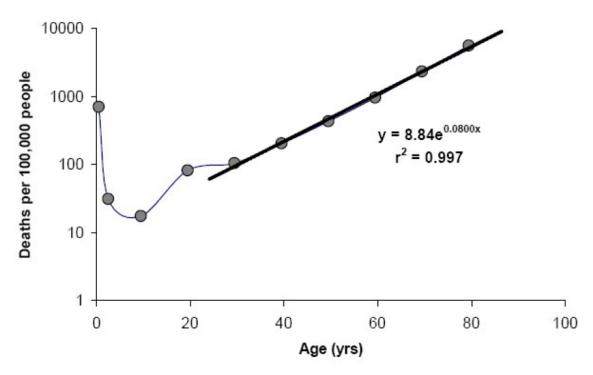
Approaching the underlying molecular mechanism of aging Non-canonical model system of aging

Journal Club on Lab Animal Science

Regina Reimann



Aging / Senescence: A risk factor for death



www.senescence.info

- 1. Exponential increase in mortality with age
- 2. Physiological changes, typically leading to a functional decline with age
- 3. Increased susceptibility to certain diseases

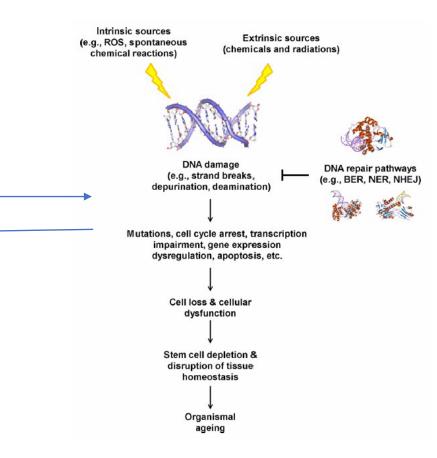
Why do we age? - Theories of aging.

1. Programmed Senescence

Birth Reproduction Senescence Maximal lifespan Humans 13.5 yrs >40 yrs Monkeys 4.5 yrs >20 yrs Mice 0.12 yrs >1 yr

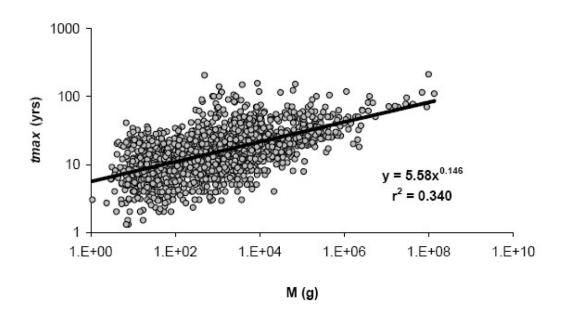
Aging occur on a fixed schedule triggered by the genetic program.

2. Damaged based theory of aging



Continuous process of damage accumulation originating in by-products metabolism.

Comparative biology of senescence

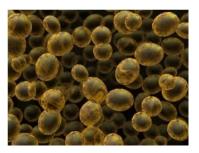


www.senescence.info

- Body size correlates with lifespan between species: larger animals live longer than smaller animals
- Brain mass correlates with tmax / in some case the size of other organs correlate
- Metabolic rate (questioned)
- Age at sexual maturation

Canonical models of senescence

Yeast



https://d3pddo38v7j30h.cloudfront.net /blog/wpcontent/uploads/2010/12/yea st-cells-web.jpg

C. elegans: 2-3 weeks



https://en.wikipedia.org/wiki/Caenorhabditis elegans

Drosophila: 2 months



https://en.wikipedia.org/wiki/Drosophila_m elanogaster

Mouse: 3 years



https://simple.wikipedia.org/wiki/Mouse

Limitation of canonical models:

- Yeast: no studies on multicellular and systemic aging.
- Nematode worms and flies: post-mitotic (no stem cell function and cancer) and lack an adaptive immune system.
- Mouse: commands considerable time and resources for lifespan studies.

Non-Canonical models of senescence

Short-lived: 4-5 month.

Turquoise killifish Nothobranchius furzeri

- · Shortest-lived vertebrate in captivity
- Wide set of aging phenotypes including cancer
- Sequenced genome
- Efficient transgenesis

Immortal



Planarian Schmidtea mediterranea

- · Potentially immortal lifespan
- · Pluripotent adult stem cell
- · Capable of whole body regeneration
- Can perform RNAi screens

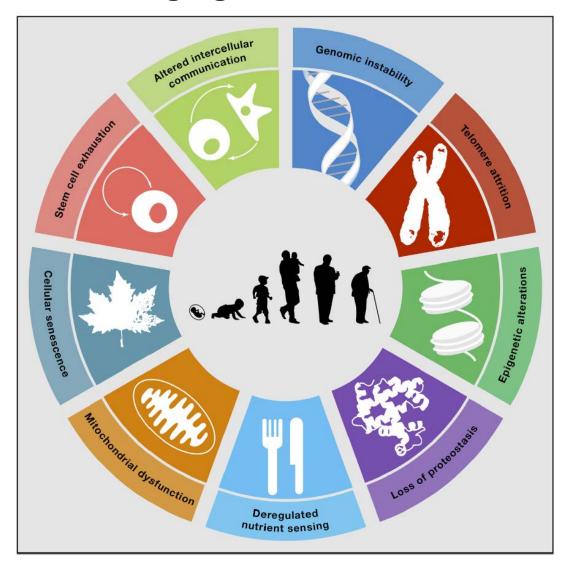
Long-lived: 31 years.



Naked mole-rat Heterocephalus glaber

- Exceptionally long-lived
- · Resistant to cancer
- Resistant to age-related diseases
- · Breeds in captivity, sequenced genome

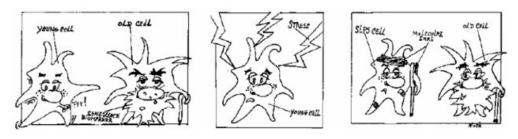
The hallmarks of aging



Cellular senescence

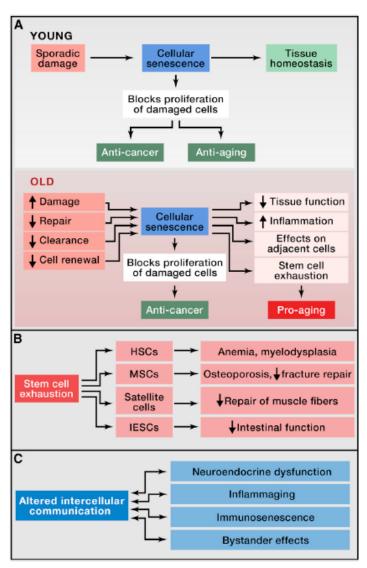
A stable arrest of the cell cycle coupled to stereotyped phenotypic changes

- Replicative senescence: telomer shortening
- Developmentally programmed senescence
- SIPS: stress-induced premature senescence
- OIS: Oncogen induced senescence



www.senescence.info

SIPS: stress-induced premature senescence



Naked mole rat: secret of a long and healthy live



Valenzano et al, 2017

- Higher protein stability
- Less increase in cysteine oxidation
- Higher level of cytoprotetive NRF2 signaling

Cancer inhibition

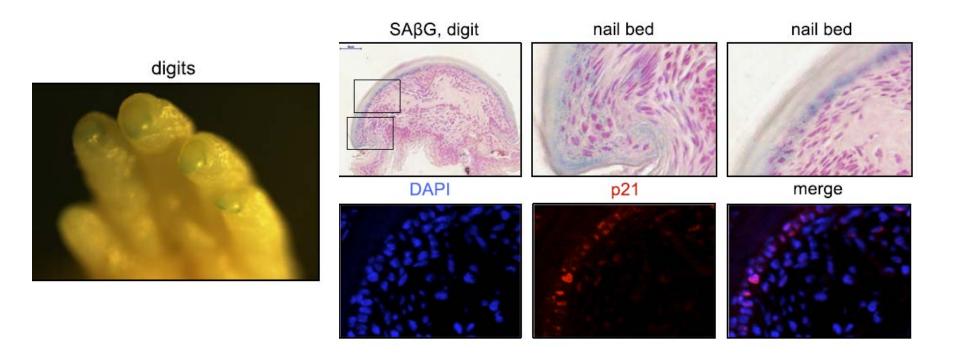
- Early contact inhibition
- p15/p16 hybrid
- High translation fidelity

Cellular senescence:

- No display of replicative senescence
- Expression of telomerase in somatic tissue

→ Other forms of cell senescence (SIPS, OIS, developmentally programmed senescence)?

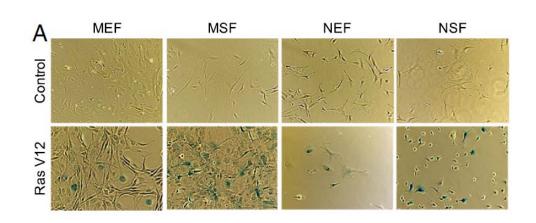
SIPS: Developmental programmed senescence in naked mole rats (NMR)



Senescence-associated β -galactosidase (SA- β -gal) staining.

β-galactosidase: Lysosomal hydrolase normally active at pH4, but it senescent cells it often happens o be activated at pH6.

SIPS: Oncogene-Induced Senescence



Transfection with:

• HRasV12: oncoprotein

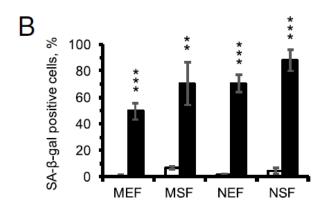
Control: GFP

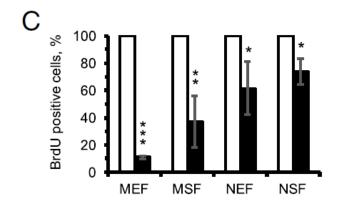
MEF: mouse embryonic fibroblast

MSF: mouse skin fibroblast

NEF: NMR embryonic fibroblast

NSF: NMR skin fibroblast



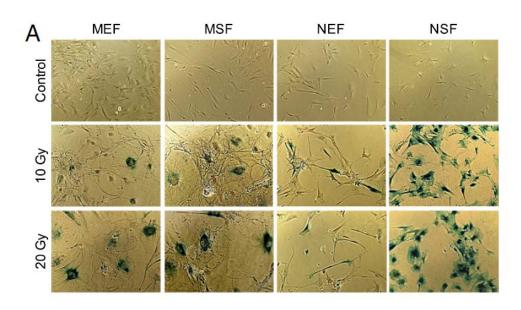


White: control Black: HRasV12

Oncogene-induced senescence, quantified by SA-β-gal.

Reduction in DNA synthesis, quantified by BrdU staining.

Attenuated senescence in response to γ-irradiation

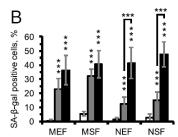


MEF: mouse embryonic fibroblast

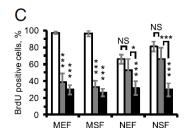
MSF: mouse skin fibroblast

NEF: NMR embryonic fibroblast

NSF: NMR skin fibroblast



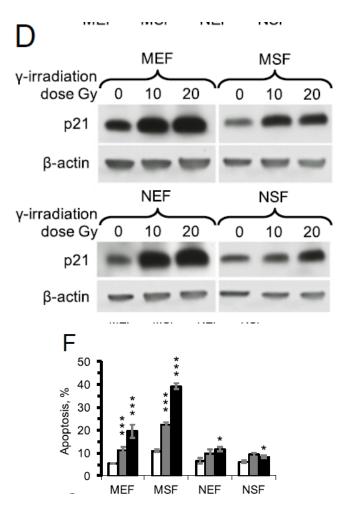
 γ -irradiation induced senescence, quantified by SA- β -gal.



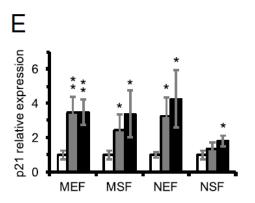
Reduction in DNA synthesis, quantified by BrdU staining.

White: control Grey: 10 Gy Black: 20 Gy

Attenuated senescence and apoptosis in response to γ-irradiation - II

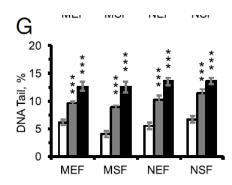


Annexin V FACS apoptosis assay



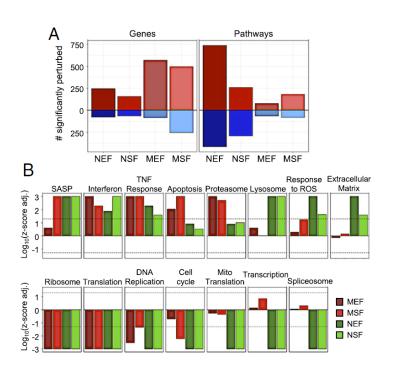
White: control Grey: 10 Gy Black: 20 Gy

P21: Cell cycle arrest in response to γ -irradiation.



DNA damage quantified with Comet assay

Gene expression change in the NMR upon γ-irradiation are less drastic, but more systematic.

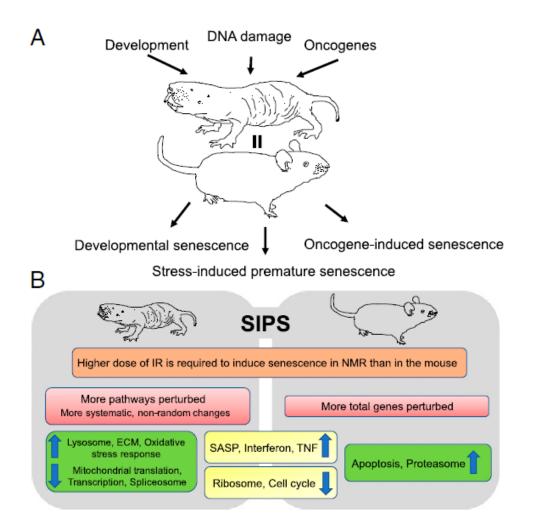


- Two times more differentially expressed genes mouse vs NMR fibroblasts
- More pathways enriched in NMR vs mouse fibroblasts
- Shared pathway alterations: immune response, cell cycle, DNA replication, translation, ribosome protein genes
- NMR unique down-regulation: transcription, spliceosome, mitochondrial translation
- NMR unique up-regulation: protein and glycoprotein metabolism, lipid metabolism, lysosomes, extracellular matrix and oxidative stress response.

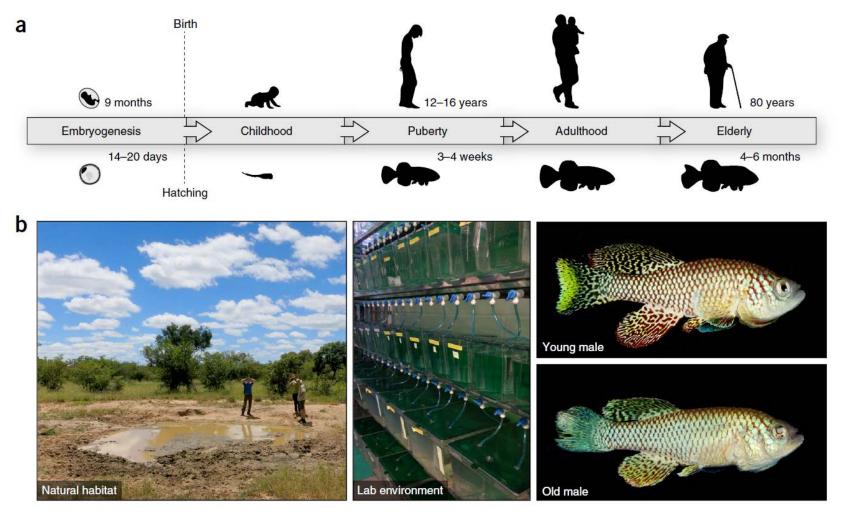
Functional enrichement of changes

- 20 Gy of γ-irradiation versus untreated
- RNA collection 12 d later (all irradiated cells displayed positive SA-β-gal staining)
- RNAseq, three biological replicates
- Uniform annotation of genes between species, 10959 gene coverage

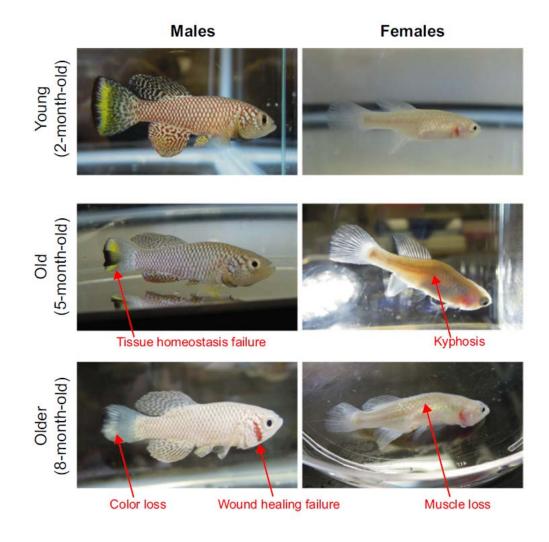
Conclusion from comparative analysis of senescence between mice and NMR



Turquioise killifish: live fast, die young

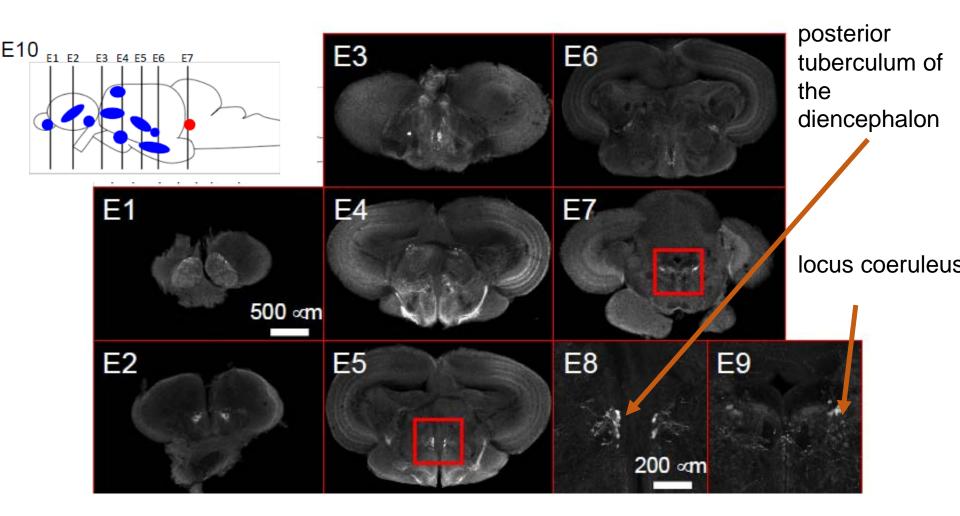


Aging phenotypes



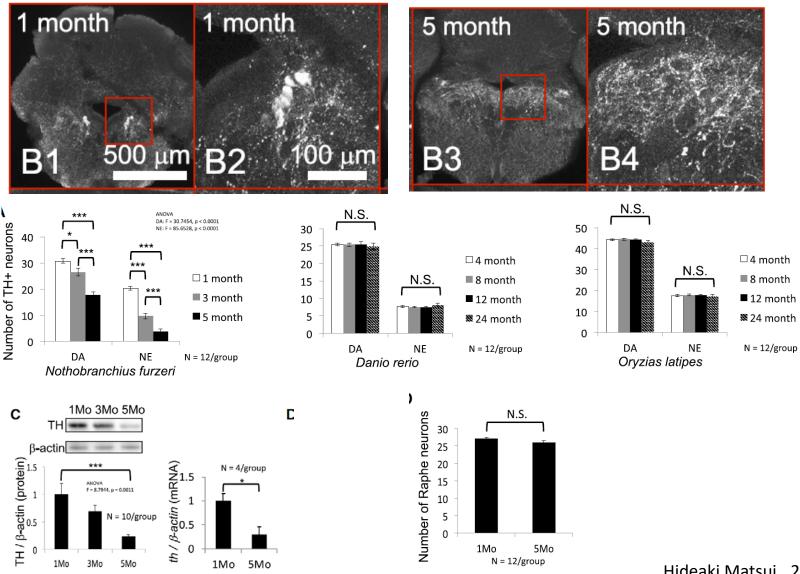
- Decline in reproduction
- Decline in fertility
- Decline in cognition
- Decline in mobility
- Decline in regeneration and tissue homeostasis
- Neural and muscular degeneration
- Cancerous lesions
- Multiple cause of death

Localisation of TH+ positive neurons and fibres

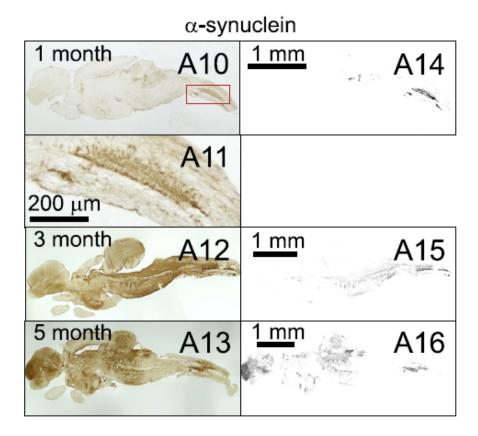


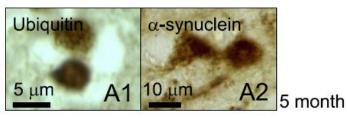
Coronal sections at different rostro–caudal levels show the localization of TH+ fibers and neurons in N. furzeri (3 month-old). / TH+: Tyrosine hydroxylase (to detect dopaminergic and noradrenergic neurons)

Degeneration of TH+ positive neurons

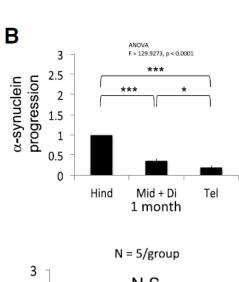


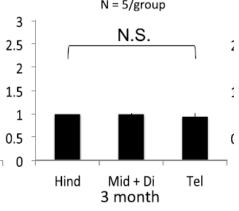
Progression of α-Synuclein pathology

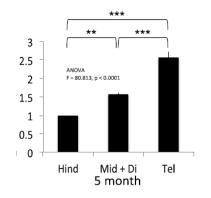




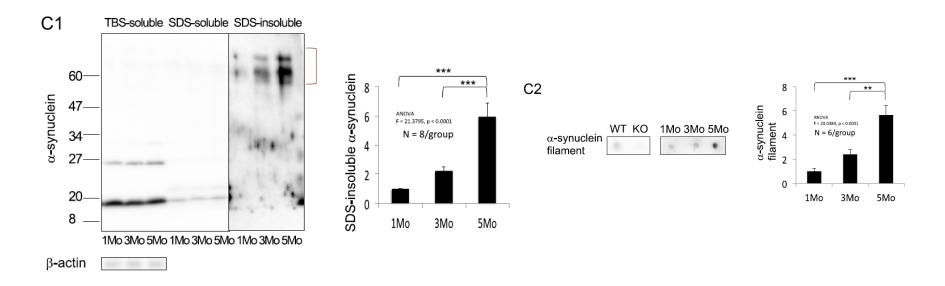
N. fuzeri α -synuclein antibody: peptide / rabbit







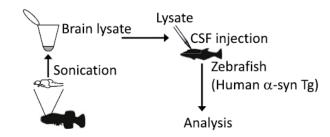
Progression of α -Synuclein pathology

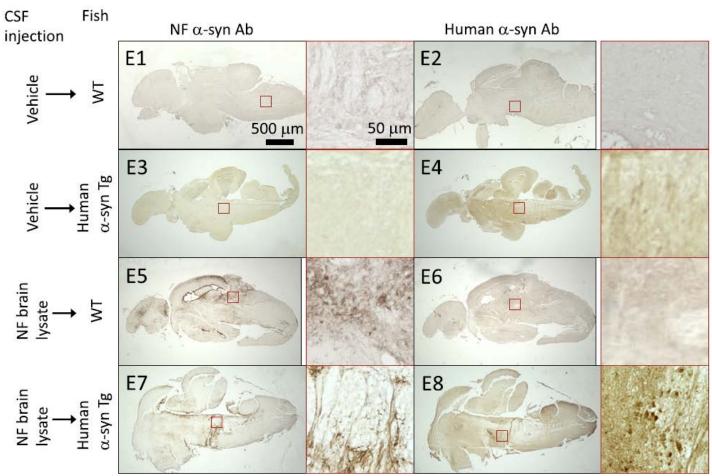


Immunoblot anaylsis of SDS-insoluble fraction.

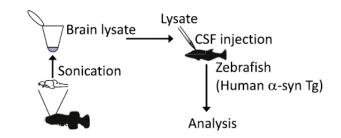
Dot-blot analysis of α -Synuclein fibrils in the brain.

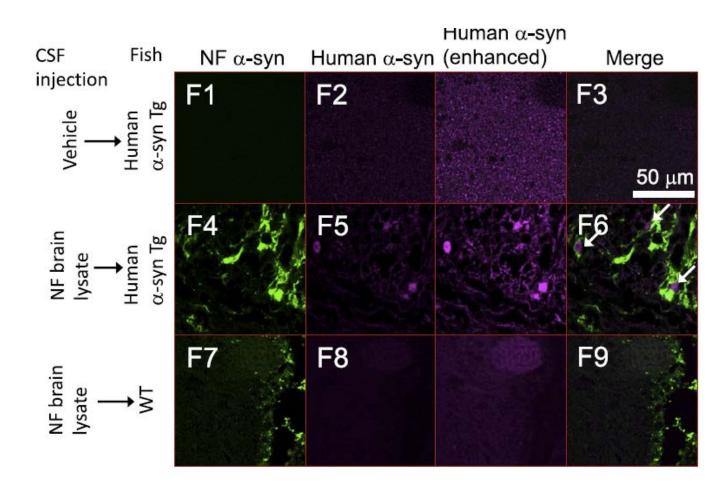
Passage of α-Synuclein seeds from Turquoise killifish to Zebrafish



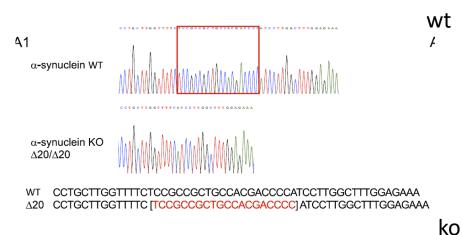


Passage of α -Synuclein seeds from Turquoise killifish to Zebrafish



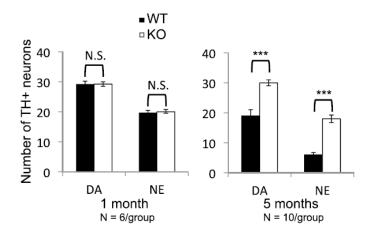


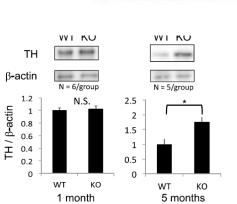
Rescue experiment-I

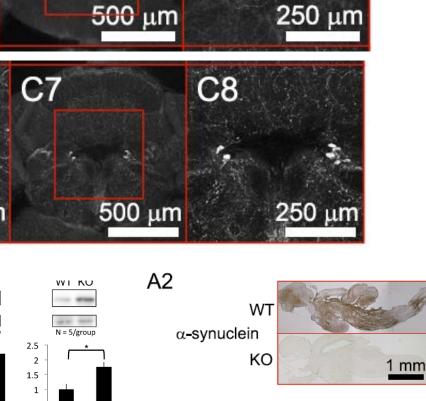


Generation of 20 bp deletion in the exone of α -Synuclein (CRISPR-Cas9).

Western blot / immunohistochemistry: no expression





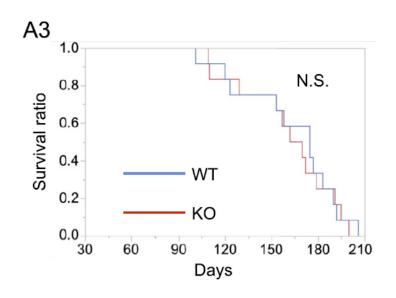


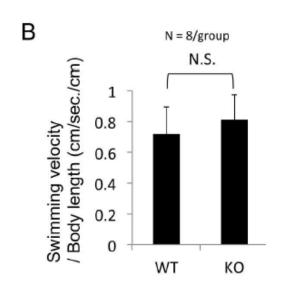
α-synuclein

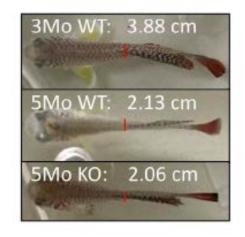
β-actin

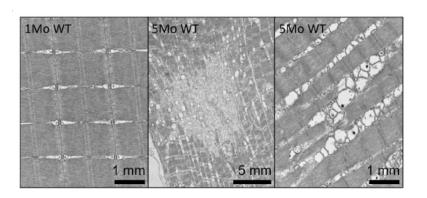
WT KO

Rescue experiment-II

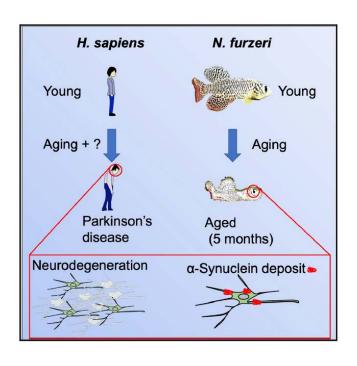






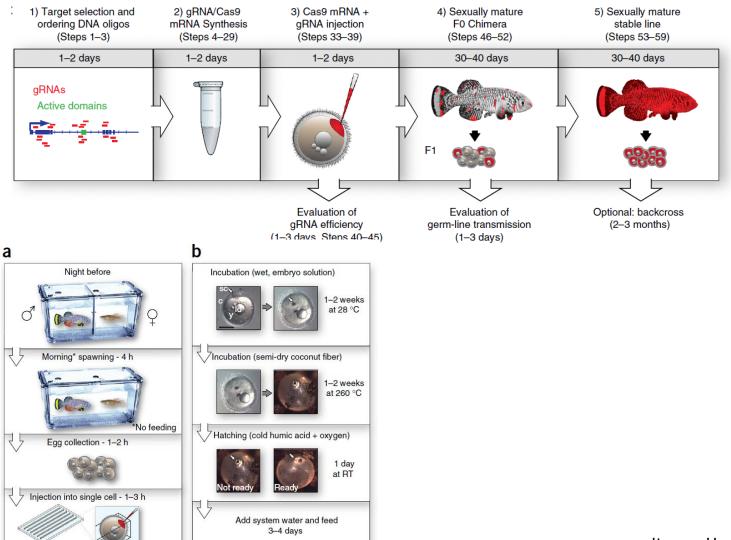


Summary: Dopaminergic degeneration in the Turquoise killifish

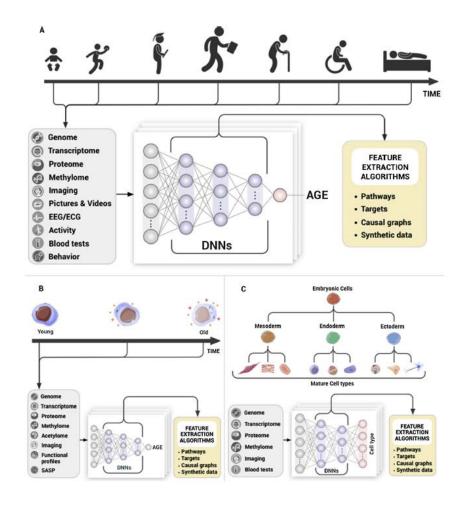


- Annual killifish reveals age-dependent degeneration of dopamine neurons
- Aged killifish shows accumulation of α -Synuclein in the brain
- Dopamine neurodegeneration is ameliorated by genetic depletion of α -Synuclein

Efficient genome engineering approaches for the Turquoise kilifish



Artificial intelligence for aging and longevity



Application of artificial intelligence of aging for biomarker development and target identification (at different levels).

Thank you for your attention!



¶

MONITORING-SHEET-FOR-PRION-INOCULATED-MICE-#

 α

 \Box

 α

CHECKED-BY:-¤ DATE:¤

Ħ	Ħ	Ħ	Ħ	Ħ	body·weight·(g)¤		Neurological·Scoring¤			comment
cage∙#¤	mouse∙#¤		inocul.·datex				Α¤	B¤	Total·(T)¤	
۵¤	۵¤	β	°μ	β	°¤	β	٩	β	Ħ	°¤
°¤	°¤	°¤	°¤	°¤	°¤	⁹ ti	٩	β	ী	°¤
°¤	٥Ħ	°¤	°¤	°¤	°¤	°¤	٩	β	Ħ	°¤
°¤	٥Ħ	°¤	°¤	°¤	°¤	⁹ ti	٩	β	'n	°¤
۵Ħ	٥Ħ	°¤	°¤	°¤	°¤	°¤	٩	٩	ীয	°¤
cage∙#¤	mouse∙#¤	mouse·line¤	<u>inocul</u> .∙date¤	researcher¤	reference-3	weight¤	Α¤	B¤	Total⋅(T)¤	å
β	۵¤	°μ	p°	β	β	β	æ	æ	ਖ	°¤
β	٥Ħ	°¤	°¤	°¤	β	⁹ t	ά	ά	শ্ব	°¤
۵¤	٥Ħ	°¤	°¤	°¤	β	°¤	β	β	শ্ব	°¤
۵¤	٥Å	°¤	°¤	°¤	β	β	ά	β	শ্ব	°¤
٥Ħ	٥Å	°¤	°¤	°¤	β	°pt	β	٩	শ্ব	°¤
cage∙#¤	mouse∙#¤	mouse·line¤	inocul.·datex	researcher¤	reference-3	weight¤	Α¤	B¤	Total⋅(T)¤	°¤
۵pt	۵pt	°¤	°¤	°μ	β	β	٩	ά	'n	°¤
۵Ħ	٥Ħ	°¤	°¤	°¤	β	β	٩	ά	Ή	°¤
۵pt	۵å	°¤	°¤	٩	β	⁹ t	٩	β	'n	°¤
۵Ä	٥Ħ	°μ	°¤	°μ	°¤	°¤	ď	ά	°μ	°¤
۵ä	۵å	٩	°¤	β	β	[®] #	β	β	'n	°¤
cage·#¤	mouse∙#¤	mouse·line¤	inoculdatex	researcher¤	reference-3	weight¤	Α¤	B¤	Total⋅(T)¤	°¤
٥Ħ	٥Ħ	°¤	°¤	°¤	°¤	°¤	٩	٩	ង	°¤
۵pt	۵pt	°¤	°¤	°¤	°¤	°¤	٩	٩	°μ	°¤
۵pt	۵å	°¤	°¤	β	βt	β	β	β	'n	°¤
٥Ħ	٥Ħ	°¤	°¤	°¤	β	β	٩	ά	Ή	°¤
۵pt	۵å	°¤	°¤	٩	β	⁹ t	٩	β	'n	°¤
cage·#¤	mouse·#¤	mouse·line¤	inocul. datex	researcher¤	reference-3	weight¤	Α¤	B¤	Total⋅(T)¤	°¤
۵pt	٥Ħ	°¤	°¤	°¤	β	°¤	٩	٩	ង	°¤
۵ħ	٥Ħ	°¤	°¤	°¤	°¤	°¤	٩	٩	°μ	°¤
۵pt	٥Ħ	°¤	°¤	°¤	β	°¤	٩	٩	°μ	°¤
۵pt	٥Ħ	°¤	°¤	°¤	β	°¤	٩¤	٩	ង	°¤
۵¤	۵pt	°¤	°¤	°¤	⁹ ¤	°¤	°¤	٩	°ta	°¤