

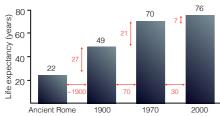
Aging



Aging is commonly characterized as a progressive, generalized impairment of function, resulting in an increasing vulnerability to environmental challenge and a growing risk of disease and death. It is also usually accompanied by a decline in fertility. Thus, aging is associated with major age-related losses in Darwinian fitness, posing the puzzle of why it has not been more effectively opposed by natural selection.

"It is remarkable that after a seemingly miraculous leat of morphogenesis, a complex metazoan should be unable to perform the much simpler task of merely maintaining what is already formed" (Williams, 1957)

How long shall we live?





Life expectancy at birth in developed countries

- US Census Bureau Middle Series: life expectancy in 2050 will be ~82 years for both sexes in the US
- US Social Security Administration: life expectancy of 78.1, 80.4 and 83.5 years for both sexes in 2066 on three alternative assumptions
- G7 Industrialized Countries: life expectancy in 2050 with a maximum of 90.9 in Japan and a minimum of 82.9 years in USA

Shall we live forever?

Maximum life span for the human species (unchanged in the last 100,000 years): 125 years

The longest-lived human being is Jeanne Calment (122.5 years), died in France, in August 1997

Maximum life span in other species:

Rat: 3 years Squirrel: 25 years Sheep: 12 years Turtle: 150 years Dog: 15-30 years Fly: 3 months Canary 15 years Bat 50 years

In animal studies, maximum life span is often taken to be the mean life span of the most long-lived 10% of a given cohort. By another definition, however, maximum life span corresponds to the age at which the oldest known member of a species or experimental group has died. Calculation of the maximum life span in the latter sense depends upon initial

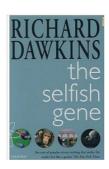
EXISTENCE OF A BIOLOGICAL CLOCK?

Why do we age?

Final part of the developmental program (aging selected because provides <u>advantage</u> to the species?)

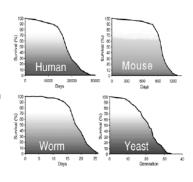
How do we age?

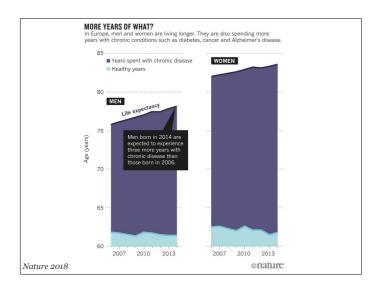
Exhaustion of the proliferative or functional capacity of all or some somatic cells (eg. in stem cells?)
Changes in biochemical composition of tissues (increased adipose tissue, lipofuscin deposit, increased ECM component cross-linking, increased glycation products)



Age-related diseases are degenerative in nature and compressed at the end of our life

- Increased mortality with age maturation
- Increased susceptibility and vulnerability to disease (centenarians live >90% of their lives in very good health and with high level of independence - marked morbidity compression toward the end of life)





Theories of Aging

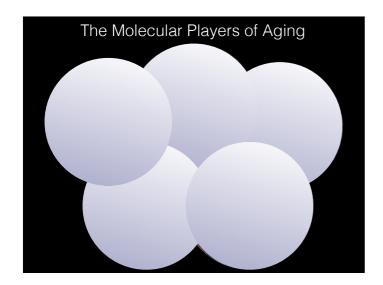
- Somatic mutation theory or Loose cannon theory or Free radical theory of aging. Damage produced by free radicals glucose, or other agents slowly disruy cellular macromolecules. This causes an age-related increase in somatic mutation and other forms of DNA damage.
- Telomere loss theory. A decline in cellular division capacit
 with age linked to the progressive shortening of telomeres a
 cells divide.
- Mitochondrial theory. Accumulation of mitochondrial DNA mutations with age.
- Altered proteins theory and waste accumulation theory.
 Accumulation over time of damaged proteins (e.g. Alzheimer: disease, Parkinson's disease, cataract, etc.). Linke to functional declines of proteasomes and chaperones
- Antagonistic pleiotropic theory. Pleiotropic genes exist having opposite effects on fitness at different ages: they are beneficial in early life, when natural selection is strong, but harmful at later ages, when selection is weak
- selected by natural selection, over successive generations they accumulate within the genome.
- Rate of living theory. Metabolic rate is inversely correlated with longevity. Smaller mammals tend to have high metabolic rates and thus tend to die at an earlier age than larger mammals.

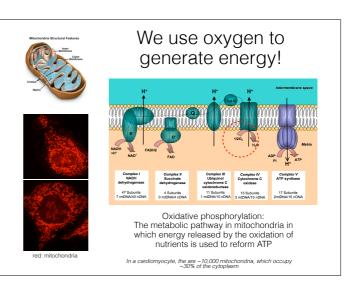
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- Weak link theory. A specific physiologic system (e.g. the neuroendocrine or the immune system) is particularly vulnerable over time and its dysfunction accelerates senescence of the whole organism



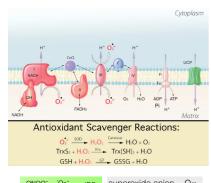
The Malmarks of Aging
 There is a simulated in the Salmarks described in the Salmar growns in statistic, before addition, adjusted, about an increase and control and produced and control and produced and control an

- Error catastrophe theory. Errors in DNA transcription or RNA translation eventually lead to genetic errors that
- Master clock theory. Aging is under genetic control (gene that controls telomere shortening? or cell division? or DNA repair?
- Disposable soma theory. Since the metabolic resources of an organism are limited (chiefly: energy), the organism should optimally allocate them between the maintenance and repair of its soma and the other functions that it must carry out in order to maximise its Darwinian fitness (growth, reproduction,...)
- 12. Combined network theories of aging. Multiplicity of aging mechanisms (e.g.: a gradual accumulation of mIDNA mutations might lead to a steady increase in the production of ROS and a gradual decline in energy production.









ROS are produced in multiple compartments:

- NADPH oxidases on the plasma membrane
- lipid metabolism in the peroxisomes
- cytosolic enzymes such as cyclooxygenases

0.2-2% of total oxygen consumption is funneled to ROS generation

The "free radical theory of aging" (Harman, 1956)

Aging and its associated degenerative diseases can be attributed to deleterious effects of free radicals on various cell components

Now better called "Oxidative stress theory of aging" (many ROS are not free radicals)

Mitochondria are the main source of ROS













Mitochondrial mutations and aging

- The mutation rate in mitochondria is 10-20 times faster than the nuclear DNA mutation rate
- Specific mutations in mitochondria could lead to defects in energy production and production of ROS by faulty electron transport
- Age-dependent declines in mitochondrial function are seen in many species including humans
- Inherited mitochondrial DNA variants are associated with aging and longevity (the J haplogroup is more represented in centenarians in Northern Italy than in younger subjects)
- Knock-in mice expressing a proof-reading deficient form of a nuclear-encoded mitochondrial DNA polymerase exhibit an increased mitochondrial mutation rate, appearance of a number of age-related phenotypes - including hair loss, kyphosis, and reduced fertility -, and shortened life span

C. elegans mutants, oxidative stress and aging

isp-1 mutants are **long-lived** (missense mutation in a component of complex III of the respiratory chain in mitochondria)

A **systematic RNAi** screen sought to inactivate over 5600 random C. elegans genes screening for long-lived animals: ~15% of the identified genes regulate mitochondrial activity

mev-1 mutants (mutation in a subunit of complex II) have increased ROS generation and are **short lived**; mice heterozygous for mitochondrial SOD2: increased incidence of nuclear DNA damage and tumor formation

clk-1 mutants are **long-lived** (lack an enzyme required in the biosynthesis of ubiquinone (coenzyme Q), an electron acceptor for both complex I and II-dependent respiration - NB: although coenzyme Q is sold as a life-extending anti-oxidant, its withdrawal from the diet of w worms increases life span by 60%!!!

You can live longer if you have mutations that makes the mitochondrion less functional and thus able to generate lower amount of ROS



Indy (I'm not dead yet): 50% increase in life span. Indy encodes a protein with sequence homology to mammalian sodium dicarbocylate cotransporters, which import Krebs cycle intermediates into cells. Indy is expressed in the midgut and the fat body, the fly functional equivalent of mammalian liver and white adipose tissue. Indy mutations create a metabolic state similar to that found in dietary restriction.

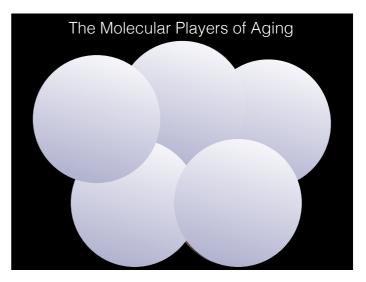
Mth (methuselah): 25% increase in life span. Family of the seven transmembrane spanning GTP-binding protein-coupled receptors (GPRC). The cognate ligand is the product of the stunted gene, encoding for a subunit of the F_1F_0 -ATP synthase of the electron transport chain (!)

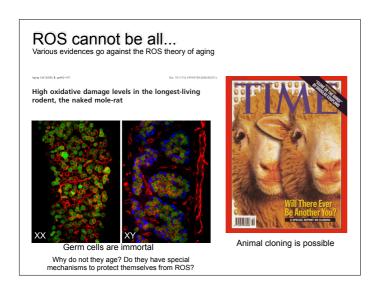


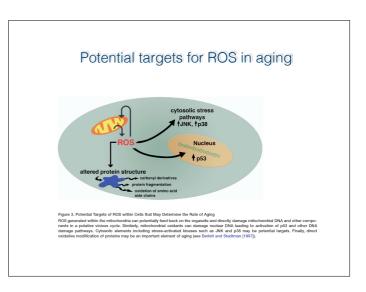
In humans, meta-analysis of randomized controlled trials showed that selenium and vitamin C have no effect while standard antioxidant supplementation (vitamins A and E and beta-carotene) actually increases mortality

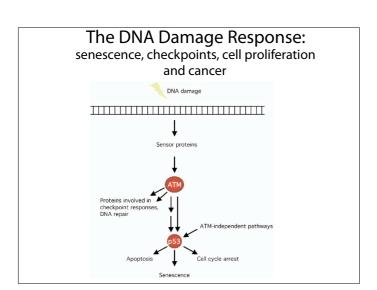


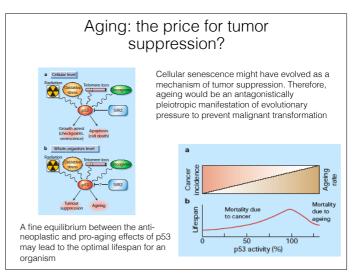


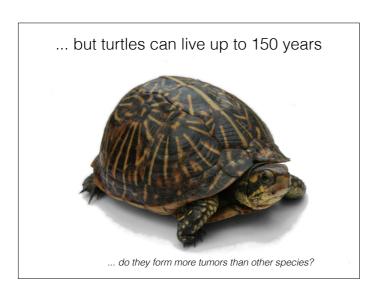


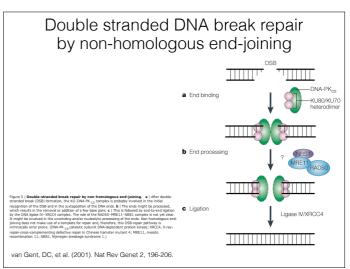


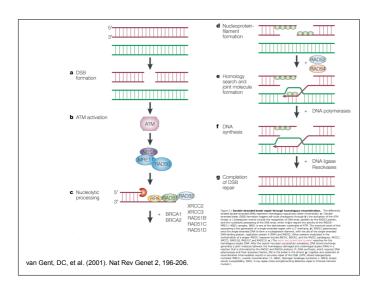












Human progeroid syndromes

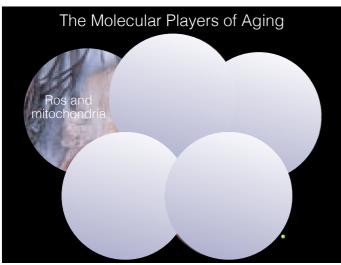
Table 1 Genetic instability syn	dromes				
Disease*	OMIM†	Gene	Function	Major phenotypes	
Werner syndrome	277,700	WRN (ref. 6)	Helicase/exonuclease	Skin atrophy, cataracts, diabetes melitus, osteoporosis, hypogonadism, afterosclerosis, cancer predisposition? ²	
Rothmund-Thomson syndrome	268,400	RecQ4 (ref. 50)	Helicase	Polklioderma, photosensithity, skeletal abnormality, cataracts, cancer predisposition (osteosarcoma) ³¹	
Cockayne syndrome, type A	216,400	CKN1 (ref. 52)	WD repeat protein	Neurodegeneration, skeletal abnormality (widened face), impaired sexual	
Cockayne syndrome, type B	133,540	ERCO6 (ref. 53)	Helicase	development, photosensitivity ⁵⁴	
Ataxia telangiectasia	208,900	ATM (ref. 55)	Kinase	Carabellar dysfunction, sensitivity to ionizing radiation, cancer predisposition ⁹⁵	
Nijmegen breakage syndrome (Atavia, tolandischaria cariant)	251,260	NSB1 (ref. 57)	Unknown	Microcephaly, growth retardation, immunodeficiency, cancer predispositio	

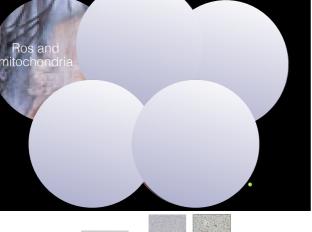
Werner syndrome

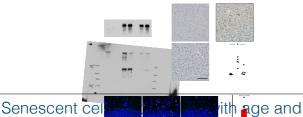




Werner syndrome is genetic recessive disorder. It is a type of progeria disease that occurs in adults ages twenty to thirty. People who are affected start to age rapidly beginning in their twenties and thirties and look as though they are twenty or more years older that what they actually are. Along with looking older patients develop other types of diseases and disorders that occur with normal aging. Werner's strikes about three in every 1 million people worldwide, although it is slightly more common in Janan.







contribute to a Local clearance of senescent cells attenuates the development of post-traumatic osteoarthritis and creates

Ok Hee Jeon^{1,8}, Chaekyu Kim^{1,2,8}, Remi-Martin Laberge^{3,4}, Marco Demaria^{3,5,0}, Sona Rathod¹, Alain P Vasserot¹, Jae Wook Chung¹, Do Hun Kim¹, Yan Pooh¹, Nathaniel David¹, Darren J Baker⁴ Jan M van Deursen⁶, Judith Campisi^{3,7} & Jennifer H Elisseeff¹

a pro-regenerative environment

Clearance of senescent glial cells prevents tau-dependent pathology and cognitive decline

Cellular senescence

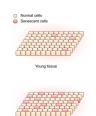
- Normal human cells have a limited ability to proliferate in vitro (Hayflick, 1965)
- Growth potential of a primary cell declines 0.2 population doublings per year of life
- Correlation between the number of senescent cells in vivo and age of
- Cells from progeria syndrome patients have limited doubling potential
- Association with several molecular
- Overexpression of telomerase overcomes senescence; overexpression of ras induces senescence

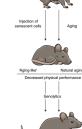
TABLE 1. Selected Alterations in Cell Phenotype with the Onset of Senescence

66, 67 69, 37

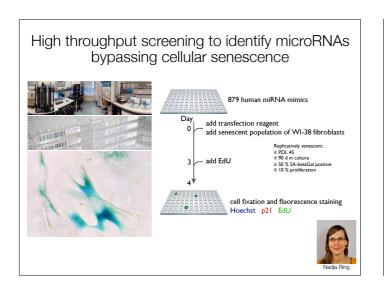
Senolytic drugs contrast aging phenotypes

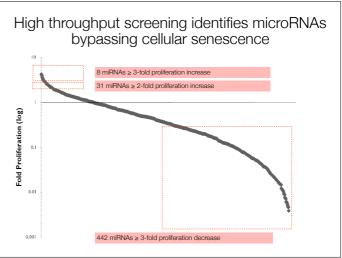
- Senolytic drugs: drugs that preferentially kill senescent cells
- In old mice, or mice 'aged' with senescent cells, as senolytic drugs (dasatinib and quercetin) increase remaining lifespan by 36%, enhance healthspan, reduce frailty and delay age-related diseases

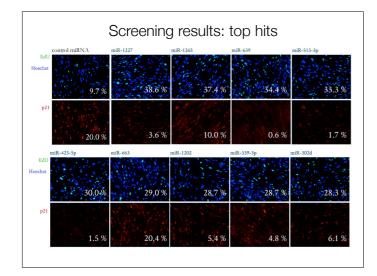


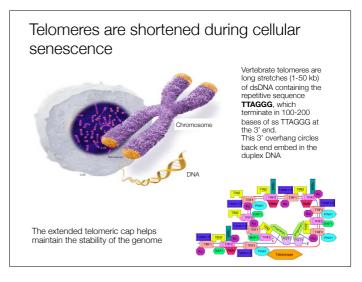


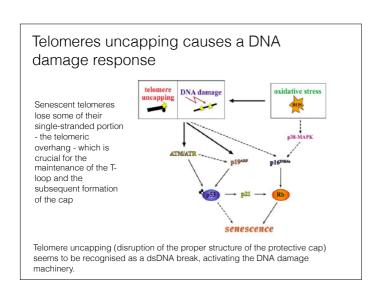
Senolytics improve physical function and increase lifespan in old age, Nature Medicine, August 2018

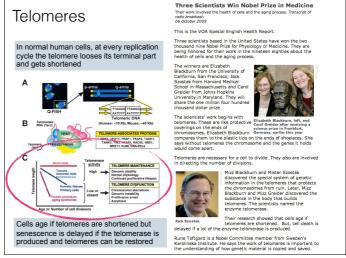














Nutritional Control of Reproductive Status in Honeybees via DNA Methylation

Is longevity controlled by a genetic (biologic) program?



C. elegans: at 20°C it lives, on average, 17 days, with a maximum of 25 days; in conditions of high density and food, it transforms into a larval form (dauer), which that does not reproduce and lives 60 days



Social insects: queens and workers are born from the same eggs fertilized by the same drone; workers live a few weeks in summer and a few months in winter; queens live several years



Some animals (turtles, deep water fishes, american lobster) age very slowly; these animals show no limit to body mass increase

Epigenetic control of longevity and reproductive status





Fertile queens and sterile workers are alternative forms of the adult female honeybee that develop from genetically identical larvae following differential

Nutritional Control of Reproductive Status in Honeybees via DNA Methylation

R. Kucharski^{*}, J. Maleszka^{*}, S. Foret and R. Maleszka[±]

Other interesting organisms



Semelparous organisms (once-only reproducing species; e.g. Pacific salmon, marsupial male rat; also called "Big Bang animals"). Die immediately after mating. Most probably, the mechanism is not active, and due to the fact that natural selection has evolved that a massive effort is made to mobilize all available resources to maximize reproductive success, even if this leaves the adult so severely depleted or damaged that death ensues. This is most likely to occur where ecological circumstances decree that the chance of surviving to breed again are very small (an extreme example of the

"disposable soma theory").

Hydra. Show slow or negligible rate of senescence. These organisms are capable of undergoing complete regeneration from almost any part of their structure, implying that germ cells permeate the body that there is no true distinction between germline and somatic tissue.





The "genes of aging"

Mutation in single genes decrease life span

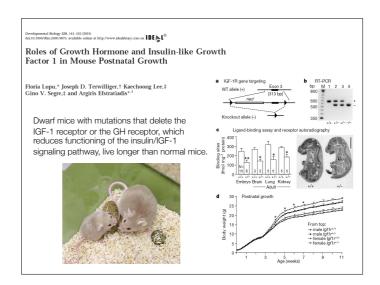
				Accelerated	
			Incressed Rate of	Fibroblast	
Mutant	Cellular Process Affected	Tissues Affected	Cancer?	Senescence?	Citations
Atm	DSB signaling/repair	Cerebellum, Gonad, Hematopoletic organs, Thymus	Yes	Yes	(fto et al., 200 Shilloh and I 2001)
Bub1b ^{HH}	Spindle assembly checkpoint	Bone, Lens, Skin, Gonad	No	Yes	(Baker et al., 2
BRCA1****(p63***	DSB repair, other	Bone, Eye, Heart, Intestine, Liver, Lymphocytic hyperplasia, Tostes +others	Yes	Yes	(Cao et al., 20
DNA-PKcs	NHEJ, other	Bone, Intestine	Yes	No	(Espejel et al.,
Ercc1, XPF	Nucleotide excision repair, crosslink repair, other	Liver (Ercc1, XPF) + Brain, Kidney, Skin, Spleen-Ercc1	No	Yes (Eroc1)	(McWhir et al. Tian et al., 2 Weeds et al
Ku80	NHEJ, other	Bone, Liver, Skin	No	Yes	(Vogel et al., 1
Dystunctional p53 (p53***)	DNA damage response	Bone, Hair, Lymphoid tissue, Skin	No	ND	(Tyner et al., 2
Dystunctional p53 (p44 Tg)	DNA damage response	Bone, Testes	No	Yes	(Major et al., 2
PASG/Lah	DNA methylation	Bone, Hair, Kidney, Skin, Thymus	No	Yes	(Sun et al., 20
PolgA	Mitochondrial DNA polymense	Bone, Hair, Heart, Hematopoletic organs, Skin, Tastes	No	ND	(Trifunovic et 2004)
Rad50 ^{US}	DSB repair	Hematopoletic organs, Testes	Yes	No	(Bender et al.
Terc	Telomere maintenance	Hematopoietic organs, Hair, Heart, Intentine, Myometrium, Skin, Testos	Conflicting results	Yes	(Espejel et al., 1
Terc/Atm	Telomere maintenance/ DSB signaling/repair	Bone, Brain, Hair, Hematopoletic organs, Intestine	No	Yes	(Wong et al.,
Terc/DNA-PKcs	Telomere maintenance/ NHEJ, other	Intestine, Various	No	ND	(Espojel et al.
Terc/Parp-1	Telomere maintenance/ DNA repair	Various	No	ND	(Espojel et al.
Terc/Ku80	Telomere maintenance/ NHEJ, other	Intestine, Various Bone, Endoorine,	No	ND	(Expejel et al.
Tero/Wrn	ro/Wm Telomere maintenance/ DNA repair		Yes	Yes	(Chang of al.,
Terc/Wrn/Birn	Telomere maintenance/ DNA repair	Bone, Endocrine, Gorsad, Hair, Intestine	Yes	Yes	(Du et al., 200
TopHibeta	Topolsomerase	Kidney, Lymphocytic inflirates, Pancreatic islets, Skin, Testes	No	ND	(Kwan et al.,)
Wm	DNA repair	None	No	Yes	(Lebel and Le 1998; Lomb al., 2000)
XPA/CSB	NER, transcription	Cerebellum	No	ND	(Mural et al.,)
Xpd ^{TTD}	NER, transcription	Bone, Hair, Ovary, Skin	Yes	ND	(de Boer et al.
Xpd ^{TT0} /XPA	NER, transcription	Bone, Hair, Skin	No	ND	ide Boer et al.

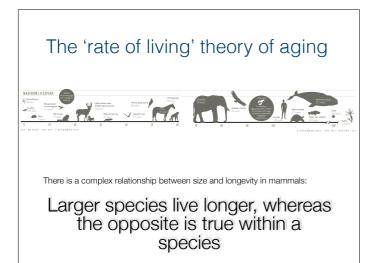
Mutants in the IIS pathway with extended lifespan in the mouse Ames and Snell Dwarf mice: miss the growth hormone-IGF-1 axis and other pituitary hormones due to mutations in the pit-1 gene Little mice: mutations in the GH-releasing hormone receptor KO mice for ligands (insulin, IGF1, IGF2) KO mice for receptors (IR, IFG1R, GHR) KO mice for immediate downstream signaling molecules (IRS proteins and other adaptor molecules including p66Shc) By Mus musculus Receptor 10F1R Receptor 10F1R

Cell growth and proliferation

and metabolic

activity





Gerontology. 2012;58(4):337-43. doi: 10.1159/000335166. Epub 2012 Jan 18

Healthy aging: is smaller better? - a mini-review.

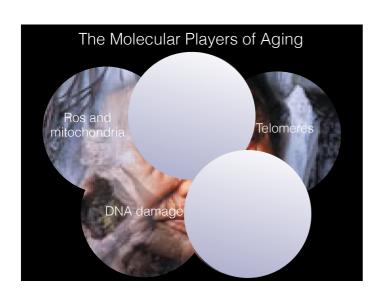
Bartke A.

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Abstract

A recent report of virtually complete protection from diabetes and cancer in a population of people with hereditary dwarfism revived interest in elucidating the relationships between growth, adult body size, age-related disease and longevity. In many species, smaller individuals outlive those that are larger and a similar relationship was shown in studies of various human populations. Adult body size is strongly dependent on the actions of growth hormone (GH) and the absence of GH or GH receptor in mice leads to a remarkable extension of longevity. Many mechanisms that may account for, or contribute to, this association have been identified. It is suggested that modest modifications of the diet at different ages may extend human healthspan and lifespan by reducing levels of hormones that stimulate growth.

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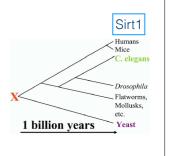


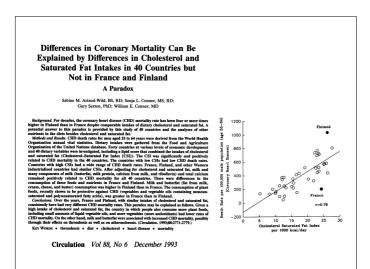
Sir2 genes and aging

Genetic studies indicate that the major genetic determinant of replicative life span in yeast is SIR2 (loss-of-function mutations shorten life span, increased gene dosage extend it).

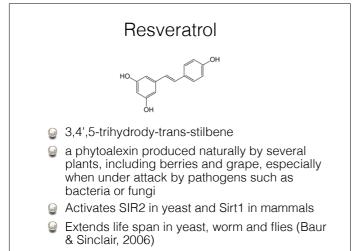
The SIR2 ortholog in *C. elegans* is a key determinant of life span in this animal

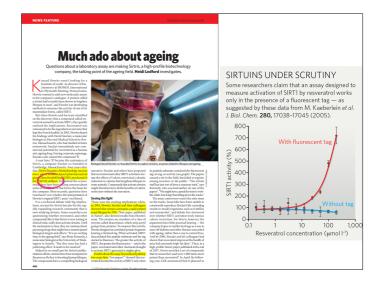
The fact that yeast and *C. elegans* diverged from a common ancestor about one billion years ago suggests that all the descendants of that ancestor (including mammals) will possess SIR2-related genes involved in regulating their life span.

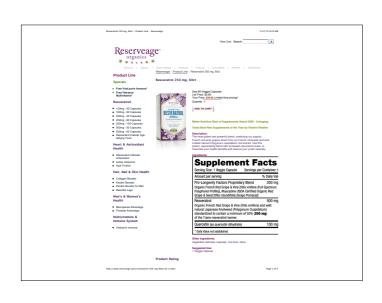


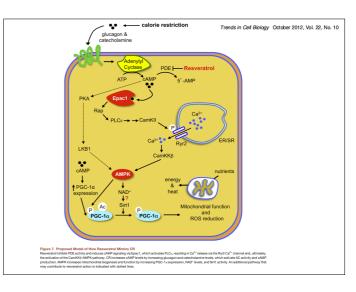




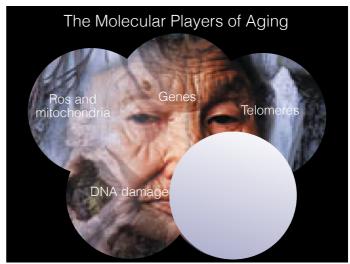




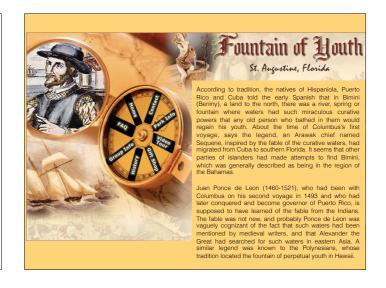








And so, what can we do?



Caloric restriction (CR) is the most effective environmental method to increase lifespan (and to prevent late-onset diseases!)

Dietary restriction extends lifespan in S. cerevisiae, C. elegans, D. melanogaster, rodents and primates.

CR = 60-70% of what an animal would eat at libitum

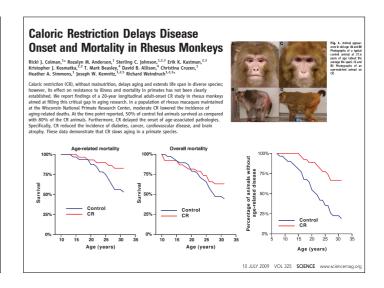
In rodents CR results in as much as a 50% increase in rodent longevity

Physiological effects of CR: acute phase followed by an adaptive period of several weeks to reach a stable, altered physiological state characterized by lower body temperature, lower blood glucose and insulin levels and reduced fat and weight.

The CR animals are more resistant to external stressors, including heat and oxidative stress; organs are typically smaller (except for the brain)

CR may represent an adaptation to scarcity in a boom and bust cycle; any organism that could slow aging and reproduction in times of scarcity and remain able to reproduce when food reappeared would enjoy a competitive advantage. Extremes examples are the formation of spored in microbes and dauer larvae in C. elegans

CR animals are resistant to disease, including cancer and infections





Caloric restriction (undernutrition without malnutrition)

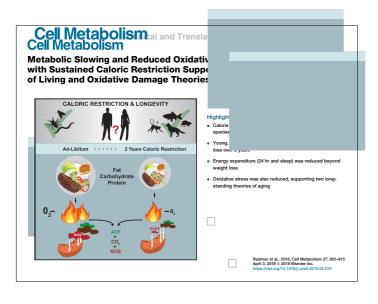
Slows multiple age-related changes, delays the onset of cancer and multiple other age-related pathologies, and extends life span

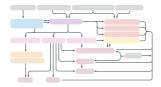
Periodic food deprivation (every-other-day intermittent feeding) may induce similar physiologic effects even when average caloric intake is not different from ad libitum intake

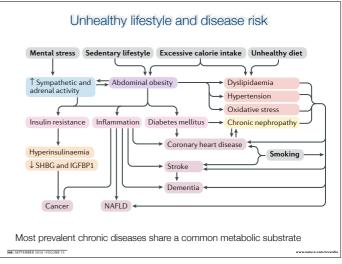
"CR mimetic drugs" e.g. metformin (hypoglycemic agent) and 2-deoxyglucose (glycolysis inhibitor)

CALERIE (Comprehensive Assessment of Long-term Effects of Restricted Intake of Energy Intake) trial: has tested effects of 2-3 years of CR (20-30% reduction) in young and middle-aged nonobese persons

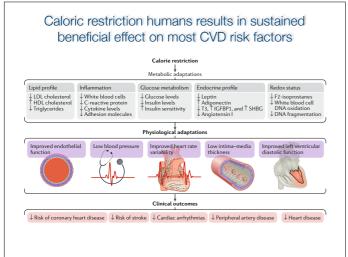
Cell Metabolism

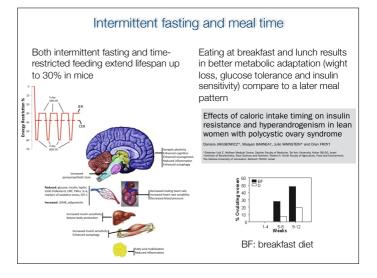


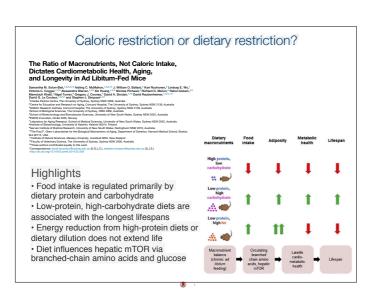


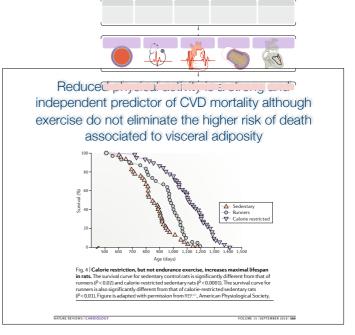


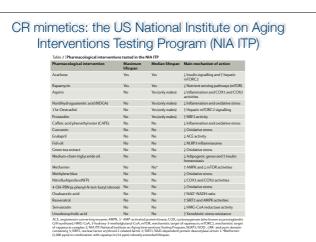












A large multi-institutional study investigating treatments with the potential to extend lifespan and delay disease/dysfunction in genetically heterogenous (outbread) mice

