

Werner's syndrome WRN protein A member of the RecQ family of DNA helicases. Other RecQ family helicase mutations produce genomic

mutations produce genomic instability diseases with progeriod symptoms: Bloom syndrome and Rothmund–Thomson syndrome.

Werner's syndrome: cellular features

- Normal human fibroblasts achieve approximately 60 population doublings in culture.
- Werner syndrome cells usually achieve only about 20 population doublings.

(lower Hayflick limit).

| | long-lived s | ential greater pecies | |
|---|---------------------------------|--|--|
| Organism + L. -mouse about 3 -human about 3 | <u>S:</u> B years 100 | Hayflick Limit: -doublings about 20 -doublings about 40-60 -doublings about 140 | |
| Species | Maximum life span (years) | Maximum doubling number | |
| Galapagos tortoise | 175 | 125 | |
| Man | 110 | 60 | |
| Horse | 46 | 82 | |
| Chicken | 30 | 35 | |
| Cat | 28 | | |
| Kangaroo | 16 | 92 46 | |
| Mink | 10 | 34 | |
| Mouse | 4 | 28 | |

| | roliferati from o | lder do | onors | | |
|--------|--------------------------------------|---------|--------------------------------------|-----------------|--|
| Fe | etal Lung | | Adult L | Adult Lung | |
| Strain | Number of population doublings | Strain | Number of population doublings | Age of donor | |
| WI-1 | 51 | WI-1000 | 29 | 87 | |
| WI-3 | 35 | WI-1001 | 18 | 80 | |
| WI-11 | 57 | WI-1002 | 21 | 69 | |
| WI-16 | 44 | WI-1003 | 24 | 67 | |
| WI-18 | 53 | WI-1004 | 22 | 61 | |
| WI-19 | 50 | WI-1005 | 16 | 58 | |
| WI-23 | 55 | WI-1006 | 14 | 58 | |

WI-1007

20

20 (14-29) 26

39

41

50

41

48

63

48

(35-63)

WI-24

WI-25

WI-26

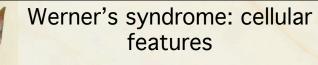
WI-27

WI-38

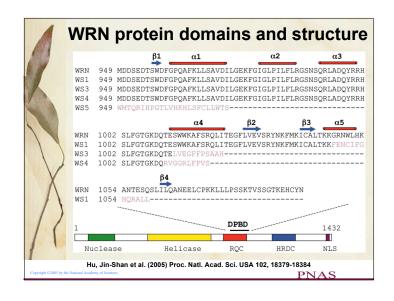
WI-44

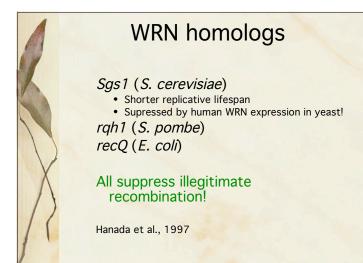
Average

range



- Sensitive to some but not all DNA damage agents:
 - Normal response: UV irradiation, ionizing radiation
 - Sensitive: carcinogen 4-nitroquinoline-1-oxide (4NQO) and to agents causing interstrand crosslinks.
- Increased rate of somatic mutations and chromosomal abnormalities such as translocations, inversions, and chromosome losses.



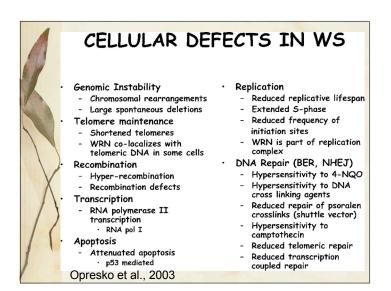


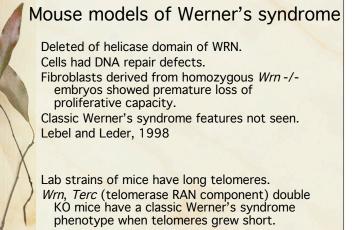
WRN protein

- Location: nucleolar
- WRN co-purifies with a 17S DNA replication complex.
- WRM binds proteins involved in DNA and RNA processing including DNA repair.

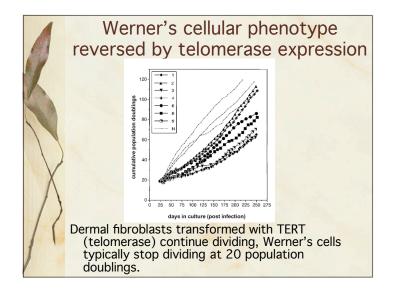
Loss of WRN:

- Transcriptional changes
- RNA pol II transcription is reduced by 40–60%.
- Deletion of telomeres from single sister chromatids.





Chang et al., 2004



Effects of WRN mutations

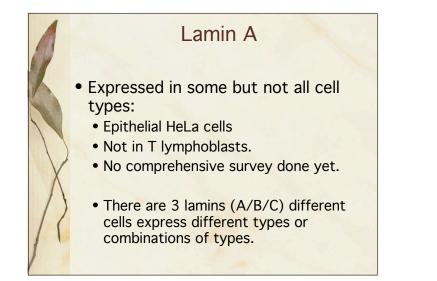
- Loss of proliferation of cell populations that replace themselves by cell division cause by cell death or early senescence.
- Renewing cell populations fail to replenish themselves and this gives rise to the Werner's syndrome phenotype.

Rothmund-Thomson syndrome

- RECQL4, a RecQ DNA helicase
- Is regulated by SIRT1
- SIRT1 deacetylates RECQL4, deactivating it and causing it to localize to the nucleolus.
- SIRT1 appears to be a regulator of several proteins involved in aging processes.

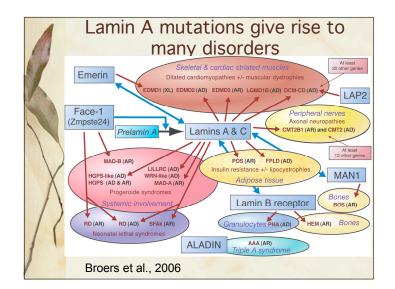
Hutchison-Gilford syndrome

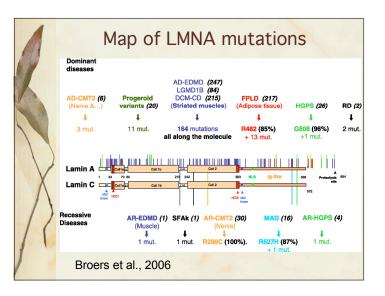
- Lamin A / LMNA protein.
- Lamins are structural protein components of the nuclear lamina, a protein network underlying the inner nuclear membrane that determines nuclear shape and size.
- The lamins constitute a class of intermediate filaments

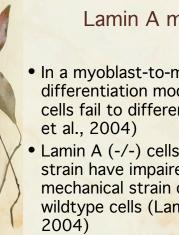


Lamin A mutations

- Truncations/missense mutations produce Hutchinson-Gilford syndrome.
- Other mutations in this gene give other disorders: muscular dystrophy, dilated cardiomyopathy, lipodystrophy





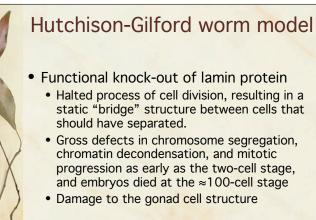


Lamin A mutations

- In a myoblast-to-myotube differentiation model, lamin A (-/-) cells fail to differentiate. (Favreau
- Lamin A (-/-) cells under mechanical strain have impaired viability under mechanical strain compared to wildtype cells (Lammerding et al.,

Lamin A mutation cellular phenotype

- Disrupted nuclear lamina, intranuclear architecture, and macromolecular interactions.
- Fibroblasts from individuals with HGPS have severe morphologic abnormalities in nuclear envelope structure.
- Heterochromatin-specific histone modifications
- Transcriptional changes.



Margalit et al., 2005

Hutchison-Gilford mouse model

- Introduced a mutation in Lamin A that causes autosomal dominant Emery-Dreifuss muscular dystrophy in humans.
- Normal at birth
- At 4 to 6 days developed severe growth retardation, dying within 4 to 5 weeks.
 - slight waddling gait, suggesting immobility of joints.
 - Loss of subcutaneous fat
 - Reduced numbers of eccrine and sebaceous glands
 - Increased collagen deposition in skin
 - Decreased hair follicle density
- Nuclear envelope abnormalities
- Decreased fibroblast Hayflick limit Mounkes et al., 2003

