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Mendelian randomization of genetically independent aging phenotypes identifies LPA and VCAM1 as biological targets for human aging

DOI: 10.1038/s43587-021-00159-8 Nature Aging, 2022, 2, 19-30.

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9	Unravelling genetic components of longevity. <i>Nature Aging</i> , 2022 , 2, 5-6		
8	Determination of cross-tissue and tissue-specific aging changes in murine proteomes.		1
7	How Important Are Genes to Achieve Longevity?. <i>International Journal of Molecular Sciences</i> , 2022 , 23, 5635	6.3	1
6	Macrophage Class A Scavenger Receptors 🖪 Functional Perspective. 2022 ,		0
5	A Novel Framework for Analysis of the Shared Genetic Background of Correlated Traits. 2022 , 13, 1694		O
4	Mendelian randomization analyses reveal causal relationships between the human microbiome and longevity.		0
3	Causal Epigenetic Age Uncouples Damage and Adaptation.		O
2	Association of PM2.5 Components with Acceleration of Aging: Moderating Role of Sex Hormones. 2023 , 57, 3772-3782		0
1	Mendelian randomization analyses reveal causal relationships between the human microbiome and longevity. 2023 , 13,		O