

Astrocytes, the new trending topic in aging and neurodegeneration

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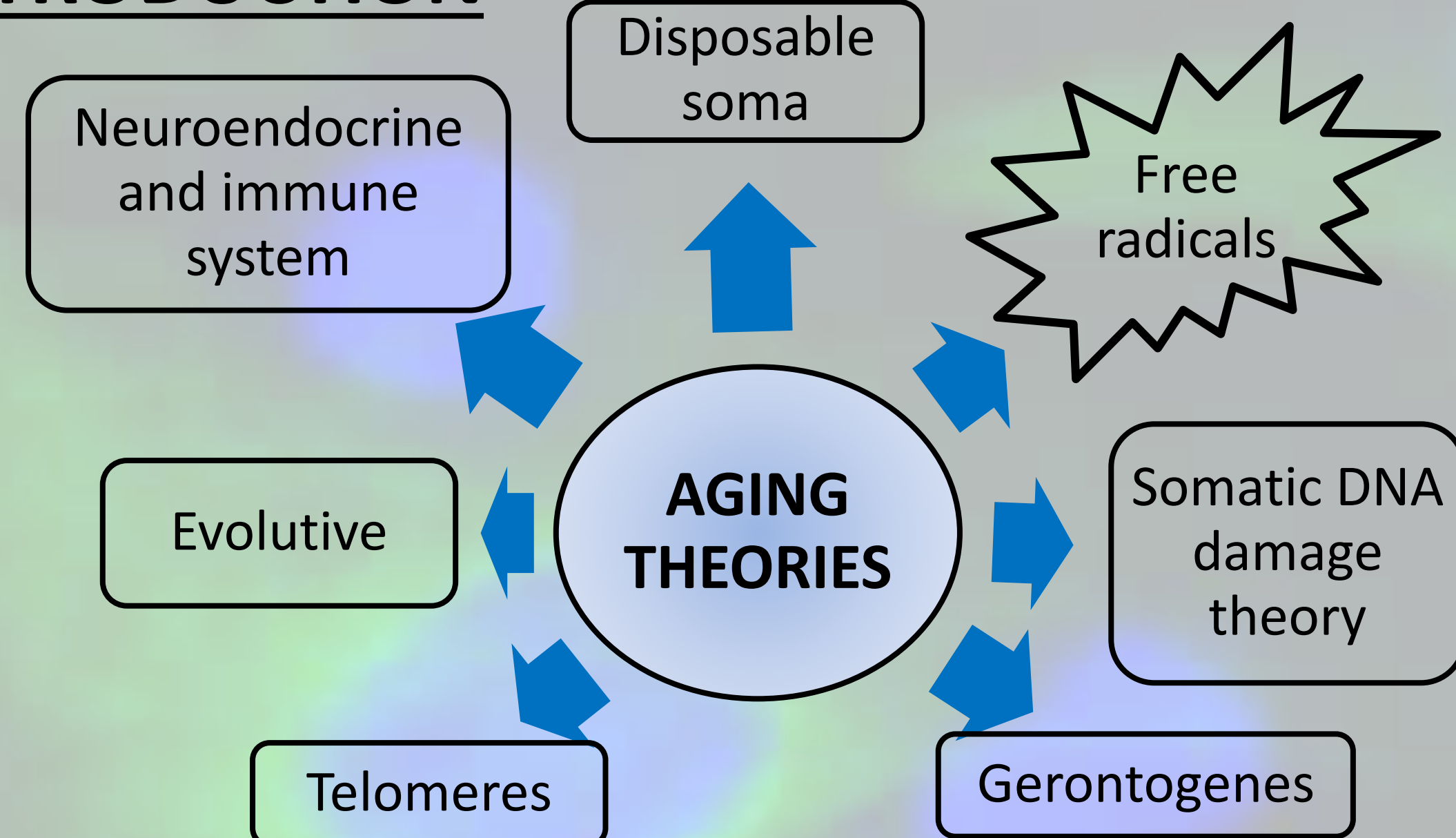
OBJECTIVES

Aging is the result of different morphological, psychological, functional and biochemical changes.

This review intends to sum up the different aging theories and the molecular changes in an aging brain. Also the aim of the project is to enhance astrocyte functions and their relation to neurodegenerative processes.

Additionally, some interesting results of the neuroprotective capacity and the changes in aging animal models are shown.

INTRODUCTION

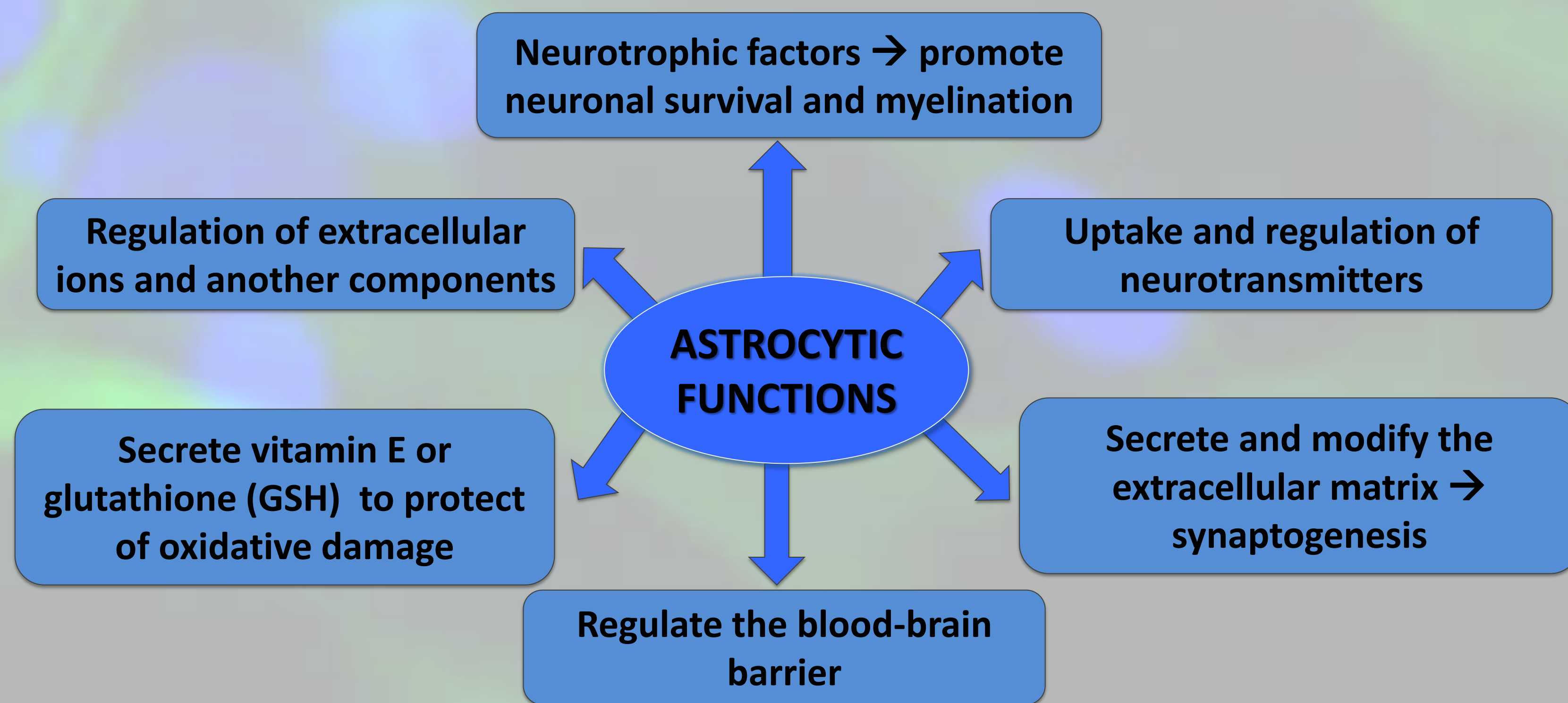


MOLECULAR CHANGES IN AGING

- Decrease levels of growth factors: nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), neurotrophin 3 (NT-3) and glial cell-derived neurotrophic factor (GDNF).
- Low energetic metabolism due to mitochondrial damage.
- Calcium homeostasis:
 - Stimulation of long-term potentiation (LTP).
 - Altered proteins → increase of apoptotic rate.
- Excitotoxicity: increase of levels of neurotransmitters in the synaptic cleft due to the decreasing levels of EAAT (excitatory amino acid transporter).

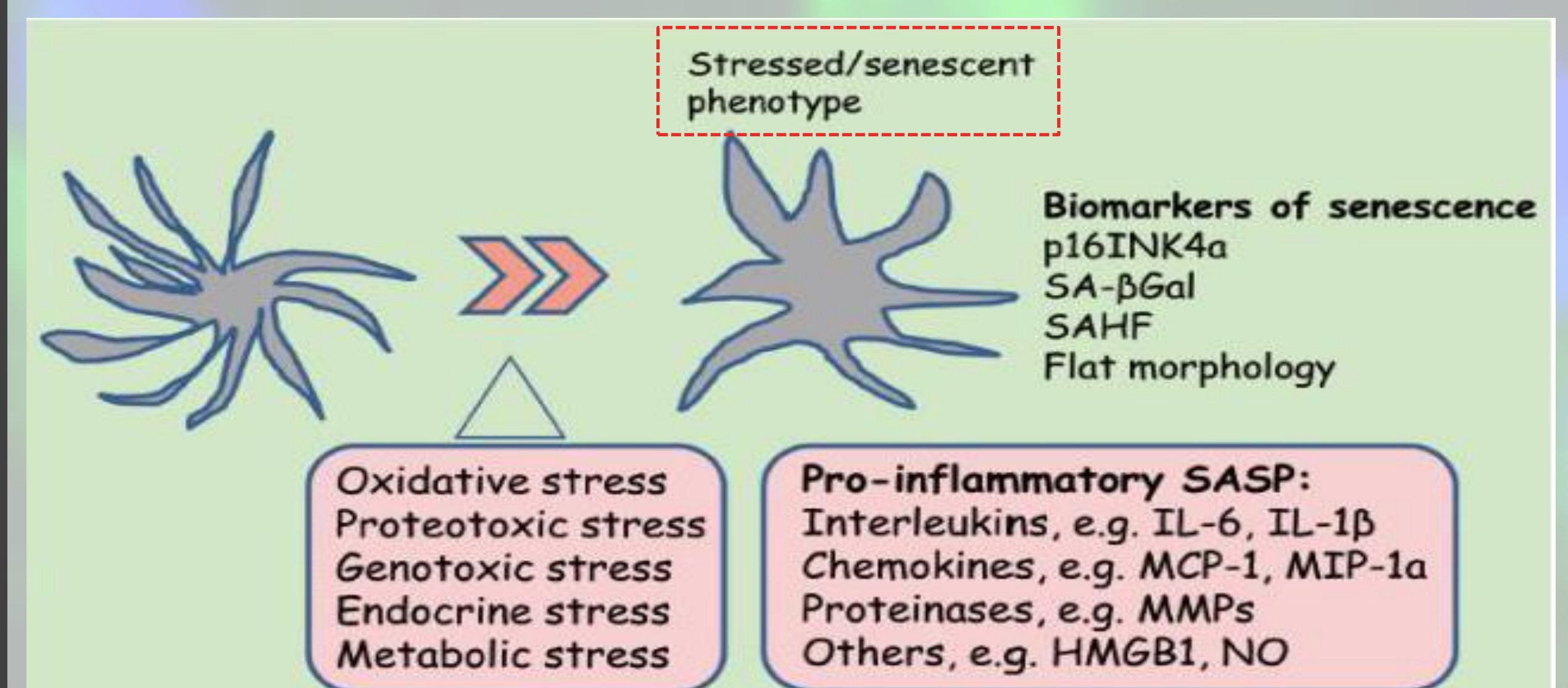
ASTROCYTES FUNCTIONS

Astrocytes are the most numerous cell type in the brain and they have a significant role in brain injury, both in the prevention of damage and later in ensuring its repair.



SENESCENCE-ASSOCIATED SECRETORY

PHENOTYPE (SASP)₁



❖ SENESCENCE-ACCELERATED MICE PRONE (SAMP8)

Murine model that manifests irreversible advancing senescence with pathological, biochemical and behavioural alterations related to aging.

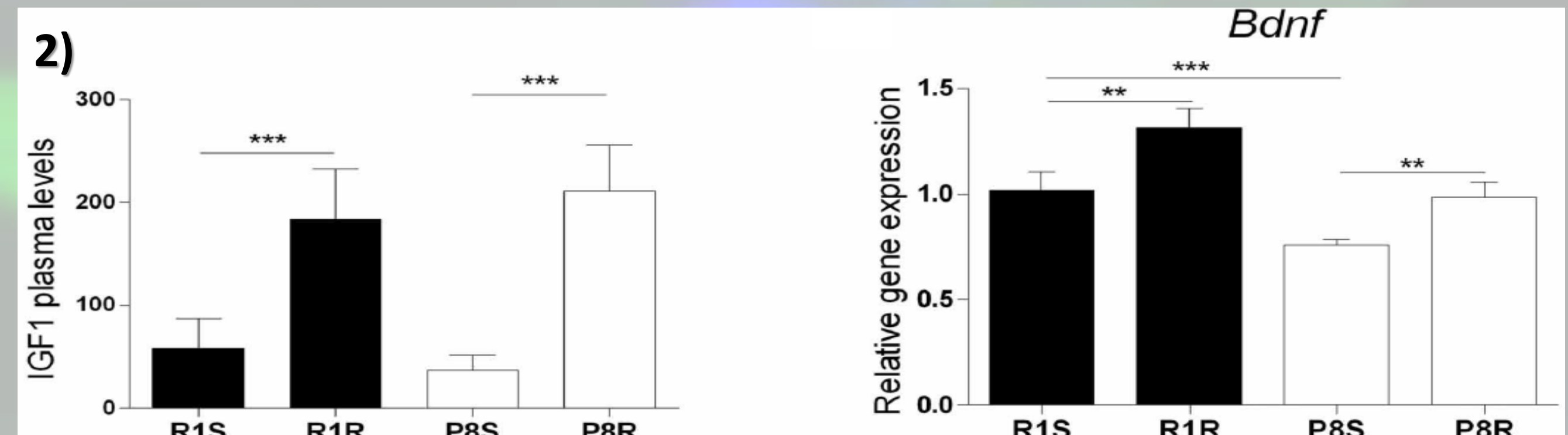
RESULTS OF THE STUDIES

1. Proteomic study of neuron and astrocyte cultures from senescence-accelerated mouse SAMP8 reveals degenerative changes.
2. Epigenetic alterations in hippocampus of SAMP8 senescent mice and modulation by voluntary physical exercise.
3. Neurons from senescence-accelerated SAMP8 mice are protected against frailty by the sirtuin 1 promoting agents melatonin and resveratrol.

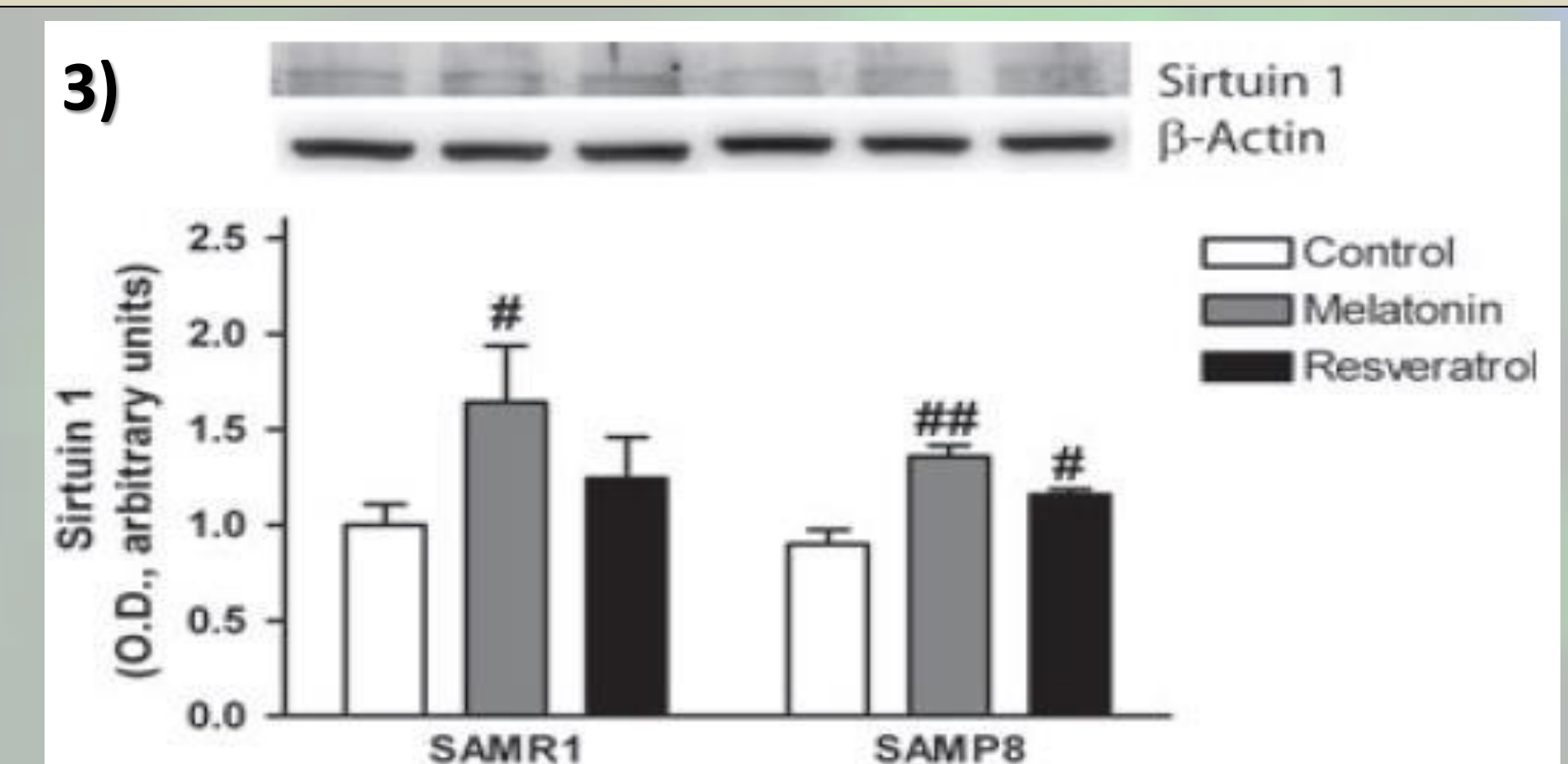
| 1) | Protein type | Protein identified | Gene name | Change SAMP8 versus SAMR1 | SAMR1 mean ± SD | SAMP8 mean ± SD | Ratio SAMR1/SAMP8 | p | Spot |
|----------------------------|--|--------------------|--------------|---------------------------|-----------------|-----------------|-------------------|----|------------------|
| Energy metabolism | Adenylate kinase isoenzyme 4 | Ak3l1 | Up | ND | 0.069 ± 0.029 | NA | NA | NA | 671 |
| | Acyl-coenzyme A thioesterase 2 | Acot2 | Up | ND | 0.149 ± 0.020 | NA | NA | NA | 273 |
| | Cytochrome c oxidase subunit 4 isoform 1 | Cox4i1 | Down | 0.054 ± 0.018 | 0.018 ± 0.010 | 3.0 | 0.0353 | | 970 |
| Biosynthesis | Dihydropteridine reductase | Qdpr | Down | 0.741 ± 0.187 | 0.340 ± 0.110 | 2.2 | 0.0330 | | 634 |
| | S-methyl-5'-thioadenosine phosphorylase | Mtap | Increased pl | 0.119 ± 0.033 | 0.092 ± 0.025 | NA | NA | | 604 ^a |
| Transduction and signaling | Protein-arginine deiminase type-2 | Padl2 | Up | ND | 0.090 ± 0.003 | NA | NA | NA | 412 |
| | Poly(rC)-binding protein 1 | Pcbp1 | Up | 0.108 ± 0.050 | 0.234 ± 0.047 | 0.5 | 0.0011 | | 336 |
| Stress response | Sodium/hydrogen exchanger 5 | Slc9a5 | Up | ND | 0.075 ± 0.021 | NA | NA | NA | 364 |
| | Aldehyde dehydrogenase | Aldh2 | Up | 0.056 ± 0.016 | 0.106 ± 0.033 | 0.5 | 0.0333 | | 597 |
| Cytoskeletal | Stomatin-like protein 2 | Stoml2 | Up | 0.075 ± 0.021 | 0.159 ± 0.065 | 0.5 | 0.0134 | | 361 |
| | Myosin light polypeptide 6 | Myl6 | Up | 0.833 ± 0.305 | 1.527 ± 0.551 | 0.5 | 0.0224 | | 903 |
| | Macrophage-capping protein | Capg | Down | 0.138 ± 0.017 | 0.061 ± 0.024 | 2.3 | 0.0002 | | 1075 |
| Miscellaneous | Actin, cytoplasmic 2 | Actg | Down | 0.123 ± 0.071 | 0.039 ± 0.022 | 3.2 | 0.0402 | | 969 |
| | Coatomer subunit epsilon | Cope | Down | 0.529 ± 0.126 | 0.254 ± 0.027 | 2.1 | 0.0053 | | 500 |

Note: The normalized mean volumes of SAMR1 and SAMP8 differential protein spots and its ratio are indicated along with the changes shown by SAMP8. Normalized volumes were compared using a Student's *t*-test. Protein parameters are described in Table S2. ND, not detectable; NA, not applicable; SAMP8, senescence-accelerated prone mouse strain 8; SAMR1, senescence-accelerated resistant mouse strain 1. ^aFirst spot number for SAMR1 and second for SAMP8.

Results of the different expressed proteins in astrocyte cultures of SAMR1 (senescence-accelerated resistant mouse strain 1, control model) and SAMP8.



Beneficial effects of 8 weeks of exercise training in SAMP8 and SAMR1 model. Increasing levels of IGF-1 and BDNF (neurotrophic factors expressed in astrocytes) in both murine models due to the regular exercise. (R=running and S= sedentary).



Increased levels of sirtuin 1, a longevity protein, with treatments of melatonin and resveratrol in both animal models, SAMP8 and SAMR1.

CONCLUSIONS

1. There is an extremely relation between astrocytes and aging processes, where these cells will experiment some molecular changes that could compromise their neuroprotective capacity or their biological functions.
2. SASP is the phenotype that astrocytes will adopt in aging and SAMP8 and SAMR1 (used like a control) are animal models in relation of this phenotype.
3. There are some studies that confirm the proteomic differences between both models mentioned and also the beneficial effects of anti-oxidant components or regular exercise, which could improve the levels of neurotrophic factors or the expression of proteins that promote longevity.

References:

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2. Cristofol R, Porquet D, Corpas R, Coto-Montes A, Serret J, Camins A et al. Neurons from senescence-accelerated SAMP8 mice are protected against frailty by the sirtuin 1 promoting agents melatonin and resveratrol. *J Pineal Res* 2012; 52(3):271-281.
3. Díez-Vives C, Gay M, García-Matas S, Comellas F, Carrascal M, Abian J et al. Proteomic study of neuron and astrocyte cultures from senescence-accelerated mouse SAMP8 reveals degenerative changes. *J Neurochem* 2009; 111(4):945-955.
4. Cosin-Tomas M, Alvarez-Lopez MJ, Sanchez-Roige S, Lanza JF, Bayod S, Sanfeliu C et al. Epigenetic alterations in hippocampus of SAMP8 senescent mice and modulation by voluntary physical exercise. *Front Aging Neurosci* 2014; 6:51.