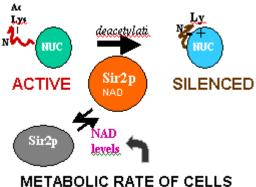
SIR2 PROTEINS REGULATE AGING IN RESPONSE TO NUTRIENTS

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Studies on the molecular basis of aging in my lab over the past nine years have led to the notion that the SIR2 gene controls the pace of aging in response to nutrient availability. SIR2 promotes longevity: sir2 deletion strains of yeast have short life spans, and strains with an extra copy of SIR2 live longer than wild type. Sir2p triggers silencing at select genomic locations, including mating type genes, telomeres, and the ribosomal DNA (rDNA). Silencing at this last locus represses the formation of toxic rDNA circles by recombination and thus potentiates a long life span.

Sir2p was shown to be an NAD-dependent histone deacetylase (see figure below). This finding explains how Sir2p creates silenced chromatin and, surprisingly, draws a link between metabolic rate and aging. We suggest that as metabolic rate slows, for example during caloric restriction, NAD is freed up from its many metabolic reactions and activates Sir2p. This promotes an increase in silencing and a longer life span. We show experimentally that SIR2 and NAD are involved in the extention of yeast life span by caloric restriction. We imagine that this mechanism has been selected because it allows organisms to stay alive longer in the face of adversity, to resume reproduction when conditions improve. It is interesting to note, in this regard, that SIR2 also controls mating and thus sporulation, a pathway of cell differentiation that allows long-term survival in the face of scarcity.



METABOLIC RATE OF CELLS

More recently we have begun experiments to determine whether SIR2 gene homologs also regulate aging in animals. Two systems have been used, C. elegans and mice. Studies in C. elegans that relate sir-2 genes to aging in that organism will be described. Models will be put forward to suggest how Sir2 proteins could be general regulators of aging.

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