

# CG020 Genomika

## Přednáška 12

### Praktické aplikace genomiky

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# Literatura

- Literární zdroje ke kapitole 12:

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**Koblan, L.W., Erdos, M.R., Wilson, C., Cabral, W.A., Levy, J.M., Xiong, Z.M., Tavarez, U.L., Davison, L.M., Gete, Y.G., Mao, X., Newby, G.A., Doherty, S.P., Narisu, N., Sheng, Q., Krilow, C., Lin, C.Y., Gordon, L.B., Cao, K., Collins, F.S., Brown, J.D., and Liu, D.R.** (2021). In vivo base editing rescues Hutchinson-Gilford progeria syndrome in mice. *Nature*.

**Li, X., Qian, X., Wang, B., Xia, Y., Zheng, Y., Du, L., Xu, D., Xing, D., DePinho, R.A., and Lu, Z.** (2020). Programmable base editing of mutated TERT promoter inhibits brain tumour growth. *Nat Cell Biol* **22**, 282-288.

# Osnova

- Lékařství
  - Molekulární diagnostika
  - Individualizovaná medicína
  - Genová terapie
- Biotechnologie
- Geneticky Modifikované Organismy
  - Transgenoze
  - Editování genomu
- Modelové organismy
- Principy PCR

# Osnova

- Lékařství
  - Molekulární diagnostika

# Molekulární Diagnostika

- Cca 10,000 onemocnění u člověka je podmíněno mutací v jediném genu
  - cystická fibróza
  - srpkovitá anémie
  - svalová dystrofie
  - $\beta$ -talasémie
  - ....
- Časná molekulární diagnostika
  - mutace nebo infekce
    - PCR
    - Hybridizace na DNA čipu
    - Cas-based

# Molekulární Diagnostika

- Mammoth Biosciences

- Spoluzakladatelka Jenifer Doudna

<https://youtu.be/Pe4ldgKGdQ>



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It has been found that binding of gRNA=targeted CRISPR to the target sequence activates ssDNA cleavage activity of Cas12. “By combining Cas12a ssDNase activation with isothermal amplification, we create a method termed DNA endonuclease-targeted CRISPR trans reporter (DETECTR), which achieves attomolar sensitivity for DNA detection. DETECTR enables rapid and specific detection of human papillomavirus in patient samples, thereby providing a simple platform for molecular diagnostics.” (Chen et al., 2018). The approach was used also for designing the CRISPR-based detection of SARS-CoV-2 (Broughton et al., 2020).

# Osnova

- Lékařství
  - Molekulární diagnostika
  - Individualizovaná medicína

# Individualizovaná Medicína

- Využívá znalost **genomu** pro:
  - Předpověď **zdravotních rizik**
  - **Diagnositku**
  - Výběr **nejvhodnějšího typu léčby**
  - **minimalizuje nežádoucí efekty léčby**
  - **prevence**



# Individualizovaná Medicína

## What is Personalized Medicine?

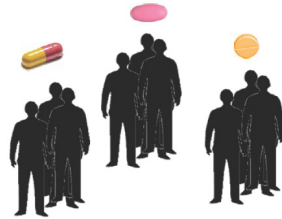
### Current Practice



One size fits all

Trial and error

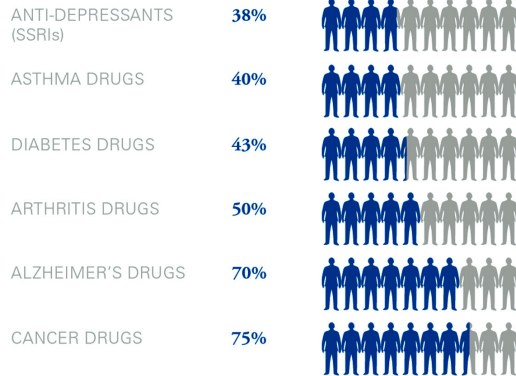
### Personalized Medicine



The **right treatment** for  
the **right person** at the  
**right time**

# Individualizovaná Medicína

## PERCENTAGE OF THE PATIENT POPULATION FOR WHICH A PARTICULAR DRUG IS INEFFECTIVE, ON AVERAGE



- Just in hospitals: about 6.7% of patients (2.2 million) experience serious adverse drug reactions



Serious adverse drug reactions in even smaller percentages of treated populations have led to the withdrawal of several drugs from the market

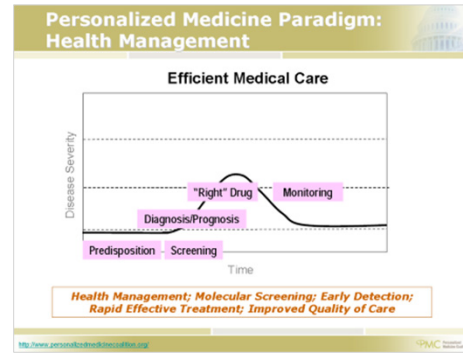
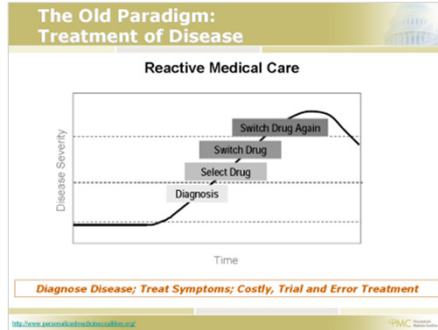
Zelnorm Vioxx Cylert

"Are good drugs going to the wrong people?"

Rezulin Baycol Lotronex\*

Source of data: Brian B. Spear, Margo Heath-Chiozzi, Jeffery Huff, "Clinical Trends in Molecular Medicine," Volume 7, Issue 5, 1 May 2001, Pages 201-204.

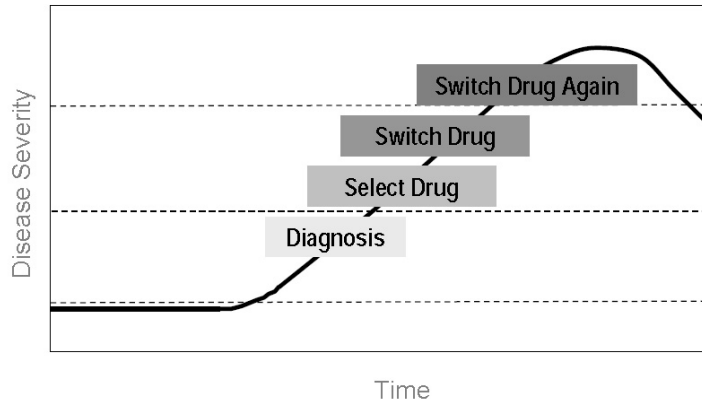
# Individualizovaná Medicína



# The Old Paradigm: Treatment of Disease



## Reactive Medical Care

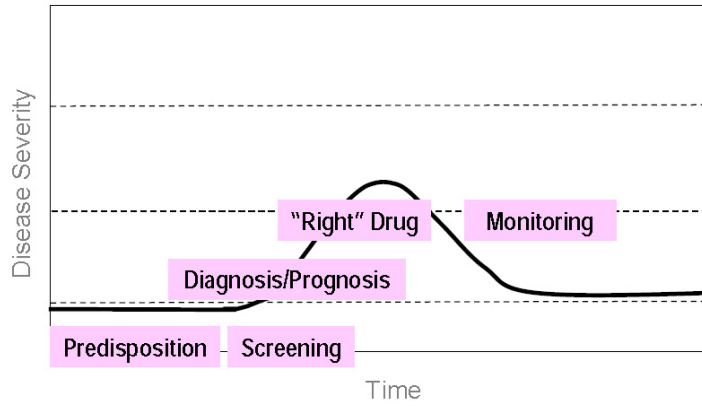


***Diagnose Disease; Treat Symptoms; Costly, Trial and Error Treatment***

# Personalized Medicine Paradigm: Health Management



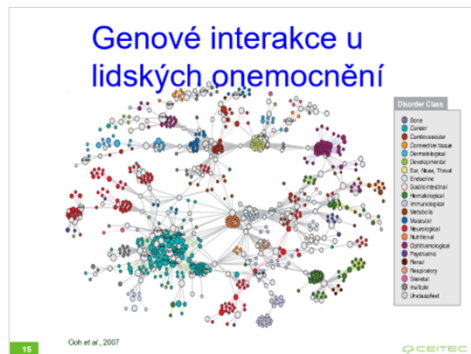
## Efficient Medical Care



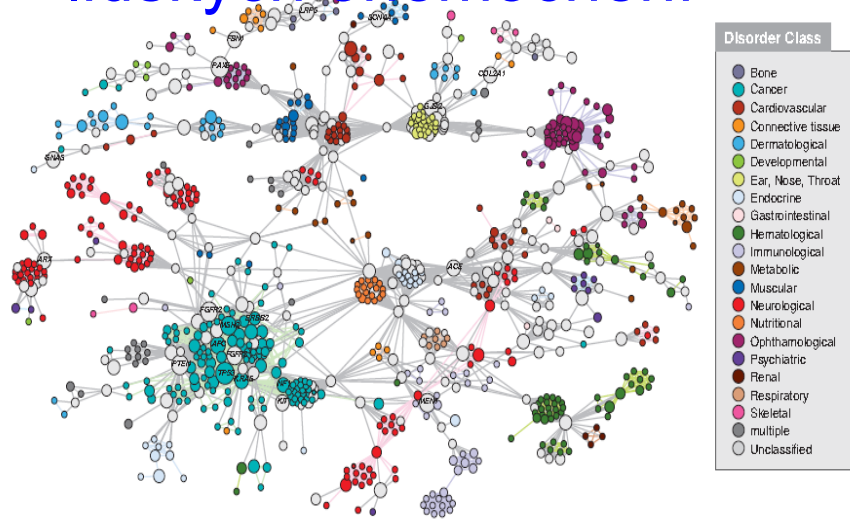
***Health Management; Molecular Screening; Early Detection;  
Rapid Effective Treatment; Improved Quality of Care***

# Individualizovaná Medicína

- Problém:
  - Mnohofaktoriální podmíněnost většiny lidských onemocnění



# Genové interakce u lidských onemocnění



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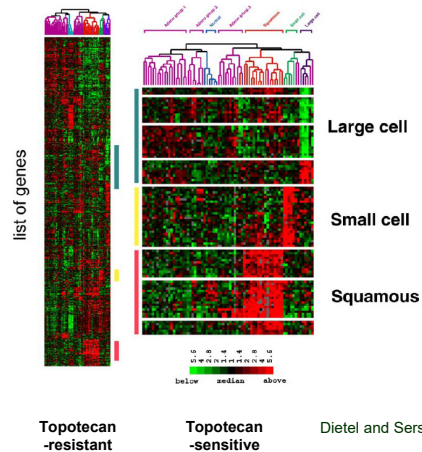
Goh et al., 2007

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In the disease gene network, each node is a gene, with two genes being connected if they are implicated in the same disorder. The size of each node is proportional to the number of disorders in which the gene is implicated (see key). Nodes are light gray if the corresponding genes are associated with more than one disorder class. Genes associated with more than five disorders, and those mentioned in the text, are indicated with the gene symbol. Only nodes with at least one link are shown.

# Individualizovaná Medicína

- Řešení problému:
  - **Systémová biologie** – využívá např. **genové klastrování** k identifikaci **genů asociovaných** s pozorovaným jevem (nemocí, poruchou, ..)



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Hierarchical clustering analysis exemplified for Topotecan-resistant (R) and Topotecan-sensitive (S) cell lines. All cell lines resistant to Topotecan (left panel) and all cell lines sensitive to Topotecan (right panel) express a unique set of genes. Each row in the cluster indicates the expression profile of a specific gene across all 19 cell lines. Each column indicates the individual cell line in which the gene is expressed.

Red, green, and black squares indicate that expression of the gene is greater than, less than, or equal to the median level of expression across all cell lines, respectively. The scale bar reflects the fold increase (red) or decrease (green) for any given gene relative to the median level of expression across all samples. Dietel and Sers, 2006.

**Topotecan**, sold under the brand name **Hycamtin** among others, is a [chemotherapeutic agent](#) medication that is a [topoisomerase inhibitor](#). It is a synthetic, water-soluble [analog](#) of the natural chemical compound [camptothecin](#). It is used in the form of its [hydrochloride salt](#) to treat [ovarian cancer](#), [lung cancer](#) and other cancer types (Wikipedia).



# Individualizovaná Medicína

- Řešení problému:
  - Systémová biologie** – využívá např. **genové klastrování** k identifikaci genů asociovaných s pozorovaným jevem (nemocí, poruchou, ..)
    - biomarkery
    - testy

**Table:** Selected Personalized Medicine Drugs, Treatments and Diagnostics as of September 2011\*

*Indications in green and relevant substructure, see cited from the therapeutic or diagnostic product label. Therapeutic product labels contain pharmacogenomic information as:*

- Information only
- Recommended
- Required
- Unhighlighted products have no pharmacogenomic information, recommendations or requirements in the label.

THERAPY	BIOMARKER/TEST	INDICATION
Metastatic (antitumor)	Chromosomal gene	<b>Avastin® adjuvant:</b> "Metastases in metastasized by breast (chemotherapy and should be used with great caution, if at all, in patients known to be or suspected of being heterozygous for the epistatic plasma-declustering gene."
Analgesic (analgesic)	CYP2C9	<b>Arbitelin:</b> "In some studies have demonstrated that cyclohexane P450 2C9 plays an important role in the metabolism of Paracetamol in some individuals."
Dyslipidemic (dyslipidemic)	US2 (ANG, CYP, A3), OTC, A3L, ABC	<b>Hydrochloride:</b> "Pharmacogenetic studies have shown that, in some individuals, the following mutation of the gene for the enzyme CYP2C9, which has been reported following initiation of adjuvant therapy in patients with early-stage breast cancer (ERCC), potentially modifies pharmacokinetic parameters."
Antitumor (antitumor) (antitumor) (antitumor)	Estrogen Receptor (ER)	<b>Breast cancer:</b> Estrogenic is indicated for adjuvant treatment of postmenopausal women with ER-positive early breast cancer. Tamoxifen is for treatment of breast cancer after surgery and for metastases in postmenopausal women. Tamoxifen is the standard therapy for estrogen receptor-positive early breast cancer in premenopausal women.
Chemotherapy	Mammaplasty®	<b>Breast cancer:</b> Diagnostic immunohistochemistry (IHC) test used for postoperative, node-negative, estrogen receptor-negative breast cancer patients who will receive hormonal therapy and are considering adjuvant chemotherapy.
Chemotherapy	Mammaplasty®	<b>Breast cancer:</b> Assess risk of breast metastasis in a 70-gene expression profile.
Chemotherapy	Oncocept DCC 16-gene signature	<b>Breast cancer:</b> A 16-gene signature (plus five reference genes) indicates whether a patient has a low, intermediate, or high risk of having a breast cancer within 10 years. Low-risk patients may be treated successfully with hormone therapy alone. High-risk patients may require more aggressive treatment with chemotherapy.
Chemotherapy	Compass™ 11-gene signature	<b>Breast cancer:</b> The test predicts "time to event" for metastasis of breast cancer, following surgery and biopsy.
Fibrotic (fibrotic)	Hormone Receptor (HR)	<b>Breast cancer:</b> Fulvestrant is indicated for the treatment of hormone receptor-positive metastatic breast cancer in postmenopausal women with disease progression following anti-estrogen therapy.
Therapeutic (therapeutic) (therapeutic) (therapeutic)	HER-2/neu receptor	<b>Breast cancer:</b> "For the treatment of patients with metastatic breast cancer whose tumors overexpress the HER-2 (Human Epidermal growth factor Receptor-2) protein and who have received one or more chemotherapy regimens for their metastatic disease." High levels of HER-2 expression have been associated with increased disease recurrence in breast cancer, but show a better response to trastuzumab.
Pharmaceutical and surgical prevention (pharmaceutical and surgical prevention)	BRCA 1/2	<b>Breast cancer:</b> Genetic surveillance and preventive treatment based on susceptibility risk for breast and ovarian cancer.
Antitumor (antitumor) (antitumor) (antitumor)	Breast Cancer Index™ (BCI) (BCI)	<b>Breast cancer:</b> Calculate a combined risk analysis for recurrence after tamoxifen treatment for ER-positive, node-negative breast cancer.

The Case for Personalized Medicine, 3rd edition, GENETEC

Aplikace přístupů systémové biologie (včetně genového klastrování na předchozím snímku) pak umožní identifikovat specifické markery a testy, které umožní testovat odpověď na různé typy terapií.

# Individualizovaná Medicína

- Další problémy
  - Etické otázky
    - Možnost zneužití znalosti genomu
    - riziko: nedostatečná ochrana dat
    - V některých zemích je uzákoněn omezený přístup pro určité typy zaměstnanců nebo pojišťovací společnosti
  - Vysoké náklady
    - Dělení medicíny na **first-class** and **low-class** služby
    - Zvětšování problému **globalizačního handicapu** – chudé země si nemohou takto pokročilý typ léčby dovolit
  - Soukromí
    - Zásadní a komplikovaná otázka
    - Jakou informaci lze považovat za soukromou?

# Osnova

- Lékařství
  - Molekulární diagnostika
  - Individualizovaná medicína
  - Genová terapie
- **Biotechnologie**

# Gene Therapy

Procedure in which the DNA sequence is inserted into the patient genome to replace or supplement the original gene

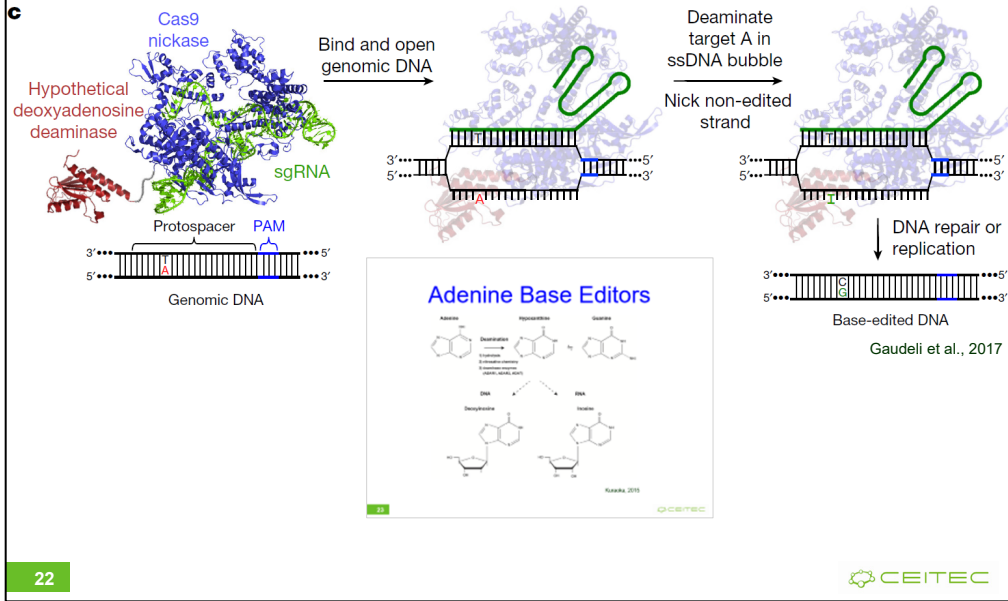
- Options:
  - replace the mutated gene
  - repair the mutation
  - deliver DNA encoding a therapeutic protein
  - antisense therapy
- In the future useful for treating e.g. hereditary diseases
- Types:
  - somatic gene therapy
  - gene therapy of germ cells

# Gene Therapy

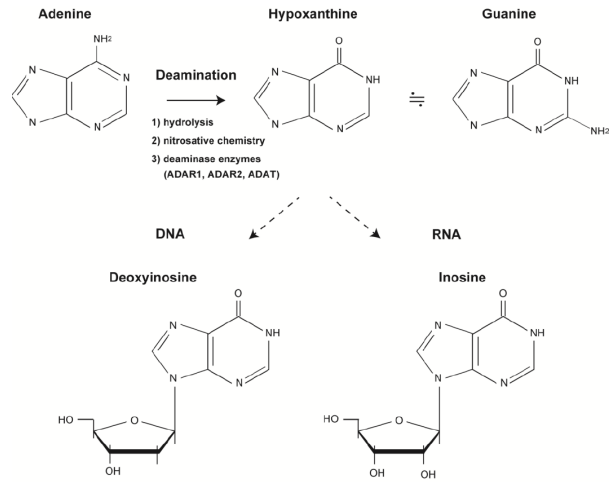


- Hutchinson–Gilford progeria syndrome
  - C•G-to-T•A mutace (c.1824 C>T; p.G608G) v genu pro laminin (*LMNA*)
  - Defekt v sestřihu RNA vede k tvorbě toxického proteoinu **progerinu**
  - Věk dožití cca 14 let
  - **In vivo oprava** pomocí ABEs potvrzena **u myší** a **lidských fibroblastů** (Koblan et al., 2021)

# Adenine Base Editors



# Adenine Base Editors



Kuraoka, 2015

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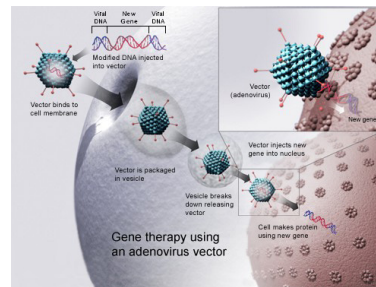
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Formation of deoxyinosine and inosine. Deamination of adenine to hypoxanthine results in the formation of deoxyinosine in DNA and inosine in RNA. Hypoxanthine, which is recognized as guanine, pairs with cytosine. Deoxyinosine and inosine are nucleosides that form when hypoxanthine is attached to a deoxyribose ring or ribose ring, respectively.

# Gene Therapy

- **Methods**

- **viral vectors**
  - retroviruses
  - adenoviruses
  - herpes simplex virus
- **non-viral methods**
  - injection of plasmid DNA into muscle
  - increased efficiency of DNA delivery
    - electroporation
    - sonoporation
    - „gene gun“ (biolistic)
    - magnetofection
- **genome editing**





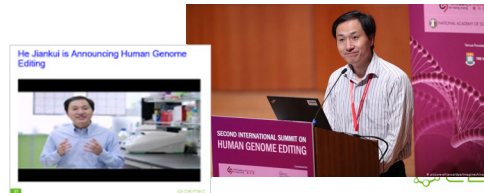
# Ethical Issues

- Regulace editace genomu v zemědělství a lidském zdraví
  - <https://crispr-gene-editing-regs-tracker.geneticliteracyproject.org/>
- International Commission on the Clinical Use of Human Germline Genome Editing
  - convened by the U.S. National Academy of Medicine (NAM), the U.S. National Academy of Sciences (NAS), and the Royal Society of the U.K. ...
  - ...to identify a number of scientific, medical, and ethical requirements that should be considered, and could inform the development of a potential pathway from research to clinical use — if society concludes that heritable human genome editing applications are acceptable
  - more details at <https://nationalacademies.org/gene-editing/international-commission/index.htm>

# Ethical Issues

- Alliance for Regenerative Medicine
  - international group representing the cell and gene therapy sector
  - put out a “statement of principles” on genome editing endorsed by 13 of the most active companies in this field
  - changing heritable DNA in sperm, eggs or a new embryo — came true in November 2018 when He Jiankui, a Chinese biophysicist, said that his lab had edited two baby girls to make them resistant to HIV infection. This mutation will be inherited by their descendants.
  - 31 clinical trials for gene edited therapies are in progress around the world, 20 of which are in oncology. None is yet close to commercialization. The US has the largest number of trials (19) followed by China (10) and the UK (6)

FT, Clive Cookson, Science Editor August 27 2019



## He Jiankui is Announcing Human Genome Editing

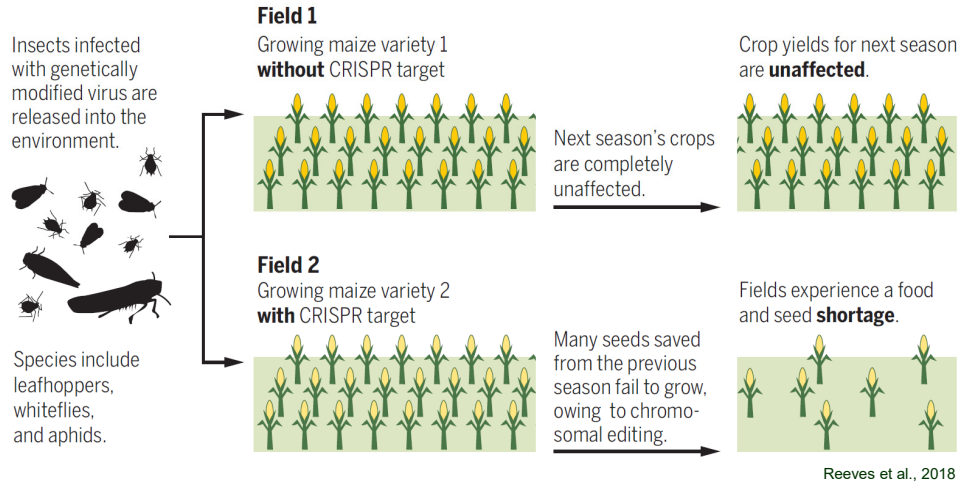


# Ethical Issues

- Genome editing as a **bioweapon**?
  - ongoing research program funded by the U.S. Defense Advanced Research Projects Agency (DARPA)
  - aims to disperse **infectious genetically modified viruses** that have been **engineered to edit crop chromosomes** directly in fields
  - the means of **delivery** of these **viral horizontal environmental genetic alteration agents (HEGAAs)** into the environment should be **insect-based dispersion**
  - Part of **scientific community** does not find the program useful for the U.S. agriculture, but points to its **possible misuse**



# Editing as a bioweapon?



# Outline

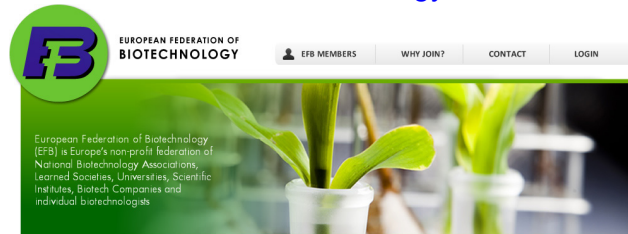
- **Medicine**
  - Molecular Diagnosis
  - Personalized Medicine
  - Gene Therapy
- **Biotechnology**

## BIOTECHNOLOGY

- It uses **living organisms, cells or parts of cells** (enzymes) for **research**, leading to **new products and applications** in **medicine, agriculture, food, environmental protection**
- Also used in developing **better/sustainable production methods** for the **chemical industry** and **other industrial processes**
- An **interdisciplinary approach** requiring knowledge of **chemistry, biology, physics, material sciences, engineering and informatics**
- The **origin** of biotechnology can be traced **4,000 years back**, when the **Sumerians** (although not knowingly) used microbes for the production of **alcoholic beverages**.

# BIOTECHNOLOGY

- **Examples**
  - effective utilization of **plant biomass** for **fuel production**
  - acquisition of starting material (**monomers**) for the **production of polymers** from living organisms instead of from fossil sources
  - **phytopharmaceuticals** – using plants in new vaccination methods such as expression of **antibodies** or **antigens** suitable for **immunization**
- **European Federation of Biotechnology**

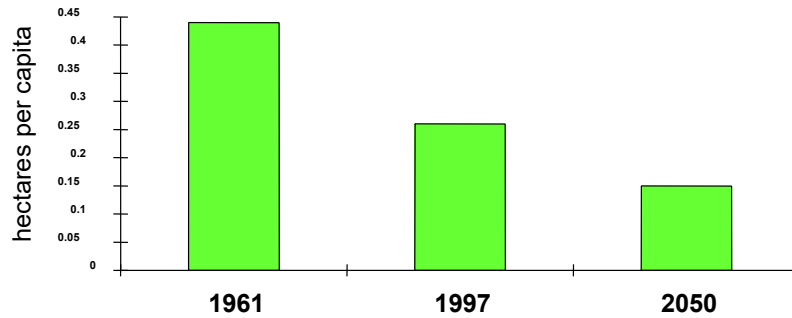




# Osnova

- Lékařství
  - Molekulární diagnostika
  - Individualizovaná medicína
  - Genová terapie
- Biotechnologie
- **Geneticky Modifikované Organismy**
  - Transgenoze

## Human Population vs Arable Land Availability



Source: UN Millennium Ecosystem Assessment

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Our civilization is built on farming, the surface area needed for feeding people has decreased by 90% over 10,000 years .

To prevent collapse, it is necessary to reduce this area from the current 0.45 ha/person to 0.2 ha/person by the year 2050. Return to original methods of agriculture would be a return to the original demands on area and therefore would be unsustainable Intensive farming = conversion of water and oil into food.

**goal of plant biotechnology** is to use all the available scientific knowledge to **breed varieties with higher yield** with lower inputs (of land, water, fertilizers, sprays ...)

# Nutrition Deficiency



<https://qz.com/africa/1064653/the-world-could-run-out-of-food-two-decades-earlier-than-thought/>

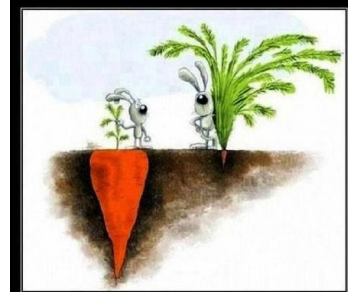
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Announced recently by Quartz server, the world could be facing a 214 trillion calorie deficit in the food production (announced **as soon as in 2027**).

# Breeding

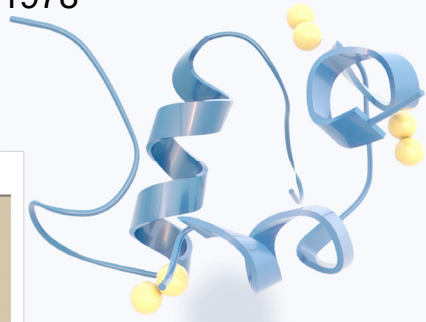
- organisms naturally vary due to **mutations**
- before the era of genetic engineering - **question of chance**
- breeding tools
  - **selection** and **crossing**
- **modern breeder** learned to **change hereditary information** – **increase the mutants allele frequency**
  - chemicals, radiation ...
- results are **incidental/non-targeted**



**Success**  
is **not always** visible at a glance

# Genetic Engineering

- Targeted modification ("targeted breeding")
  - ability to transfer genes = **transgenesis**
- the first practical application: production of **human insulin** in bacteria - 1978



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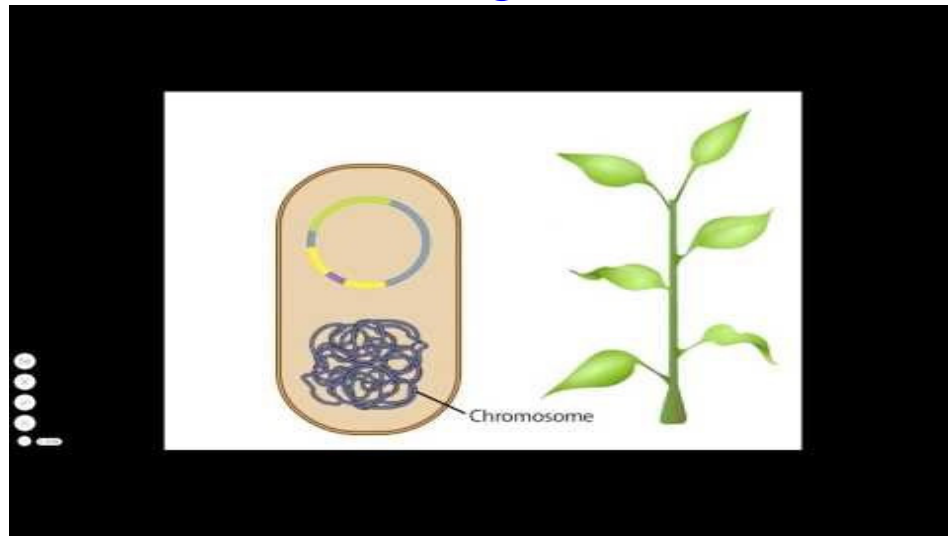
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Insulin, a peptide hormone composed of two polypeptide chains connected via two disulfide bridges (in yellow, the third one is not involved in the linkage).

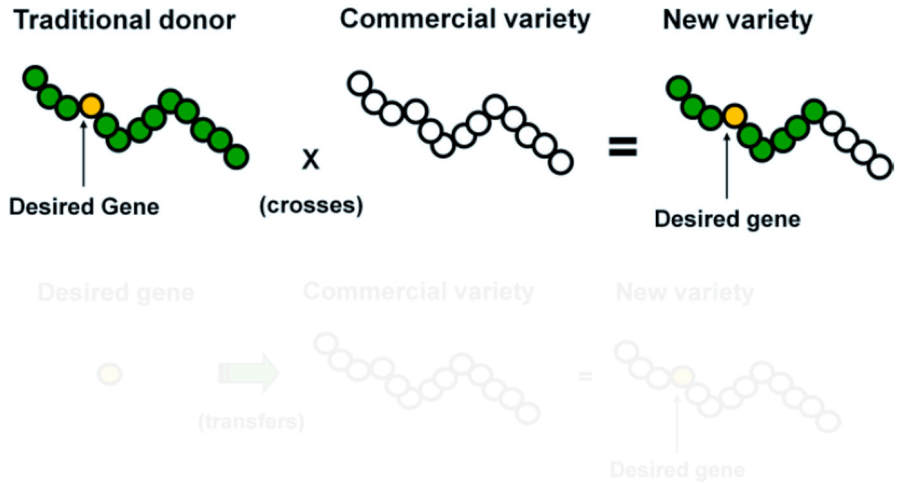
# The Story of Insulin



# Plant Transgenesis



# Breeding Vs. Genetic Engineering





## Genetically Modified Organisms (GMOs)

- Organisms carrying **modified genetic information** – either **own** or **foreign** (from another organism), enabling **targeted changes** in the organism and its use for **specific purposes**
- **GMOs**
  - plants
  - bacteria
  - animals

<http://www.gmo-compass.org/>

## Genetically Modified Plants

- resistance to **pests**
- **herbicide** resistance
- resistance to **drought**
- resistance to **cold**
- resistance to **salinity**
- more efficient **nitrogen utilization**
- increasing **nutritional quality**



<http://ipbo.vib-ugent.be/>

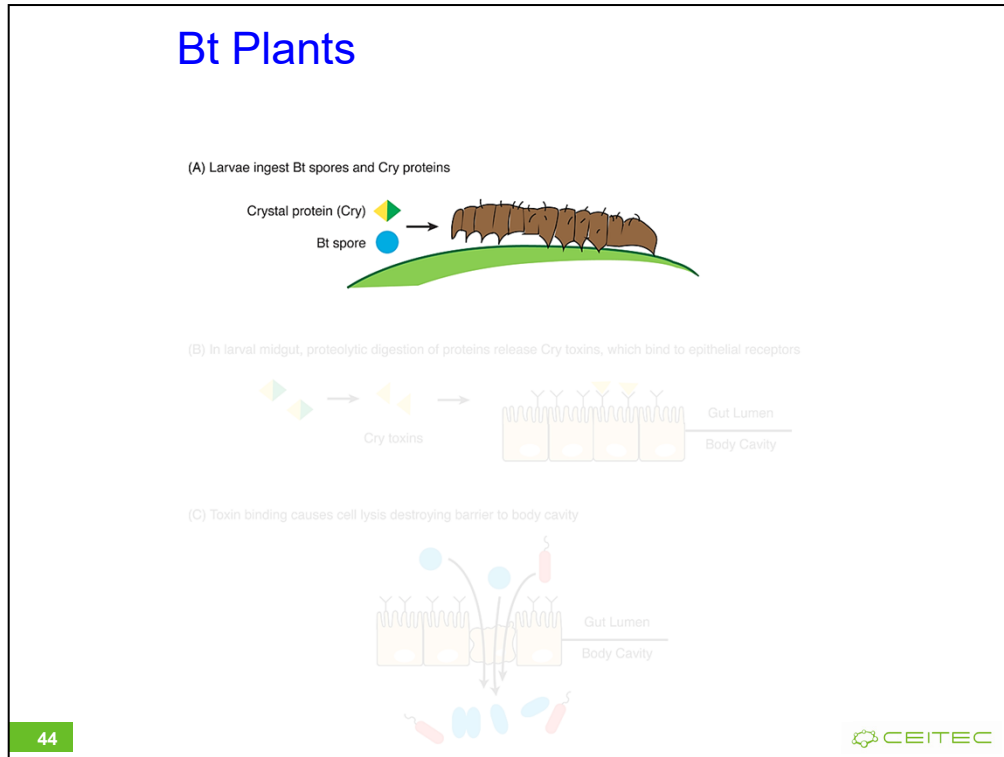
## Bt Plants

- resistance to **insect pests**
- corn, cotton, rice
- genes from *Bacillus thuringiensis* (**Bt**)
- Expression of crystalline delta-endotoxins - **Crystal (Cry) proteins**
- increasing **yields**, **reducing** the amount of **chemical sprays**



© Gary Muehlbauer  
European corn borer damage and fungal infection in non-Bt (left) and Bt hybrids (right)

## Bt Plants



When the Cry protein reaches the gut, it is partially degraded, releasing a smaller and potentially toxic part of the protein [6]. But this toxin will only be active if it finds the right matching protein receptor sticking off the cells lining the gut of a larval insect. This is the most important aspect of the Cry toxin mechanism. Much in the same way that a certain key will only open a certain lock, the Cry toxin can only exert its toxic effect on a particular cell receptor. Consequently, the toxin tends to only impact insects within a particular taxonomic order.

Once the toxin is bound, the process is fairly straightforward. The toxin recruits other Cry toxins to the same cell and together they form a hole in cell's membrane that ultimately causes the cell to burst [6]. The cumulative effect of this happening to many cells is the irreversible destruction to the midgut membrane, compromising the barrier between the body cavity and gut. Without this barrier, *Bt* spores and other native gut bacteria can infiltrate and grow within the nutrient-rich body of the insect [4-5].

What makes *Bt* such a great candidate for pesticide and GM applications is that while these Cry toxins are highly effective against insects, they have been shown to be safe for consumption by mammals. Tests by the EPA have demonstrated that Cry proteins, like any other benign dietary protein, are very unstable in the acidic stomach environment. Furthermore, an oral toxicity test, which involves giving mice exceptionally high doses of purified toxic *Bt* proteins, showed no significant health impacts. In their 2001 reassessment of several *Bt* Cry proteins, the EPA concluded from these findings that "there is reasonable certainty that no harm will result from aggregate exposure to the U.S. population, including infants and children, to the Cry1AB and Cry1F proteins and the genetic material necessary for their production." Similar conclusions were drawn about the Cry1Ac protein of *Bt* cotton [7]. Other mouse studies on have shown that even high doses of truncated Cry proteins, such that only the toxic region is conserved, have no deleterious effects [8]. A paper in Annual Review of Entomology from 2002 also makes the strong point that, in addition to no demonstrated toxicity of *Bt* toxins, their use provides important health benefits to livestock and humans by preventing certain insect-caused crop diseases that produce toxic and carcinogenic compounds [13].

## Ht Plants

- resistance to systemic herbicides
- glyphosate
  - interferes with the synthesis of aromatic amino acids; animals without the appropriate enzymatic apparatus = harmless
  - blocks the enzyme 5-enolpyruvylshikimate-3-phosphate synthase (EPSPS) in chloroplasts – affects green plants
  - ineffective for bacterial EPSPS - evolutionarily divergent
  - soya, maize, sugar beet, canola, cotton, alfalfa - added enzyme for tolerance
  - company Bayer (Monsanto), trade name Roundup

## Ht Plants

- resistance to systemic herbicides
- glufosinate (phosphinothricin)
  - prevents processing of ammonium - toxic
  - *Streptomyces hygroscopicus* synthesizes and transforms it: acetylation by the enzyme phosphinothricin acetyltransferase – coding gene isolated in 1987 - named *bar*
  - trade names: Basta, Liberty, Finale, Radical ...

## Multiresistant Plants

- Bt resistance + herbicide
- multiresistant corn - the majority of total production in the USA
- example of multiresistant corn:
  - three Bt genes for resistance to air pests
  - three Bt genes for resistance against soil pests
  - two genes for herbicide resistance

## Disease-Tolerant Plants

- **viruses** - no chemical agents available
- gene encoding **non-infectious viral envelope** protein - increases resistance to viral infection
  - **banana; papaya** - Hawaii, Southeast Asia
  - **cassava** - a basic food ingredient for more than **500 million people** + animal feed



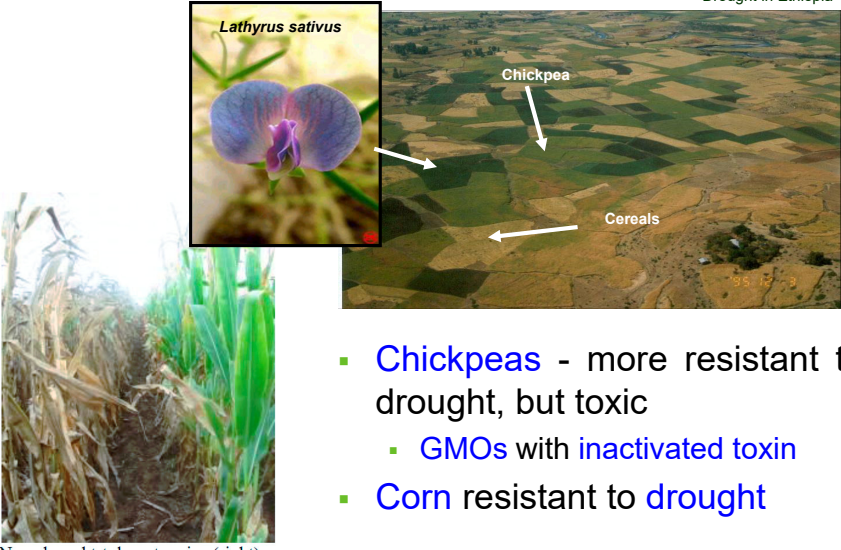
Left: Papaya with Papaya ringspot disease  
Right: Biotech Papaya resistant

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Hlízy manioku (cassava) tvoří základní potravinovou složku pro více než 500 milionů lidí. Rovněž se využívá jako krmivo - zkrmuje se v podobě maniokové moučky hlavně prasatům, skotu, ovcím a kozám.



## Disease- and Stress-Tolerant Plants



Drought in Ethiopia

*Lathyrus sativus*

Chickpea

Cereals

- Chickpeas - more resistant to drought, but toxic
  - GMOs with inactivated toxin
- Corn resistant to drought

New drought-tolerant maize (right) needs less water.

CEITEC

Hrachor – *Lathyrus sativus*

Cizrna – Chickpea

Obiloviny - Cereals

## Nitrogen Use Efficiency

- use of nitrogen from fertilizers
  - rice with gene from barley - 3x higher nitrogen utilization under oxygen deficiency



The effect of Nitrogen Use Efficiency (NUE) in rice growth with reduced N applications. Left: rice engineered

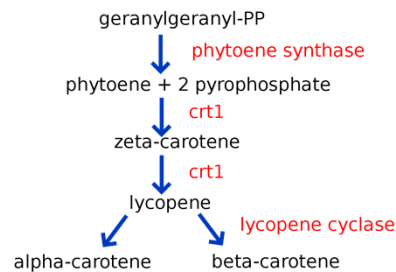
50

When crops are supplied with excess nitrogen fertilizer to gain maximal yields the excess nitrogen is converted into the gas nitrous oxide (N<sub>2</sub>O) and also leaches into rivers. N<sub>2</sub>O has 300x the Global Warming Potential of CO<sub>2</sub> and nitrogen fertilizer runoff creates marine dead zones, such as in the Gulf of Mexico at the mouth of the Mississippi river. Crops that have the ability to grow well with less nitrogen, because of enhanced uptake or similar characteristics, result in less N<sub>2</sub>O release and less N runoff. This lessens the effect of fertilizer nitrogen on global warming and lake and marine pollution.

## Improved Nutrition Value

- **Golden rice**

- several genes from maize encoding enzymes for the biosynthesis of  $\beta$ -carotene (precursor of vitamin A)



- **Canola and Soybean**

- improved oil properties: stable, resistant to high temperatures, long storage



Golden Rice  
شیراز

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While most biotech crops have characteristics that enhance their cultivation, those with enhanced consumer characteristics are being developed. For example many children in SE Asia develop blindness because of a deficiency of vitamin A. Golden rice is engineered with genes from maize to be high in the precursor of vitamin-A that when eaten is converted to vitamin-A in order to prevent blindness in developing countries. High oleic soybean and canola oil are now available. Oil with this fat profile is more stable, allowing for greater heat tolerance and longer shelf life.

## GMO Animals

- Transgenic cats
  - lentiviruses are sensitive to restriction factors
    - specific restriction factor: rhesus macaque TRIMCyp + eGFP
  - uniform expression, no mosaicism and no silencing in F1 generation
  - lymphocytes of transgenic animals resistant to replication of FIV



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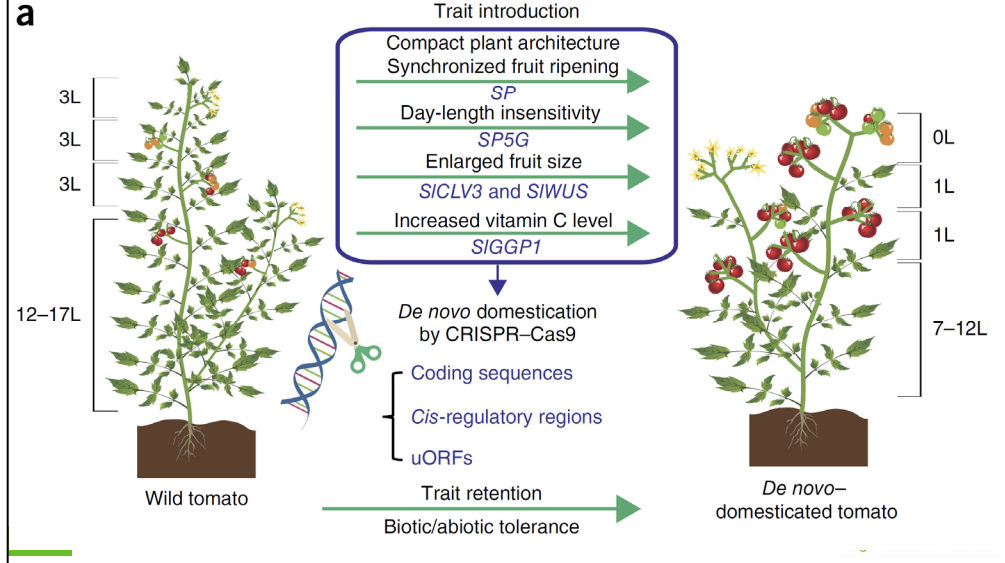
Wongsrikeao et al., 2011, Nature Methods

**Feline immunodeficiency virus (FIV)**, the “cat AIDS” is a [Lentivirus](#) that affects [cats](#) worldwide, with 2.5% to 4.4%<sup>[1][2]</sup> of [felines](#) being infected (Wikipedia)

# Osnova

- Lékařství
  - Molekulární diagnostika
  - Individualizovaná medicína
  - Genová terapie
- Biotechnologie
- Geneticky Modifikované Organismy
  - Transgenoze
  - Editování genomu

# Gene Editing in Plant Domestication



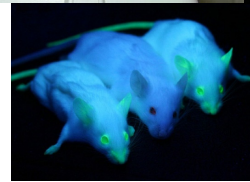
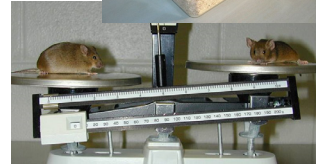
# Osnova

- Lékařství
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  - Editování genomu
- Modelové organismy

# *Mus musculus*

## house mouse

- **Low requirements** for area
- Relatively **large number of offspring** (3-14, 6-8 on average)
- **Genome size is close to the size of human genome** (about 3000 Mbp), the number of genes as well (about 24K)
- **20 chromosomes** (19+1)
- **Suitable for a wide range of physiological experiments** (anatomical and physiological similarity to human)
- Possibility to **obtain (quite easily) KO mutants and transgenic lines**



More info about mouse at <http://www.informatics.jax.org/greenbook/index.shtml>.



# Mus musculus

## house mouse

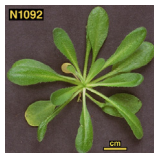
- Genome known since 2002 (<http://www.ncbi.nlm.nih.gov/projects/genome/assembly/grc/mouse/>)

The screenshot shows the 'Mouse Genome Overview' page from the Genome Reference Consortium. The page features a navigation bar with links like 'GRC Home', 'Data', 'Help', 'Report an Issue', 'Contact Us', 'Credits', and 'Curators Only'. Below the navigation, there are tabs for 'Mouse Overview', 'Mount Issues under Review', 'Mouse Assembly Data', and 'Report a problem'. The main content area is titled 'Mouse Genome Overview' and includes a paragraph explaining the GRC's goal to provide the best possible reference assembly for mouse. It mentions the use of multiple representations (alternate loci) for complex regions and the release of regional fixes (patches). A diagram shows 19 chromosomes (1-18, X, Y) with red and orange markers indicating regions with alternate loci and patches, respectively. A text box highlights the 'Next assembly update' for GRCm38.p1, scheduled for March 2013. The page also includes a 'Getting Data' section with links to FTP sites for GRCm38.p1, GRCm38, and MGSv37. On the right side, there is a 'GRC Blog' section with recent posts and a 'Recently Resolved Mouse Issues' section listing specific assembly problems and their resolutions.

# *Arabidopsis thaliana*

mouse-ear cress

- **Low requirements** for cultivation area
- High **number of seeds** (20.000 per plant and more)
- **Small and compact genome**, (125 MBp, about 25.000 genes, average size 3 kb)
- **5 chromosomes**
- Suitable for **wide range of physiological experiments**
- **High natural variability** (approximately 750 ecotypes (Nottingham Arabidopsis Seed Stock Centre))



Columbia 0

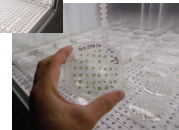


Landsberg 0



Wassilewskija 0

<http://seeds.nottingham.ac.uk/>



# Arabidopsis thaliana

mouse-ear cress

- Genome known since 2000 (<http://www.arabidopsis.org/>)

The screenshot shows the homepage of the Arabidopsis Information Resource (TAIR). The page features a navigation menu with links for Home, Help, Contact, About Us, Login/Register, Search, Browse, Tools, Portals, Download, Submit, News, and ABRIC Stocks. The main content area is titled "The Arabidopsis Information Resource" and provides a detailed description of the database, including its scope and the types of data it contains. A prominent banner in the center encourages users to "try our new online submission form" and submit molecular function, biological process, or interacting partner information for a gene. The right sidebar contains several news items, including "Breaking News" about a new set of confirmed T-DNA lines, "New from ABRIC Education and Outreach" regarding a re-designed website, and a "2012 MAASC Report Now Available" from the Arabidopsis Steering Committee. The footer of the page includes the CEITEC logo.

# Osnova

- Lékařství
  - Molekulární diagnostika
  - Individualizovaná medicína
  - Genová terapie
- Biotechnologie
- Geneticky Modifikované Organismy
  - Transgenoze
  - Editování genomu
- Modelové organismy
- Principy PCR



# Polymerase Chain Reaction

# Klíčové koncepty

- Techniky využívající pokročilé genetické a genomické přístupy zásadním způsobem mění naše možnost dosahovat požadovaných cílů v medicíně i zemědělství.
- Možnost programovatelné editace genomu slibuje zásadní obrat v léčbě zejména dědičně podmíněných chorob a ve šlechtění nových odrůd i ras
- Je nezbytná přísná kontrola s jasně nastavenými pravidly pro všechny, ale nikoliv úplný zákaz

# Diskuse