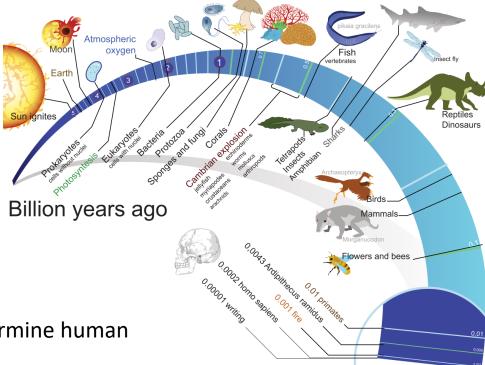
MUNI MED

Evolutionary medicine

Petr Müller

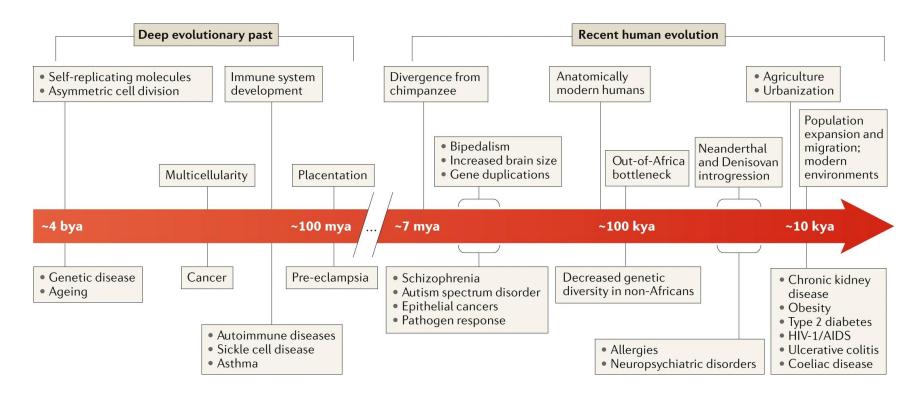
Molecular and Cellular Pathophysiology

Evolutionary medicine is the study of how evolutionary processes have produced human traits/disease and how evolutionary principles can be applied in medicine.



- Timeline of evolutionary events that determine human diseases
- Evolution and non-genetic adaptation to environmental changes - lifestyle changes and the effect of cultural evolution
- Evolutionary pressure and adaptation in other animal species
- Evolutionary trade-offs and and civilization diseases

A timeline of evolutionary events \rightarrow



A timeline patterns of human disease risk \rightarrow

Evolutionary medicine and genetic diseases

How evolutionary medicine explains complex genetic diseases

- 1. natural selection does not result in perfect bodies but operates on relative reproductive fitness
- 2. mismatch between our biological legacy and our modern environments
- **3.** trade-offs, the idea that there are combinations of traits that cannot be simultaneously optimized by natural selection
- 4. evolutionary conflicts. Traits expressed by complex metazoans are a balanced compromise between different genetic elements and bodily systems

NATURE REVIEWS | GENETICS

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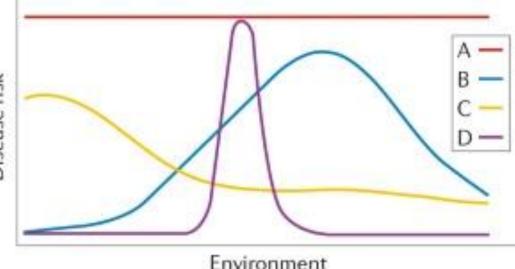
The influence of evolutionary history on human health and disease

Mary Lauren Bentono^{1,2}, Abin Abraham^{3,4}, Abigail L. LaBella⁵, Patrick Abbot⁵, Antonis Rokas^{6,1,3,5} and John A. Capra^{6,1,5,6}

The evolutionary necessity of disease / the impact of environment

Reaction norms

Representations of how the expressed phenotype for a genotype varies in response to a range of environments.



genotypes lead to disease in all environments Most diseases fall between these extremes (lines B and C) specific pairing of environment and genotype

> EVOLUTIONARY MEDICINE

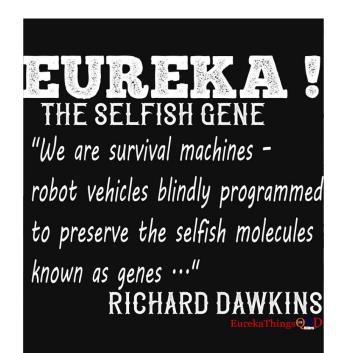
Viewing disease through the lens of evolution provides a flexible and powerful framework for defining and classifying disease.

Gene-centered view of evolution

natural selection does not result in perfect bodies but operates on relative reproductive fitness

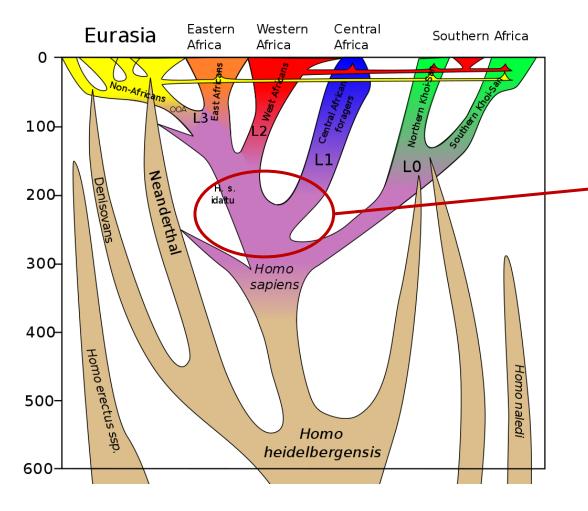
"Selfish gene theory"





- Altruism, cooperation, suicide
- Transposons, genetic waste information
- Sexual selection vs. Natural selection

Interbreeding between archaic and modern humans



Higher genetic diversity cohabitation of non-relatives cooperation

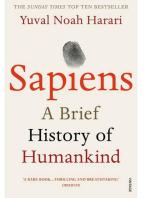
HLA-B*73 introgressed from Denisovans into modern humans in western Asia

Tibetan people EGLN1 and EPAS1 gene variant, associated with hemoglobin concentration

A model of the phylogeny of *H. sapiens* over the last 600,000 years (vertical axis).

Cultural evolution

is the idea that human cultural change—that is, changes in socially transmitted beliefs, knowledge, customs, skills, attitudes, languages, and so on—can be described as a Darwinian evolutionary process





Slaves to wheat: How a grain domesticated us

Unlike animals, the survival of humans is currently much less determined by their genetic information.

Much more important to human evolutionary fitness has become information obtained non-genetically

Neolithic revolution, cooperation and cultural evolution



THE LANCET

Dietary carbohydrate intake and mortality: a prospective cohort study and meta-analysis

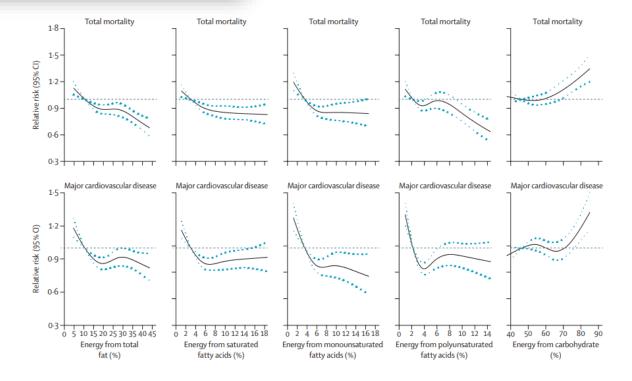
Sara B Seidelmann, Brian Claggett, Susan Cheng, Mir Henglin, Amil Shah, Lyn M Steffen, Aaron R Folsom, Eric B Rimm, Walter C Willett, Scott D Solomon



Associations of fats and carbohydrate intake with cardiovascular disease and mortality in 18 countries from five continents (PURE): a prospective cohort study

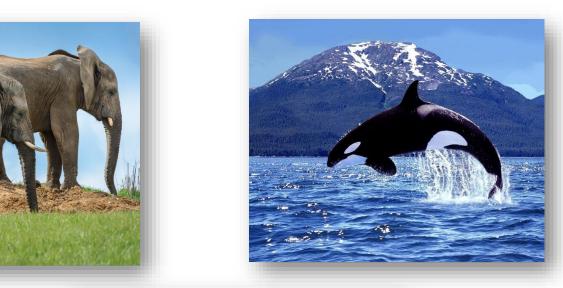
Mahshid Dehghan, Andrew Mente, Xiaohe Zhang, Sumathi Swaminathan, Wei Li, Viswanathan Mohan, Romaina Iqbal, Rajesh Kumar,





Mechanisms of evolutionary adaptations in different animal species The traits related to common human diseases

- Cancer
- Ageing
- Pathogen/infection resistance

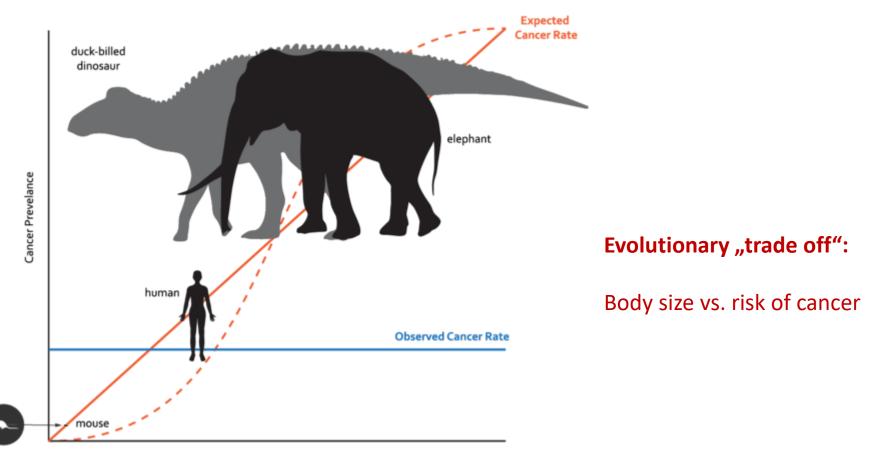






Cancer and Peto's paradox

- the incidence of cancer does not appear to correlate with the number of cells in an organism
- In order to build larger and longer-lived bodies, organisms required greater cancer suppression.



Lifespan x Body Mass

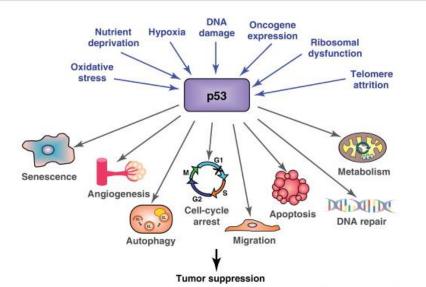
Gene Quantity in Cancer

HUMANS

VS.

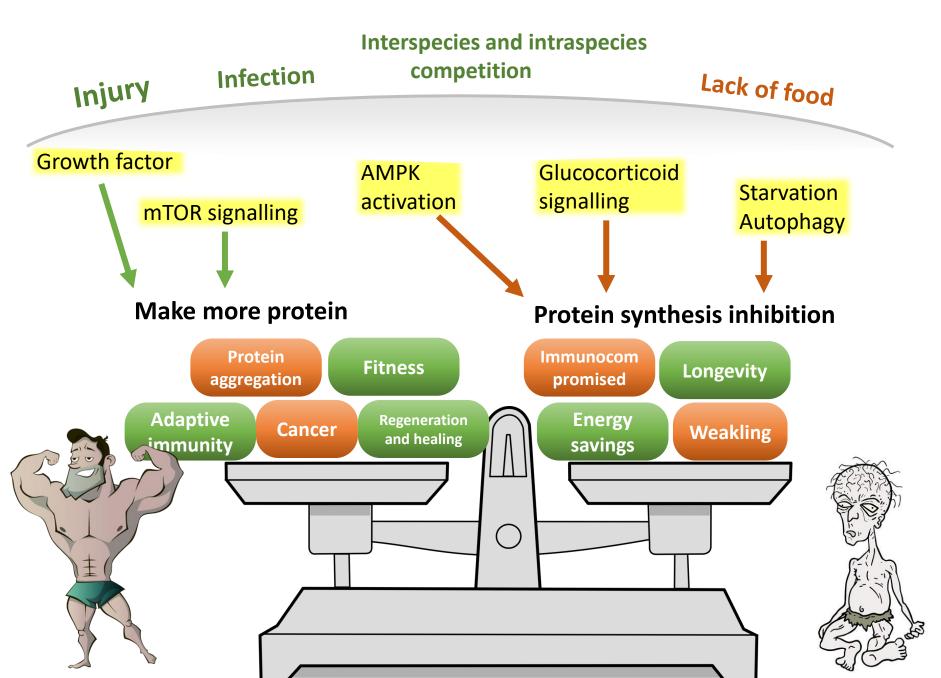


ELEPHANTS

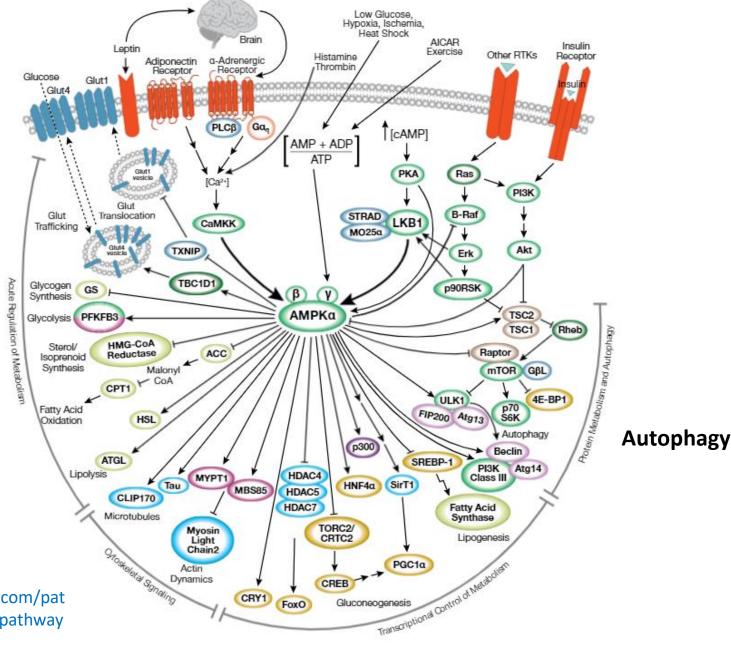


Mice altered to express "always-on" active TP53 exhibited increased tumor suppression ability, but also showed signs of premature aging. (TP53 cannot be the only explanation)

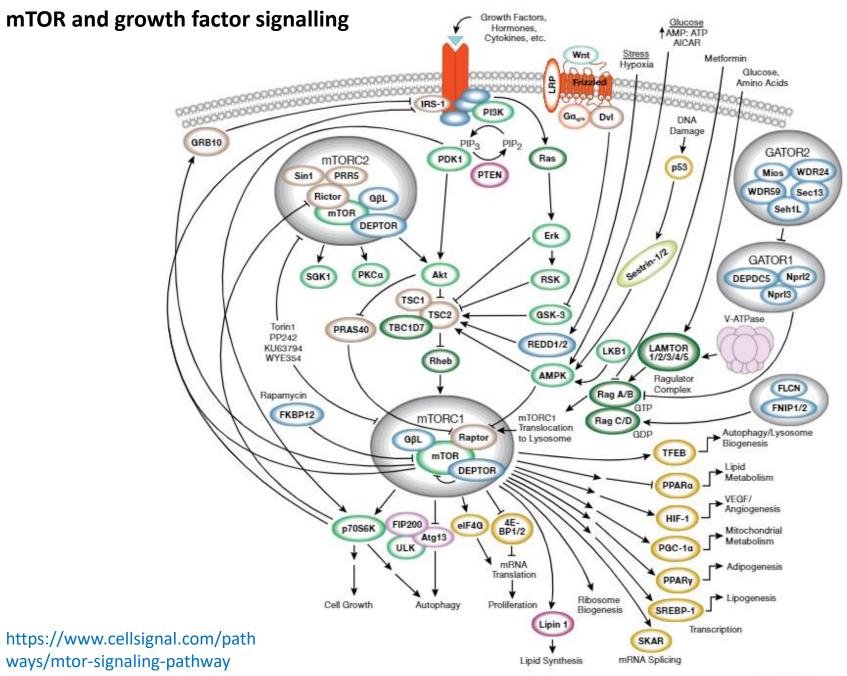
Balance of protein production and its regulation



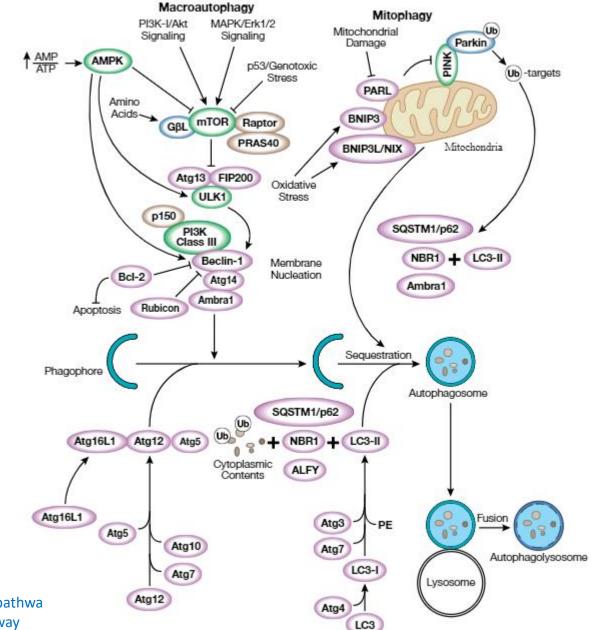
AMPK signalling



https://www.cellsignal.com/pat hways/ampk-signaling-pathway



Autophagy

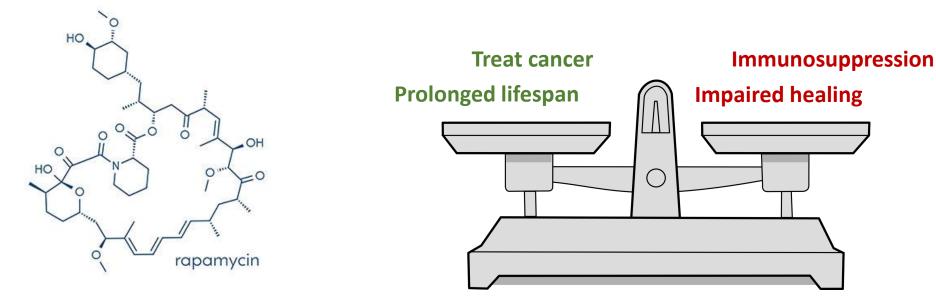


https://www.cellsignal.com/pathwa ys/autophagy-signaling-pathway

How can we affect protein homeostasis?

- Georges Nógrády was trying to understand why the inhabitants of Easter Island, despite walking around barefoot
- The Ayerst Pharmaceuticals team was able to identify a new antifungal compound in the soil samples that was produced by the bacterium Streptomyces hygroscopicus
- Identification of the mTOR Signaling Network
- Rapamycin's eventual development into a clinical compound (Rapamune), used to prevent organ transplant rejection and treatment for some cancers

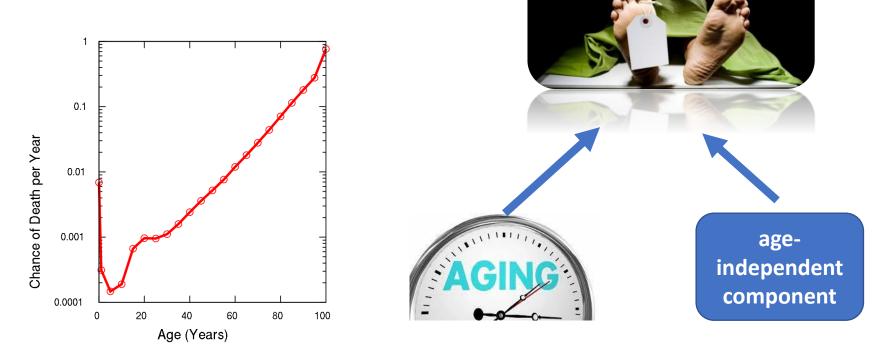




https://www.bio-rad-antibodies.com/blog/history-of-rapamycin.html

Gompertz–Makeham law of mortality

Estimated probability of a person dying at each age, for the U.S. in 2003. Mortality rates increase exponentially with age after age 30.



Probability of death

The Gompertz–Makeham law states that the human death rate is the sum of an **age-dependent component** (the Gompertz function, named after Benjamin Gompertz), which increases exponentially with age and an **age-independent component** (the Makeham term, named after William Makeham).

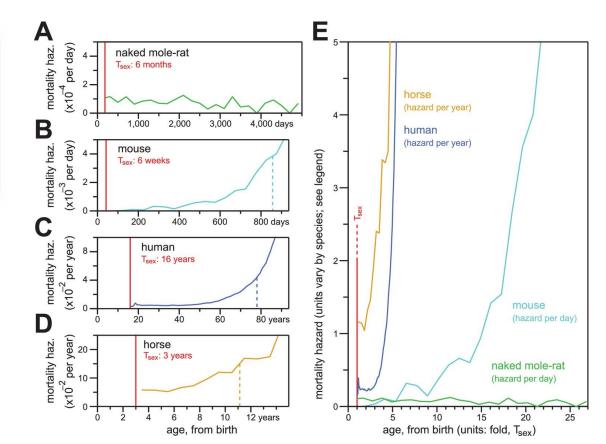
Naked mole rats defy the biological law of aging (Heterocephalus glaber)



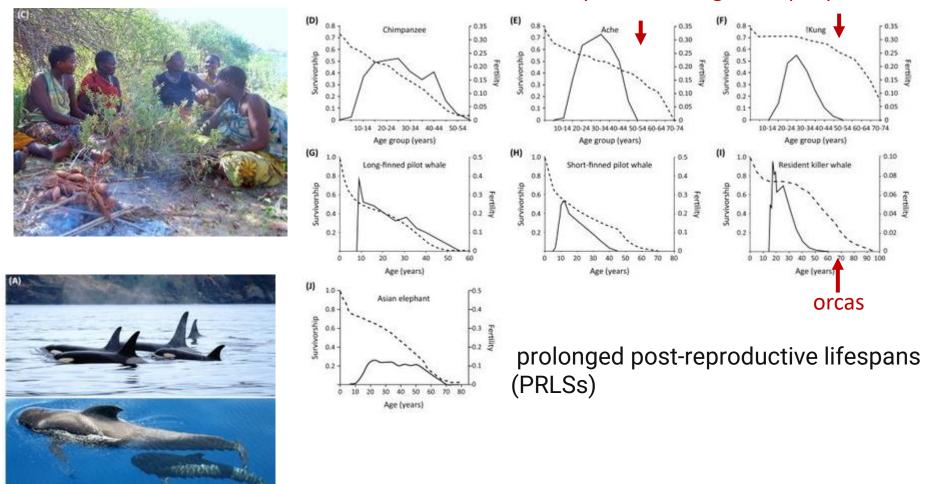


In contrast to the mortality hazards of other mammals, which increased with chronological age, the mortality hazard of naked mole-rats remained constant.

- rarely get cancer
- resistant to some types of pain
- survive up to 18 minutes without oxygen.



The evolution of prolonged life after reproduction



primitive indigenous people

Mechanisms of innate immunity

(fast but non-specific response)

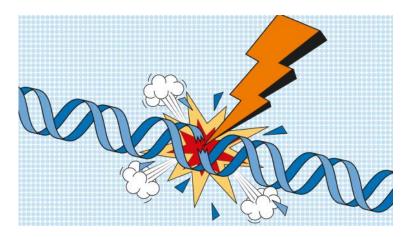
Detection of pathogenic microorganisms

- Membrane receptors
- Intracellular receptors of foreign nucleic acids
- Cytokine signalling

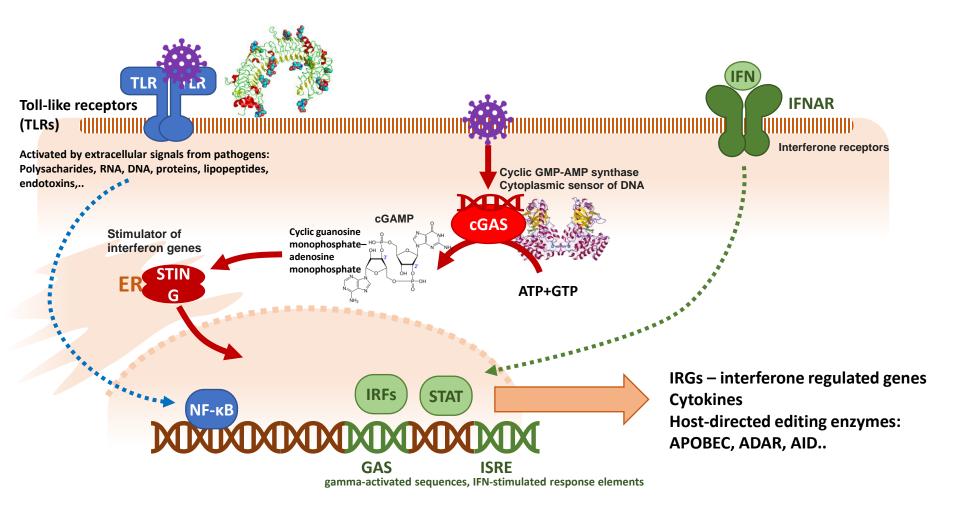
Intracellular signalling pathways



- Expression of cytokines
- Activation of specific immune response
- Elimination of microorganisms
- Use of gene

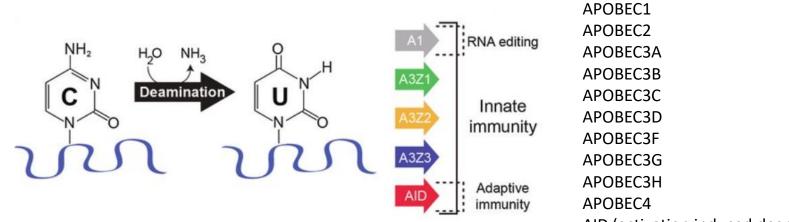


Mechanisms of innate immunity



APOBEC family members

- APOBEC ("apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like") is a family of evolutionarily conserved cytidine deaminases.
- Discovered due to their ability to eliminate HIV infection
- When misregulated, are a major source of mutation in numerous cancer types.
- AID is a part of adaptive immunity; it is responsible for hypermutation of variable immunoglobulin regions in lymphocytes



AID (activation induced deaminase)



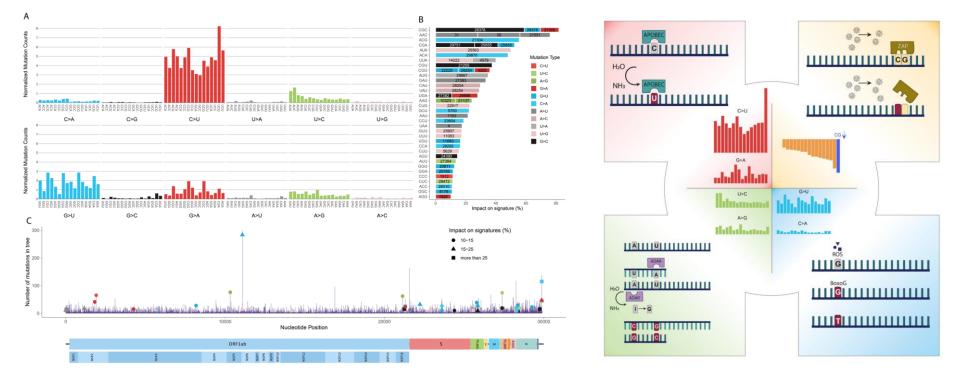
MDPI

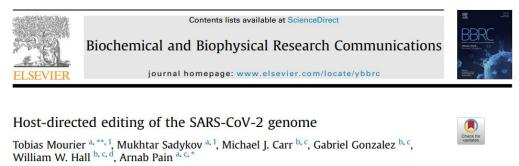
Article

The Mutation Profile of SARS-CoV-2 Is Primarily Shaped by the Host Antiviral Defense

Cem Azgari 🗓, Zeynep Kilinc 💿, Berk Turhan 💿, Defne Circi 💿 and Ogun Adebali *💿

The results suggest that the heterogeneous mutation patterns are mainly reflections of host (i) antiviral mechanisms that are achieved through APOBEC, ADAR, and ZAP proteins, and (ii) probable adaptation against reactive oxygen species.





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^b National Virus Reference Laboratory (NVRL), School of Medicine, University College Dublin, Belfield, D04 V1W8, Dublin, Ireland

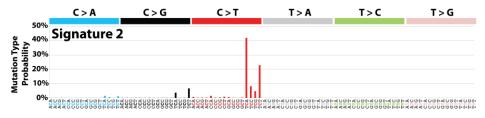
^c Research Center for Zoonosis Control, Global Institution for Collaborative Research and Education (GI-CoRE), Hokkaido University, N20 W10 Kita-ka, Sapporo, 001-0020, Japan

^d Global Virus Network (GVN), 801 W. Baltimore St., Baltimore, MD, 21201, USA

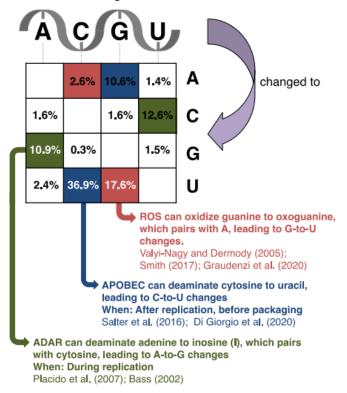


Signatures of Mutational Processes in Human

Signature 2 has been attributed to activity of the AID/APOBEC family of cytidine deaminases.



SARS-CoV-2 genome



578 | Nature | Vol 583 | 23 July 2020

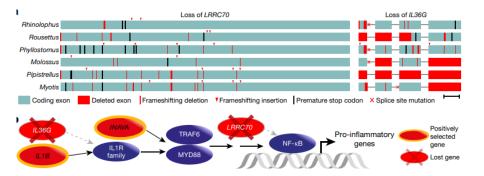
Six reference-quality genomes reveal evolution of bat adaptations

https://doi.org/10.1038/s41586-020-2486-3	David Jebb ^{12,225} , Zixia Huang ⁴²⁵ , Martin Pippel ^{1,225} , Graham M. Hughes ⁴ , Ksenia Lavrichenko ⁴ , Paolo Devanna ⁸ , Sylke Winkler ¹ , Lars S. Jermiin ^{4,6,7} , Emilia C. Skirmunt ⁴ , Aris Katzourakis ⁸ , Lucy Burkit-Gray ¹ , David A. Ray ⁹ , Kevin A. M. Sullivan ¹⁰ , Juliana G. Roscio ^{1,2,3} , Bogdan M. Kirilenko ^{12,3} , Liliana M. Dávalos ^{11,27} , Angelique P. Corthals ¹³ , Megan L. Power ⁴ , Gareth Jones ¹⁴ , Roger D. Ransome ¹⁴ , Dina K. N. Dechmann ^{33,627} , Andrea G. Locatelli ⁴ , Sébastien J. Puechmaille ⁴¹⁹ , Olivier Fedrig ²⁰³ , Erich D. Jarvis ^{203,123} , Michael Hiller ^{12,22663} , Sonja C. Vernes ^{5,22,2685} , Eugene W. Myers ^{13,24,2655} & Emma C. Teeling ^{4,2065}
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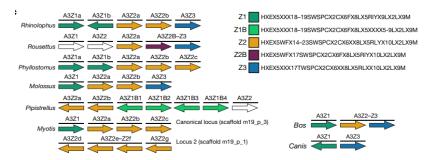
Article



Loss of genes in NF-KB signalling pathway



Expansion of the APOBEC3 gene locus



https://twitter.com/bat1kgenomes?s=20