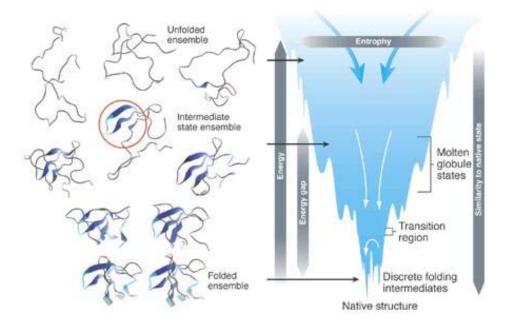
Aging and proteostasis



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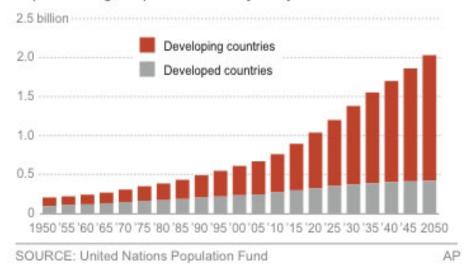


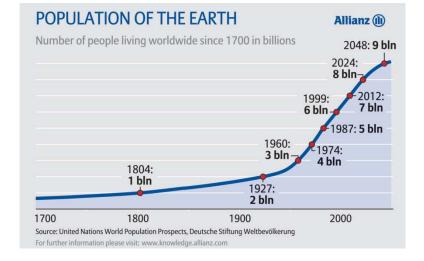
https://en.wikipedia.org/wiki/Jeanne_Calment

Aging population

Global rise in aging population

The number of people in the world aged 60 and older is expected to grow past 2 billion by the year 2050.





The Hallmarks of Aging



Cell 153, 1194-1217 (2013)

What is aging?

Aging is characterized by a progressive loss of physiological integrity, leading to impaired function and increased vulnerability to death.

Aging is a complex biological process that has prompted many new theories in regard to its process and origin. (programmed & non-programmed)

Many studies of aging have focused on molecular changes across the lifetime of an organism with the reasonable assumption that a series of progressive events collectively contribute to the aging process.

Proteostasis: protein homeostasis (through recycling)

A term coined in a landmark Science paper in 2008, is a broad concept that is relevant to all fields of study that involve protein synthesis, folding, processing, and turnover (Science 319:916–919, 2008).

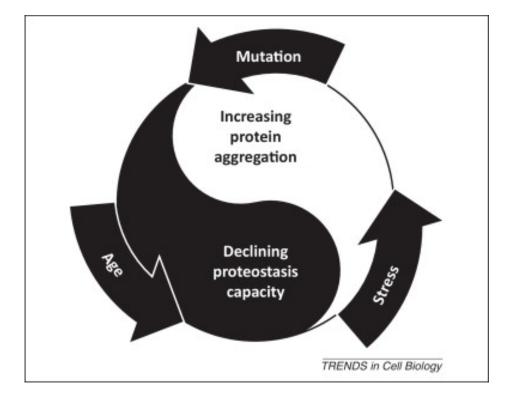
The decline of proteostasis with aging

When protein homeostasis degenerates and collapses, cells get old . . . and sick. Hallmark features of cellular aging include a buildup of proteotoxic stress, mistranslation of nascent polypeptides, and concentration of damaged, misfolded proteins and protein aggregates.

This accumulation of cellular waste is toxic to cells. That's when things start to fall apart. <u>http://irp.nih.gov/catalyst/v21i6/the-sig-beat</u>

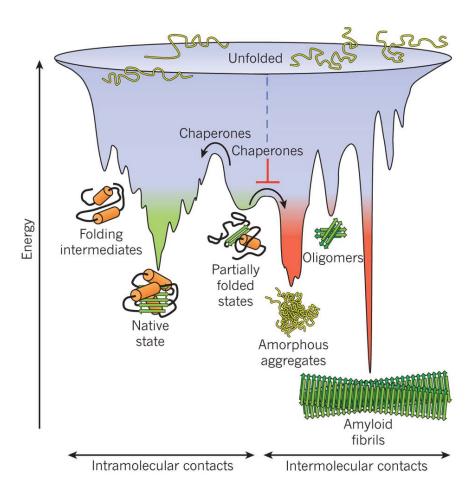


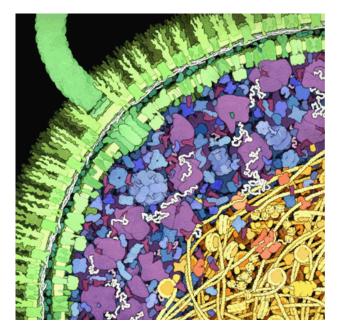
Self-propagating cycle of proteostasis decline in disease and aging



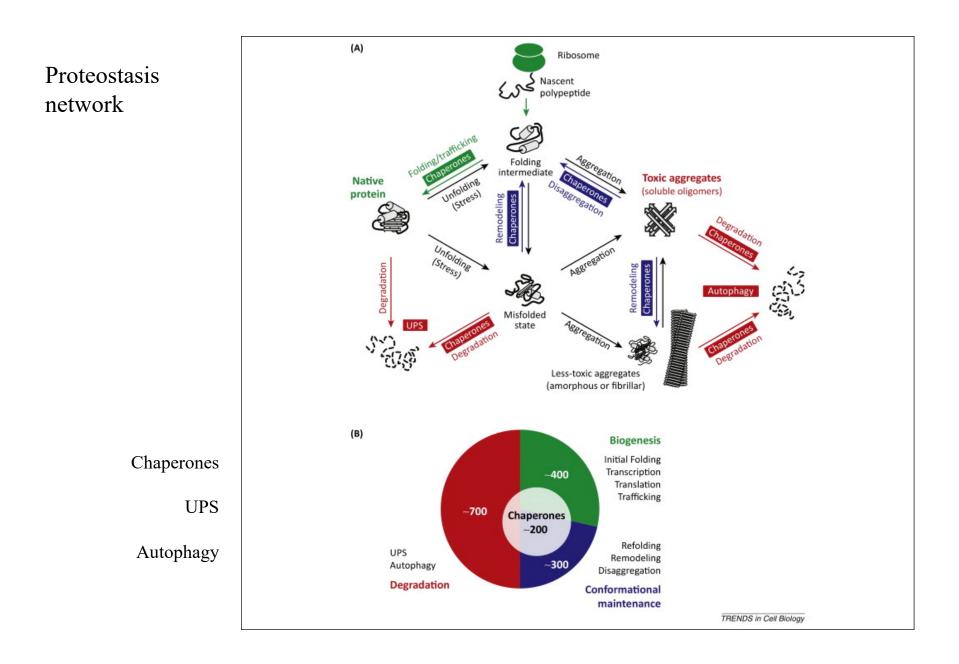
Chronic expression of aberrantly folded protein species caused by disease, aging, or external stress reduces proteostasis capacity by sequestering or otherwise inhibiting proteostasis network (PN) components. This results in misfolding and aggregation of endogenous proteins. These additional misfolded species in turn engage the PN (PN titration model), thereby further reducing the available proteostasis capacity and driving a positive feedback loop that eventually leads to proteostasis collapse. Trends in Cell Biology September 2014, Vol. 24, No. 9

Competing reactions of protein folding and aggregation





http://www.nature.com/nature/journal/v475/n7356/full/nature10317.html



How does impaired proteostasis in one cell or tissue affect the organism?

Recent studies show that unlike thought before, stress responses and proteostasis are controlled at the organismal level by inter-tissue communication.

Can impaired proteostasis be a therapeutic target?

Pathways and modifiers influencing proteostasis. Genetic, epigenetic, physiological and environmental stressors affect proteostasis and cause the accumulation of misfunctional proteins. Small molecule modulators of the activities of the proteostasis network pathways (small molecule proteostasis regulators) facilitate chaperone-mediate refolding and/or induce the degradation of misfolded and damaged proteins therefore rebalancing cellular proteostasis. In parenthesis are indicated some of the genes responsible for proteostasis maintenance. Curr Top Med Chem. 2012; 12(22): 2623–2640.

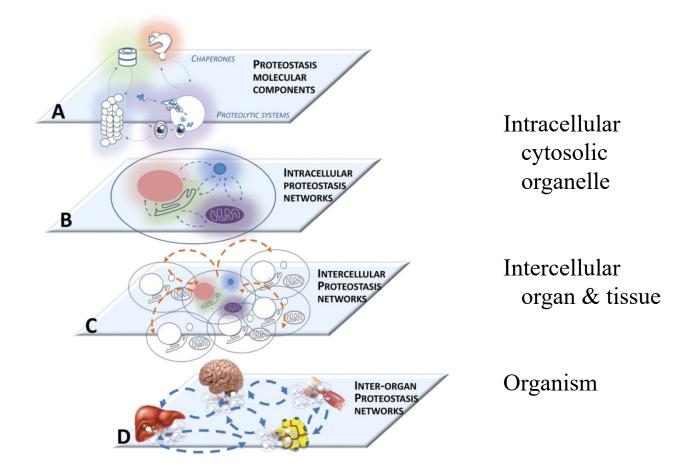
Proteostasis modifiers

Error-prone synthesis Mutations Polymorphisms Post-translational modifications Environmental stress Physiological stress Metabolic stress

PROTEOSTASIS

HSR (HSPA1A, DNAJ) UPR (HSPA5, HSP90B1, CALR) Oxidative stress response (GCLM, HO1, SOD1) Autophagy (ATG) UPS, ERAD (EDEM) Organellar Ca²⁺, HDAC

Small molecule proteostasis regulators Schematic of the different levels of integration of the proteostasis networks



J Gerontol A Biol Sci Med Sci 2014 June;69(S1):S33-S38

Organismal proteostasis: role of cell-nonautonomous regulation and transcellular chaperone signaling

The HSR and other cell stress responses such as the unfolded protein response (UPR) can function autonomously in single-cell eukaryotes and tissue culture cells; however, within the context of a multicellular animal, the PN is regulated by cell-nonautonomous signaling through specific sensory neurons and by the process of transcellular chaperone signaling.

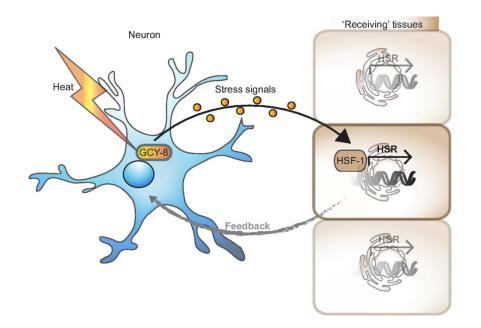
These newly identified forms of stress signaling control the PN between neurons and nonneuronal somatic tissues to achieve balanced tissue expression of chaperones in response to environmental stress and to ensure that metastable aggregation-prone proteins expressed within any single tissue do not generate local proteotoxic risk. Transcellular chaperone signaling leads to the compensatory expression of chaperones in other somatic tissues of the animal, perhaps preventing the spread of proteotoxic damage. Genes Dev. 2014 Jul 15; 28(14): 1533–1543.

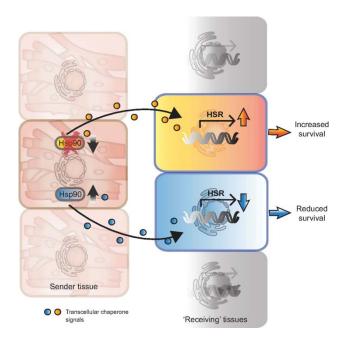
Signaling category	Cell-nonautonomous response	Sender tissue	Signal	References
Neuron to nonneuronal soma	HSR	AFD neuron	AFD-dependent neuropeptide?	Prahlad et al. 2008
	UPR	ASH/ASI neurons	Unknown	Sun et al. 2011
	Noncanonical UPR (innate immunity)	ASH/ASI neurons	Unknown	Sun et al. 2012
	UPR ^{mito}	Neurons	Neuronal "mitokine"	Durieux et al. 2011
	UPRER	Neurons	Neurotransmitter?	Taylor and Dillin 2013
Direct signaling between nonneuronal tissues	Gonadal longevity pathway	Steroidogenic tissues	Dafachronic acid	Antebi et al. 1998; Motola et al. 2006; Yamawaki et al. 2010
	HSR and systemic hsp epxression	Intestine, muscle, neurons	Transcellular stress factor (TSF)	van Oosten-Hawle et al. 2013
	FOXO-to-FOXO ⁻ signaling	Intestine, muscle, hypodermis	mdt-15-dependent lipid signals	Zhang et al. 2013
	Innate immune responses	Gonad, intestine	Secreted immune peptides	Ermolaeva et al. 2013
	Longevity	Gonad to soma	lipl-4-dependent fatty acids	Lapierre et al. 2011
	Longevity	Gonad to soma	Oleic acid	Goudeau et al. (2011)
	Longevity	Gonad to soma	let-7 microRNA family	Shen et al. 2012
	FOXO-to-FOXO	Intestine	IGF-like signals	Murphy et al. 2003
Indirect signaling between nonneuronal tissues (neuronal feedback)	Thermotaxis	Muscle, intestine	Estrogen	Sugi et al. 2011
	Longevity and survival	Gonad, neurons, intestine	mir-71	Boulias and Horvitz 2012
	FOXO/4EBP signaling (D. melanogaster)	Muscle	IGF?	Demontis and Perrimon 2010
	Mitophagy/ILS (D. melanogaster)	Muscle	Secreted IGFBP (impL2)	Owusu-Ansah et al. 2013

 Table 1.
 Cell-nonautonomous signaling responses

Categorization for cell-nonautomonous signaling responses in C. elegans (unless otherwise noted).

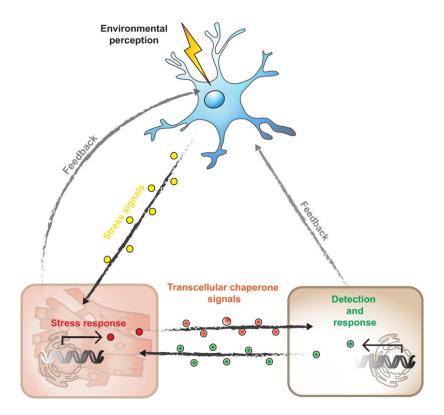
Cell non-autonomous control of the heat shock response (HSR) by neurons. Organismal control of heat shock transcription factor 1 (HSF-1) transcriptional activity in peripheral tissues by the thermosensory AFD neuron via a guanylyl cyclase GCY-8-dependent signaling cascade. A steroid signallingdependent feedback loop from peripheral cells reports HSR activity back to the AFD neurons.





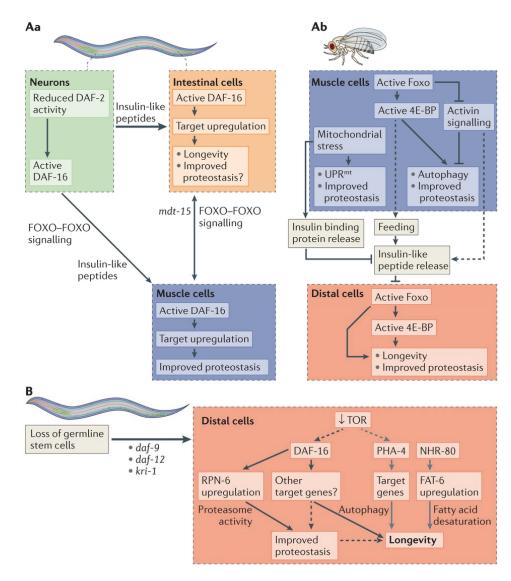
Transcellular chaperone signaling regulates the organismal HSR via

tissue-to-tissue crosstalk. An imbalance of proteostasis in a single tissue, through reduced (orange) or elevated (blue) expression of Hsp90, is detected and responded to in a different tissue via transcellular chaperone signaling. Because of the key role of Hsp90 as a negative regulator of the HSF-1-dependent HSR, the HSR is either induced (orange) or repressed (blue) at a cell non-autonomous level, leading to different outcomes for organismal survival during stress conditions. Cell non-autonomous control of proteostasis via the nervous system and transcellular chaperone signaling.

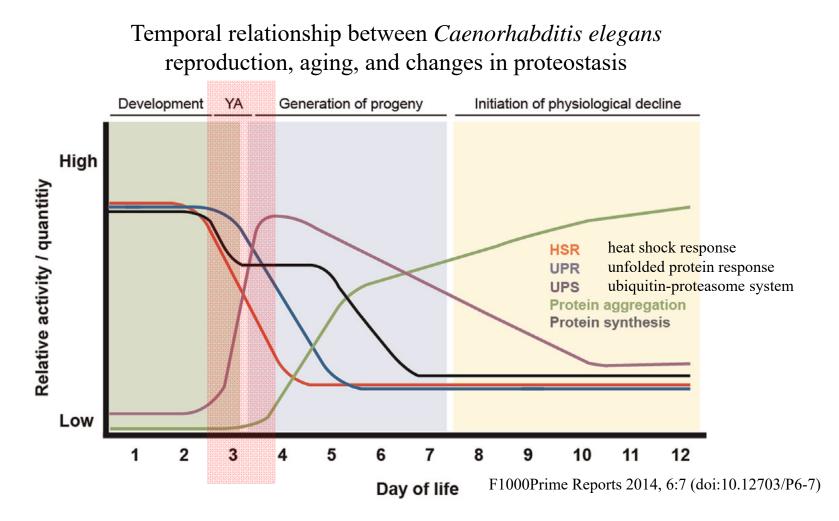


Sensory neurons perceive environmental stimuli and integrate the environmental challenge to fine-tune proteostasis in peripheral tissues. Non-neuronal tissues report altered proteostasis conditions back to the nervous system. At the same time, tissue-to-tissue signals via transcellular chaperone signaling can override the neuronal component to control cell-specific proteostasis. This allows proteostasis crosstalk between somatic cells independent of neural control.

Cell non-autonomous regulation of proteostasis by insulin-like signalling and gonadal signalling



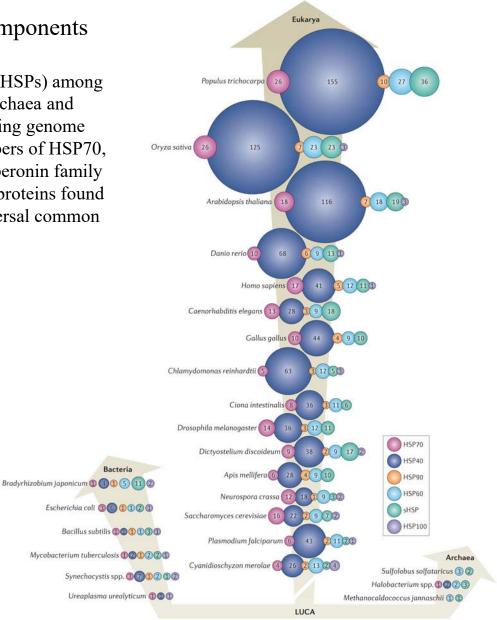
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What is the molecular mechanism(s) responsible for PN remodeling and the upstream signals that promote these changes?

Diversity in proteostasis network components

Shown is the evolution of heat shock proteins (HSPs) among the extant three domains of life — Bacteria, Archaea and Eukarya. Vertical arrows correspond to increasing genome size and number of expressed genes. The numbers of HSP70, HSP40 and HSP90 chaperones, of HSP60 chaperonin family subunits, small HSPs (sHSPs) and of HSP100 proteins found in each species are indicated. LUCA, last universal common ancestor.



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Pharmacological chaperones: a class of molecules that bind and inhibit the unstable or misfolded proteins.

