potential gene drive systems while formal national guidelines are developed. Although we cannot claim to represent all researchers, we share a commitment to the safe and responsible development of gene drive technology. Although we differ in our assessments of the types of precaution needed, we recognize that any single confinement strategy could fail. We therefore unanimously recommend that future studies use a combination of stringent confinement strategies (see the table) whenever possible and always use safeguards adequate for preventing the unintentional release of synthetic gene drive systems into natural populations.

RECOMMENDATIONS. RNA-guided gene drive systems are created by the germline a DNA cassette and a single synthetic guide RNA (sgRNA)

Potentially stringent confinement strategies for gene drive research

Multiple stringent confinement strategies should be used whenever possible.

TYPE	STRINGENT CONFINEMENT STRATEGY	EXAMPLES
Molecular	Separate components required for genetic drive Target synthetic sequences absent from wild organisms	sgRNA and Cas9 in separate loci (8) Drive targets a sequence unique to laboratory organisms (3,4,8)
Ecological	Perform experiments outside the habitable range of the organism Perform experiments in areas without potential wild mates	<i>Anopheles</i> mosquitoes in Boston <i>Anopheles</i> mosquitoes in Los Angeles
Reproductive	Use a laboratory strain that cannot reproduce with wild organisms	<i>Drosophila</i> with compound autosomes*
Barrier	Physical barriers between organisms and the environment •Remove barriers only when organisms are inactive •Impose environmental constraints •Take precautions to minimize breaches due to human error	Triply nested containers, >3 doors (6) Anesthetize before opening (6) Low-temperature room, air-blast fans Keep careful records of organisms, one investigator performs all experiments (6)

*An example of reproductive confinement would be *Drosophila* laboratory strains with a compound autosome, where both copies of a large autosome are conjoined at a single centromere. These strains are fertile when crossed inter se but are sterile when outcrossed to any normal or wild-type strain because all progeny are monosomic or trisomic and die early in development.

ACADEMICS, GOVERNMENTS AND NGOS DELIBERATE

UN Biological Weapons Convention Meeting of Experts Geneva August 2014

International Experts Group on Biosecurity Regulation, Berlin 2015

National Academy of Sciences, Consensus Committee, Washington DC 2016

European Union Synergene Amsterdam, June 2016

OECD Gene Editing Workshop, Ottawa October 2016

IUCN World Conservation Congress Hawaii September 2016

UN Convention on Biodiversity COP13 Cancun December 2016

UN Convention on Biodiversity SBSTTA-22 and SBI-2 Montreal July 2018

NIH Novel and Exceptional Technologies Advisory Committee Gene Drive Group 2021

Outreach Network for Gene Drive Research August 2021

Gene Drives on the Horizon

Advancing Science, Navigating Uncertainty,
and Aligning Research with Public Values



The National Academies of
SCIENCES • ENGINEERING • MEDICINE



IUCN
World
Conservatio
Congress
Hawai'i 2016

COP13-COPMOP8-COPMOP2
CANCUN, MEXICO 2016



MAINSTREAMING BIODIVERSITY FOR WELL-BEING
CONVENTION ON BIOLOGICAL DIVERSITY



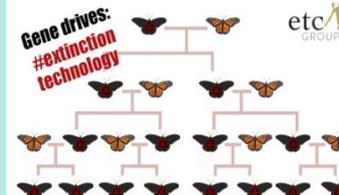
Convention on
Biological Diversity



PRIORITY PASS

MEETING	VENUE	DATES
SBSTTA-22 Twenty-second meeting of the Subsidiary Body on Scientific, Technical and Technological Advice	MONTREAL, CANADA	7/2/2018 7/7/2018
SBI-02 Second meeting of the Subsidiary Body on Implementation	MONTREAL, CANADA	7/9/2018 7/13/2018

BIOLOGICAL WEAPONS CONVENTION



GENE DRIVES FOR VECTOR CONTROL: CURRENT GOVERNANCE AND POLICY DEBATES AND THE IMPLICATIONS FOR RESEARCH

An overview of ongoing policy processes and debates from an African perspective, and of upcoming opportunities for experts from the continent to engage with the

5TH AUGUST 2021 | 15:00 - 17:00 EAT / 12:00 - 14:00 GMT

Slide 29

GOVERNING BENEFITS AND RISKS OF EMERGING BIOTECHNOLOGIES

RETROSPECTIVE AND CURRENT CASES

10:05-10:25 AM 17 FEBRUARY 2022

Professor Kenneth A. Oye

MIT Political Science and Data Systems and Society

OUTLINE

Risk Governance Under Uncertainty: Permissive, Precautionary, Adaptive
Retrospective and Current Biotechnology Cases

- Agriculture: Genetically Engineered Plants and Animals
- Industry: Synthesis of Biofuels, Fragrances, and Drugs
- Medicine: SCGT, Regenerative Medicine, HGGT
- Environment: Gene Drives

A Closing Conundrum: Governing Risks with Information Hazards

The research, conferences and workshops on which this presentation is based have been supported by the Netherlands Rijksinstituut voor Volksgezondheid en Milieu; the US Defense Advanced Research Projects Agency, US National Science Foundation Molecular and Cellular Biology, the MIT Center for Biomedical Innovation, the International Risk Governance Council and Richard Johnson.

INFORMATION HAZARDS: A CONUNDRUM

Information should be shared among biotechnology geeks to validate work and to provide building blocks for scientific advance.

Information should be shared with biosecurity policy wonks and biosafety officers to enable effective risk management.

Information should not be shared with malevolent or negligent actors to reduce the risk of harmful applications.

Compartmentalization of information is difficult / impossible.

Risk Analysis, Vol. 39, No. 5, 2019

DOI: 10.1111/risa.13235

Perspective

Information Hazards in Biotechnology

**Gregory Lewis,^{1,*} Piers Millett,¹ Anders Sandberg,¹ Andrew Snyder-Beattie,¹
and Gigi Gronvall²**

With the advance of biotechnology, biological information, rather than biological materials, is increasingly the object of principal security concern. We argue that both in theory and in practice, existing security approaches in biology are poorly suited to manage hazardous biological information, and use the cases of Mousepox, H5N1 gain of function, and Botulinum toxin H to highlight these ongoing challenges. We suggest that mitigation of these hazards can be improved if one can: (1) anticipate hazard potential before scientific work is performed; (2) consider how much the new information would likely help both good and bad actors; and (3) aim to disclose information in the manner that maximally disadvantages bad actors versus

PAST EXAMPLES?

How to make mammal-to-mammal transmissible strains of avian flu (2012)

Airborne Transmission of Influenza A/H5N1 Virus Between Ferrets

SANDER HERFST, EEFJE J. A. SCHRAUWEN, MARTIN LINSTER, SALIN CHUTINIMITKUL, EMMIE DE WIT, VINCENT J. MUNSTER, ERIN M. SORRELL, THEO M. BESTEBROER,

DAVID F. BURKE, [...] RON A. M. FOUCHIER +4 authors [Authors Info & Affiliations](#)

SCIENCE • 22 Jun 2012 • Vol 336, Issue 6088 • pp. 1534-1541 • DOI: 10.1126/science.1213362

Experimental adaptation of an influenza H5 HA confers respiratory droplet transmission to a reassortant H5 HA/H1N1 virus in ferrets

[Masaki Imai](#), [Tokiko Watanabe](#), [Masato Hatta](#), [Subash C. Das](#), [Makoto Ozawa](#), [Kyoko Shinya](#), [Gongxun Zhong](#), [Anthony Hanson](#), [Hiroaki Katsura](#), [Shinji Watanabe](#), [Chengjun Li](#), [Eiryu Kawakami](#), [Shinya Yamada](#), [Maki Kiso](#), [Yasuo Suzuki](#), [Eileen A. Maher](#)

[Nature](#) **486**, 420–428 (2012) | [Cite this](#)



EDITORIAL
July/August 2020 Volume 11 Issue 4 e01868-20
<https://doi.org/10.1128/mBio.01868-20>

Rethinking Gain-of-Function Experiments in the Context of the COVID-19 Pandemic

Michael J. Imperiale ^a, Arturo Casadevall ^b

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^bDepartment of Molecular Microbiology and Immunology, Johns Hopkins Bloomberg School of Public Health, Baltimore, Maryland, USA

Slide 32

CURRENT EXAMPLES?

How to synthesize horsepox (2018), How to make COVID variants (2021)

PLOS BIOLOGY

PLOS Biology | <https://doi.org/10.1371/journal.pbio.3001091> February 25, 2021

METHODS AND RESOURCES

A plasmid DNA-launched SARS-CoV-2 reverse genetics system and coronavirus toolkit for COVID-19 research

 **PLOS** | ONE

Published: January 19, 2018 • <https://doi.org/10.1371/journal.pone.0188453>

RESEARCH ARTICLE

Construction of an infectious horsepox virus vaccine from chemically synthesized DNA fragments

Ryan S. Noyce¹, Seth Lederman², David H. Evans^{1*}

¹ Department of Medical Microbiology & Immunology and Li Ka Shing Institute of Virology, University of Alberta, Edmonton, Alberta, Canada, ² Tonix Pharmaceuticals, Inc., New York, New York, United States of America

Technologists have a duty to engage with risks of their own work*

*Note: with info hazards responsible conduct is even more critical

Duty: Those proposing/conducting/funding research have a duty to evaluate actions with reference to both legal standards and ethical norms

Concrete Obligations that Follow from Duty:

- 101 Align own work with law/norms based on existing knowledge
- 201 Encourage others to align their work with laws/norms
- 301 Identify important gaps in knowledge and fill gaps through research
- 401 Identify gaps in legal coverage and join in debate on gaps

Engage early in order to allow time to:

- 101 Modify own work to fit law/norms before tech designs lock in
- 201 Organize to improve conduct of others before bad conduct locks in
- 301 Initiate research on benefits/risks in time to inform choices
- 401 Evaluate and influence policy before policy designs lock in

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