

Are Your Mitochondria Ready for a Space Odyssey?

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Trends in Endocrinology and Metabolism Are your mitochondria ready for a space odyssey? --Manuscript Draft--

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Abstract:	Anticipating very long space trips, NASA has performed pan-OMIC analyses on in- flight samples from astronauts, mice and cells. Results revealed major mitochondrial dysfunctions responsible for alterations in metabolism, immunity and circadian rhythm, which should prompt the evaluation of countermeasures to reduce the risks of future space odyssey, especially towards Mars.

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1	Spotlight
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3	Are your mitochondria ready for a space odyssey?
4	Salim Khiati ¹ , Dominique Bonneau ^{1,2} , Guy Lenaers ^{1,*}
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11	Keywords: space odyssey, OMICS, mitochondria, genome.
12	
13	Abstract (53 words)
14	Anticipating very long space trips, the NASA has performed pan-OMIC analyses on in-
15	flight samples from astronauts, mice and cells. Results revealed major mitochondrial
16	dysfunctions responsible for alterations in metabolism, immunity and circadian rhythm,
17	which should prompt the evaluation of countermeasures to reduce the risks of future
18	space odyssey, especially towards the planet Mars.

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20 Core text (800 words)

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22	Over the next decades, space flight will increase dramatically, including for recreational
23	purposes. The duration of space cruises will also increase, since it will take at least 400 to 500
24	days to make a round-trip flight to the planet Mars [1].
25	This odyssey will undoubtedly induce physiological changes on the passengers that must be
26	anticipated to reduce health problems. In a surprising study, entitled "Comprehensive multi-
27	OMICs analysis reveals mitochondrial stress as a central biological hub for spaceflight
28	impact", by Da Silveira et al. published in November 2020 in Cell, NASA and associated
29	teams addressed the molecular and biochemical consequences of space flights on cellular,
30	animal and human samples collected during previous missions. Their study revealed drastic
31	alterations of mitochondrial physiology as the main origin of most secondary metabolic and
32	physiological manifestations occurring during space flights.
33	The multi-OMICS approach including epigenomic, transcriptomic, proteomic and
34	metabolomic analyses were performed on cultured (fibroblasts and endothelial) and primary
35	(T cells and hair follicles) human cells, on multiple organs from the C57BL/6 and BALB/c
36	inbred mouse strains, as well as on urine and blood samples from 59 astronauts, the latter
37	including the NASA twin study data [2].
38	The main results emerging from the study of this remarkable collection of data concerned
39	changes in mitochondrial physiology occurring during space flights.
40	Indeed, OMIC signatures revealed a drastic uncoupling of the nuclear and mitochondrial
41	genome cross-talk, with a systemic down-regulation expression of nuclear genes linked to
42	energy production (i.e. electron transport, oxidative phosphorylation (OXPHOS) and ATP
43	synthesis), while a progressive 2 to 9 fold longitudinal overexpression of mitochondrial genes,

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paralleling increased mtDNA copy number, was observed, in particular in the blood cell
fraction of astronauts, proportional to the time spent in space.

46 This nucleo-mitochondrial genome uncoupling is responsible for misassembly of respiratory 47 chain complexes, decreased mitochondrial oxidative metabolism and increased production of 48 reactive oxygen species (ROS). In parallel, a metabolic shift from glycolysis to OXPHOS 49 respiration, as well as increased beta-oxidation of long-chain fatty acids and Krebs cycle were 50 observed in mouse skeletal muscle, but not in liver. In addition, the imbalance of the 51 OXPHOS complex subunits led to proteostasis failures triggering the mitochondrial unfolded protein response (UPR^{mt}), and eventually the activation of Gadd34 and of the integrated stress 52 53 response (ISR) in glycolytic muscle, while ISR biomarkers were downregulated in oxidative 54 muscles together with a downregulation of the FGF21 and GDF15 mitokines. Alterations of 55 mitochondrial physiology were further associated to the enrichment of genes associated to 56 DNA repair as well as excretion of 8OH-dG in urine, witnessing possible alterations of the 57 mitochondrial genome, and resulting in increased secretion of damage-associated molecular 58 patterns (DAMPS) and VEGF, IGF1, IL1 α , IL1 β and IL-1ra in blood, and up-regulation of 59 IFN-dependent innate immune pathways in internal organs. In addition, genes involved in cell 60 cycle were upregulated in all mouse tissues except T-cells, most likely to compensate for a 61 higher level of apoptosis, as were genes involved in circadian rhythm in all internal organs, 62 except liver, although animals were maintained in 12h/12h-light/dark cycle. Finally, the 63 decreased in the synthesis of 1.25 vitamin D and the increase in the metabolism of NAD are 64 further associated to the worsening of mitochondrial functions and alterations of DNA 65 integrity, respectively. Notably, epigenomic methylation signatures reflected the major transcriptomic modifications induced by space flight. Ultimately, most of these deleterious 66 67 changes reverted to normal in the weeks following the end of flight.

Altogether, these cellular modifications are witnessing processes related to premature aging, which could originate from major alterations of the mitochondrial genome, as loss of mtDNA integrity is clearly responsible for aging in mice [3,4] and humans [5]. In this regard, future assessments of mtDNA integrity in space-exposed samples are warranted to further reveal the initial mechanism responsible for the molecular changes reported in the Da Silveira paper, as mtDNA instability could be triggered both by the exposure to low doses of irradiation [6] and by the consequences of microgravity [7].

75 The effects of accelerated aging of space flight further result in somatic changes, including 76 loss of bone density, reduction in muscle mass and strength. Stiffening of the arteries, 77 overactivation of the immune response, increased cardiovascular risks and possibly thrombo-78 vascular events were also observed. Alopecia, a marker of premature aging in animal models, 79 is also affecting astronauts. Therefore, if astronauts (or future space tourists) do not wish to 80 land on Mars as "centenarians", countermeasures are needed to minimize the consequences of 81 long-duration space flights. In addition to daily muscular exercise, nutritional and 82 pharmacological supplementation targeting mitochondrial physiology are mandatory. The 83 "Mitochondrial Space Odyssey Cocktail" should be composed of anti-oxidants such as 84 CoQ10, to counteract the oxidative stress and promote mitochondrial electron transfer, 85 vitamin B3 to sustain NAD-related mitochondrial activity and DNA repair, and vitamin D to 86 maintain bone mineral density and arteries plasticity. And why not a bottle of red wine to 87 promote mitochondrial turnover and rejuvenation [8]?

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113 Figure Legend

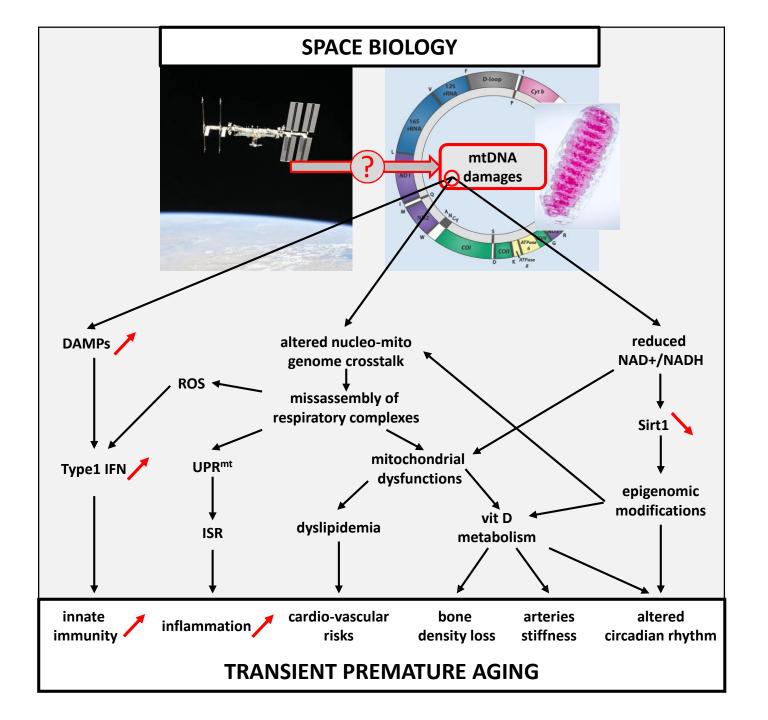
114	Synopsis of the molecular processes affecting mitochondrial physiology and the related
115	somatic consequences on health.

- 116 Exposure to low doses radiation and microgravity most probably induces mitochondrial
- 117 genome alterations that are responsible for the uncoupling of the nuclear and mitochondrial
- 118 genome cross-talk, the overproduction of DAMPs and changes in NAD metabolism, resulting
- 119 in downstream somatic effects reflecting a transient premature systemic aging process.
- 120
- 121

122 Abbreviations:

- 123 DAMP: damage-associated molecular patterns; IFN: interferon; ISR: integrated stress
- 124 response; mtDNA: mitochondrial DNA; NAD: nicotinamide adenine dinucleotide; OXPHOS:
- 125 oxidative phosphorylation; ROS: reactive oxygen species; UPR^{mt}: mitochondrial unfolded
- 126 protein response; vit D: vitamin D.

Figure



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