

SUPPLEMENTARY TABLE

Supplementary Table 1. Role of studied genes in determining the stress response.

Target gene	Characteristics	Effects
Catalase (<i>Cat</i>)	Enzymes catalyzes the decomposition of hydrogen peroxide in cells [1].	Is involved in response to photooxidative [2], endoplasmic reticulum [3], oxidative stresses [4] and to hydrogen peroxide stimulus [5]. Overexpression of human catalase, which targets mitochondria, increases life span in <i>Mus musculus</i> by about 20% [6].
frataxin (<i>fh</i>)	Protein essential for iron-sulfur clusters synthesis, which are necessary for the production of ATP by the respiratory chain, as well as in other biological processes such as steroidogenesis [7–9].	Is involved in response to oxidative stress [7], to hydrogen peroxide stimulus [10], to iron ion homeostasis [9] and to the hypoxia-induced response [11]. Overexpression of frataxin in mitochondria increases the antioxidant capacity, resistance to oxidative stress, and life span in <i>Drosophila melanogaster</i> females [12].
Growth arrest and DNA damage-inducible 45 (<i>Gadd45</i>)	Participates in the regulation of the cell cycle and oviposition; required to activate MAPKKK, JNK activity [8, 13].	Is involved in response to oxidative, thermal and genotoxic stresses [14, 15]. Overexpression of the <i>D-Gadd45</i> gene in the nervous system leads to a significant increase in the life span in <i>Drosophila melanogaster</i> [16].
Heat shock protein 68 (<i>Hsp68</i>)	Protein necessary for response to temperature or stressing cell stimuli [17].	Is involved to starvation stimulus [18]. Overexpression in somatic cells led to an increase in the average life span of <i>Drosophila melanogaster</i> by 20% [18, 19].
Heat shock protein 83 (<i>Hsp83</i>)	Protein necessary for response to temperature or stressing cell stimuli [17] orthologous to human HSP90 gene (https://www.ncbi.nlm.nih.gov/gene/38389).	Is involved in response to heat, oxidative stresses and ionizing radiation [20–22], also in regulation of circadian sleep/wake cycle [23].
Ku80 (<i>Ku80</i>)	Ku70/Ku80 heterodimer, or Ku, is the central component of the nonhomologous end joining (NHEJ) pathway of double strand break repair [24]. Orthologous to human XRCC5 gene (https://www.ncbi.nlm.nih.gov/gene/7520).	Is involved in gamma radiation and X-ray responses [8, 25]. Deletion of Ku80 leads to signs of premature aging such as osteopenia, atrophic skin, hepatocellular degeneration, and age-related mortality in <i>Mus musculus</i> [26]. Also, <i>Mus musculus</i> showed a decrease in life span by 40% [27].
Peroxiredoxin V (<i>PrxV</i>)	Encodes an atypical member of the thiol-specific peroxidase family, which form intramolecular disulfide bonds during the catalytic cycle [28].	Is involved in response to oxidative stress, cell redox homeostasis and hydrogen peroxide catabolic process [8, 29]. Overexpression of PrxV caused an increase in the average and median life span in <i>Drosophila melanogaster</i> under normal conditions. Against the dPrxV mutants (- / -) were more susceptible to oxidative stress, had a higher incidence of apoptosis, and a shorter average life span [30].

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