

# CITATION REPORT

List of articles citing

**Genome-wide association meta-analysis of human longevity identifies a novel locus conferring survival beyond 90 years of age**

**DOI: 10.1093/hmg/ddu139**

**Human Molecular Genetics, 2014, 23, 4420-32.**

**Source:** <https://exaly.com/paper-pdf/59366090/citation-report.pdf>

**Version:** 2024-04-28

This report has been generated based on the citations recorded by exaly.com for the above article. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

#	Paper	IF	Citations
215	Why genes extending lifespan in model organisms have not been consistently associated with human longevity and what it means to translation research. <b>2014</b> , 13, 2671-3		42
214	Genetics and Genomics of Human Longevity. <b>2014</b> ,		
213	Associations between STR autosomal markers and longevity. <b>2015</b> , 37, 95		2
212	Genetic evidence for common pathways in human age-related diseases. <i>Aging Cell</i> , <b>2015</b> , 14, 809-17	9.9	59
211	Genome-Wide Scan Informed by Age-Related Disease Identifies Loci for Exceptional Human Longevity. <b>2015</b> , 11, e1005728		86
210	The Genomic Basis of Postponed Senescence in <i>Drosophila melanogaster</i> . <b>2015</b> , 10, e0138569		26
209	Genetic factors and epigenetic mechanisms of longevity: current perspectives. <b>2015</b> , 7, 1339-49		7
208	Genetics and underlying pathology of dementia. <b>2015</b> , 25, 113-24		22
207	Genetics and pharmacology of longevity: the road to therapeutics for healthy aging. <b>2015</b> , 90, 1-101		31
206	Current research in aging: a report from the 2015 Ageing Summit. <b>2015</b> , 5, 187-90		2
205	Genome-wide association studies of late-onset cardiovascular disease. <b>2015</b> , 83, 131-41		34
204	Genes and life-style factors in BELFAST nonagenarians: Nature, Nurture and Narrative. <b>2015</b> , 16, 587-97		16
203	Longevity Genes. <b>2015</b> ,		2
202	Models to explore genetics of human aging. <b>2015</b> , 847, 141-61		3
201	Exome and whole genome sequencing in aging and longevity. <b>2015</b> , 847, 127-39		5
200	GWAS of longevity in CHARGE consortium confirms APOE and FOXO3 candidacy. <b>2015</b> , 70, 110-8		188
199	MARK-AGE biomarkers of ageing. <b>2015</b> , 151, 2-12		145

198	Genome-Wide Association Study and Linkage Analysis of the Healthy Aging Index. <b>2015</b> , 70, 1003-8	13
197	The APP A673T frequency differs between Nordic countries. <b>2015</b> , 36, 2909.e1-4	7
196	Genetic factors associated with longevity: a review of recent findings. <b>2015</b> , 19, 1-7	65
195	Normal Aging and Dementia. <b>2016</b> ,	5
194	Intakes of Folate and Vitamin B12 and Biomarkers of Status in the Very Old: The Newcastle 85+ Study. <b>2016</b> , 8,	14
193	Novel loci and pathways significantly associated with longevity. <b>2016</b> , 6, 21243	105
192	Genetic variants near MLST8 and DHX57 affect the epigenetic age of the cerebellum. <b>2016</b> , 7, 10561	55
191	Genome sequencing in a case of Niemann-Pick type C. <b>2016</b> , 2, a001222	9
190	Analysis of FOXO1A and FOXO3A gene allele association with human longevity. <b>2016</b> , 52, 416-422	2
189	The MC1R Gene and Youthful Looks. <b>2016</b> , 26, 1213-20	42
188	Employing biomarkers of healthy ageing for leveraging genetic studies into human longevity. <b>2016</b> , 82, 166-74	21
187	Blood lipids influence DNA methylation in circulating cells. <b>2016</b> , 17, 138	118
186	Replication of Genome-Wide Association Study Findings of Longevity in White, African American, and Hispanic Women: The Women's Health Initiative. <b>2017</b> , 72, 1401-1406	10
185	Copy number variation associates with mortality in long-lived individuals: a genome-wide assessment. <i>Aging Cell</i> , <b>2016</b> , 15, 49-55	9.9 15
184	ImmunoChip analysis identifies association of the RAD50/IL13 region with human longevity. <i>Aging Cell</i> , <b>2016</b> , 15, 585-8	9.9 15
183	Tick tock: DNA methylation, the epigenetic clock and exceptional longevity. <b>2016</b> , 8, 1577-1582	6
182	Genome-wide Association Study of Parental Life Span. <b>2017</b> , 72, 1407-1410	5
181	Variants near CHRNA3/5 and APOE have age- and sex-related effects on human lifespan. <b>2016</b> , 7, 11174	55

180	Genetic analysis of variation in lifespan using a multiparental advanced intercross Drosophila mapping population. <b>2016</b> , 17, 113	17
179	Human longevity: Genetics or Lifestyle? It takes two to tango. <b>2016</b> , 13, 12	90
178	Exome-wide Association Study Identifies CLEC3B Missense Variant p.S106G as Being Associated With Extreme Longevity in East Asian Populations. <b>2017</b> , 72, 309-318	11
177	Genetics: Healthy ageing, the genome and the environment. <b>2016</b> , 12, 378-80	3
176	Common variants in SIRT1 and human longevity in a Chinese population. <b>2016</b> , 17, 31	14
175	Lack of association between polymorphisms in the SIRT6 gene and longevity in a Chinese population. <b>2016</b> , 30, 79-82	5
174	Genetics of Human Aging. <b>2016</b> , 327-358	
173	Longitudinal RNA-Seq Analysis of Vertebrate Aging Identifies Mitochondrial Complex I as a Small-Molecule-Sensitive Modifier of Lifespan. <b>2016</b> , 2, 122-32	88
172	Association of common variants in TOMM40/APOE/APOC1 region with human longevity in a Chinese population. <b>2016</b> , 61, 323-8	12
171	Understanding age-related diseases: report of the 2015 Ageing Summit. <b>2016</b> , 47, 5-9	1
170	A novel strain of Lactobacillus mucosae isolated from a Gaotian villager improves in vitro and in vivo antioxidant as well as biological properties in D-galactose-induced aging mice. <b>2016</b> , 99, 903-914	32
169	Human-specific derived alleles of CD33 and other genes protect against postreproductive cognitive decline. <b>2016</b> , 113, 74-9	50
168	Genetic Variants in KLOTHO Associate With Cognitive Function in the Oldest Old Group. <b>2016</b> , 71, 1151-9	33
167	A Genetic Network Associated With Stress Resistance, Longevity, and Cancer in Humans. <b>2016</b> , 71, 703-12	23
166	How the effects of aging and stresses of life are integrated in mortality rates: insights for genetic studies of human health and longevity. <b>2016</b> , 17, 89-107	21
165	Living long and ageing well: is epigenomics the missing link between nature and nurture?. <b>2016</b> , 17, 33-54	19
164	Increasing Sibling Relative Risk of Survival to Older and Older Ages and the Importance of Precise Definitions of "Aging," "Life Span," and "Longevity". <b>2016</b> , 71, 340-6	46
163	Conserved regulators of cognitive aging: From worms to humans. <b>2017</b> , 322, 299-310	16

162	Assessment of health status by molecular measures in adults ranging from middle-aged to old: Ready for clinical use?. <b>2017</b> , 87, 175-181	8
161	Associations of triglyceride levels with longevity and frailty: A Mendelian randomization analysis. <b>2017</b> , 7, 41579	12
160	Limitations and risks of meta-analyses of longevity studies. <b>2017</b> , 165, 139-146	23
159	Rapamycin additively extends lifespan in short- and long-lived lines of the nematode <i>Caenorhabditis remanei</i> . <b>2017</b> , 90, 79-82	3
158	Demographic, phenotypic, and genetic characteristics of centenarians in Okinawa and Honshu, Japan: Part 2 Honshu, Japan. <b>2017</b> , 165, 80-85	14
157	Genetic interplay between human longevity and metabolic pathways - a large-scale eQTL study. <i>Aging Cell</i> , <b>2017</b> , 16, 716-725	9.9 11
156	Cancer-Incidence, prevalence and mortality in the oldest-old. A comprehensive review. <b>2017</b> , 164, 113-126	36
155	Disease variants alter transcription factor levels and methylation of their binding sites. <b>2017</b> , 49, 131-138	252
154	Identification of context-dependent expression quantitative trait loci in whole blood. <b>2017</b> , 49, 139-145	240
153	Four Genome-Wide Association Studies Identify New Extreme Longevity Variants. <b>2017</b> , 72, 1453-1464	60
152	Genomewide Association Scan of a Mortality Associated Endophenotype for a Long and Healthy Life in the Long Life Family Study. <b>2017</b> , 72, 1411-1416	4
151	The genetics of human longevity: an intricacy of genes, environment, culture and microbiome. <b>2017</b> , 165, 147-155	61
150	Cognitive status in the oldest old and centenarians: a condition crucial for quality of life methodologically difficult to assess. <b>2017</b> , 165, 185-194	24
149	Differential Aging Analysis in Human Cerebral Cortex Identifies Variants in TMEM106B and GRN that Regulate Aging Phenotypes. <b>2017</b> , 4, 404-415.e5	65
148	Demographics, phenotypic health characteristics and genetic analysis of centenarians in China. <b>2017</b> , 165, 86-97	38
147	Genome-wide meta-analysis associates HLA-DQA1/DRB1 and LPA and lifestyle factors with human longevity. <b>2017</b> , 8, 910	78
146	What has GWAS done for HLA and disease associations?. <b>2017</b> , 44, 195-211	35
145	Bayesian association scan reveals loci associated with human lifespan and linked biomarkers. <b>2017</b> , 8, 15842	39

144	The Business of Anti-Aging Science. <b>2017</b> , 35, 1062-1073	80
143	Why and how are we living longer?. <b>2017</b> , 102, 1067-1074	47
142	Cohort Profile: The 1895, 1905, 1910 and 1915 Danish Birth Cohort Studies - secular trends in the health and functioning of the very old. <b>2017</b> , 46, 1746-1746j	23
141	Human longevity: 25 genetic loci associated in 389,166 UK biobank participants. <b>2017</b> , 9, 2504-2520	83
140	Genetic Factors Associated with Longevity in Humans. <b>2017</b> ,	1
139	Quantitative analysis of population-scale family trees with millions of relatives. <b>2018</b> , 360, 171-175	94
138	Genetics of Human Longevity From Incomplete Data: New Findings From the Long Life Family Study. <b>2018</b> , 73, 1472-1481	15
137	The genetic component of human longevity: New insights from the analysis of pathway-based SNP-SNP interactions. <i>Aging Cell</i> , <b>2018</b> , 17, e12755	9.9 14
136	Genetic cartography of longevity in humans and mice: Current landscape and horizons. <b>2018</b> , 1864, 2718-2732	18
135	Lowbush cranberry acts through DAF-16/FOXO signaling to promote increased lifespan and axon branching in aging posterior touch receptor neurons. <b>2018</b> , 40, 151-162	10
134	High-sensitivity C-reactive protein, low-grade systemic inflammation and type 2 diabetes mellitus: A two-sample Mendelian randomization study. <b>2018</b> , 28, 795-802	10
133	Phenome and genome based studies into human ageing and longevity: An overview. <b>2018</b> , 1864, 2742-2751	21
132	Hidden heterogeneity in Alzheimer's disease: Insights from genetic association studies and other analyses. <b>2018</b> , 107, 148-160	27
131	Why Does the Shift from "Personalized Medicine" to "Precision Health" and "Wellness Genomics" Matter?. <b>2018</b> , 20, E881-890	23
130	Genetics of Human Longevity Within an Eco-Evolutionary Nature-Nurture Framework. <b>2018</b> , 123, 745-772	46
129	Uncovering Natural Longevity Alleles from Intercrossed Pools of Aging Fission Yeast Cells. <b>2018</b> , 210, 733-744	7
128	Autosomal genetic variation is associated with DNA methylation in regions variably escaping X-chromosome inactivation. <b>2018</b> , 9, 3738	12
127	Facing up to the global challenges of ageing. <b>2018</b> , 561, 45-56	342

126	Sex Differences in Genetic Associations With Longevity. <b>2018</b> , 1, e181670	40
125	Apolipoprotein E in Cardiovascular Diseases: Novel Aspects of an Old-fashioned Enigma. <b>2018</b> , 49, 522-529	16
124	A locus at 7p14.3 predisposes to refractory celiac disease progression from celiac disease. <b>2018</b> , 30, 828-837	16
123	Cardiac ageing: extrinsic and intrinsic factors in cellular renewal and senescence. <b>2018</b> , 15, 523-542	59
122	Age and Age-Related Diseases: Role of Inflammation Triggers and Cytokines. <b>2018</b> , 9, 586	406
121	Weighted mining of massive collections of [Formula: see text]-values by convex optimization. <b>2018</b> , 7, 251-275	2
120	Genome-wide identification of directed gene networks using large-scale population genomics data. <b>2018</b> , 9, 3097	13
119	Genetics of Human Aging. <b>2018</b> , 1025-1039	
118	A decade in psychiatric GWAS research. <b>2019</b> , 24, 378-389	40
117	Whole-Exome Sequencing of an Exceptional Longevity Cohort. <b>2019</b> , 74, 1386-1390	9
116	A meta-analysis of genome-wide association studies identifies multiple longevity genes. <b>2019</b> , 10, 3669	102
115	Genetic Syndromes and Aging. <b>2019</b> , 211-239	
114	Genetic background, epigenetic factors and dietary interventions which influence human longevity. <b>2019</b> , 20, 605-626	17
113	The impact of APOE genotype on survival: Results of 38,537 participants from six population-based cohorts (E2-CHARGE). <b>2019</b> , 14, e0219668	31
112	Obesity Paradox in Caucasian Seniors: Results of the PolSenior Study. <b>2019</b> , 23, 796-804	13
111	Genetic Support for Longevity-Enhancing Drug Targets: Issues, Preliminary Data, and Future Directions. <b>2019</b> , 74, S61-S71	1
110	Biomarkers of Human Aging. <b>2019</b> ,	5
109	The Danish Twin Registry: An Updated Overview. <b>2019</b> , 22, 499-507	18

108	Identification of 12 genetic loci associated with human healthspan. <b>2019</b> , 2, 41	49
107	Gene-Lifestyle Interactions in Longevity. <b>2019</b> , 91-109	1
106	Apolipoprotein E gene in physiological and pathological aging. <b>2019</b> , 178, 41-45	12
105	A nonsynonymous mutation in PLCG2 reduces the risk of Alzheimer's disease, dementia with Lewy bodies and frontotemporal dementia, and increases the likelihood of longevity. <b>2019</b> , 138, 237-250	50
104	Developments in molecular epidemiology of aging. <b>2019</b> , 3, 411-421	8
103	Genomic and Epigenomic Potential With Age: Genome, Epigenome, and the Epigenetic Clock. <b>2019</b> , 445-459	
102	The Genetics of Aging: A Vertebrate Perspective. <b>2019</b> , 177, 200-220	95
101	Adult Cardiac Stem Cell Aging: A Reversible Stochastic Phenomenon?. <b>2019</b> , 2019, 5813147	24
100	Neuropathology-driven Whole-genome Sequencing Study Points to Novel Candidate Genes for Healthy Brain Aging. <b>2019</b> , 33, 7-14	1
99	Exonic Variants in Aging-Related Genes Are Predictive of Phenotypic Aging Status. <b>2019</b> , 10, 1277	1
98	A Prospective Analysis of Genetic Variants Associated with Human Lifespan. <b>2019</b> , 9, 2863-2878	23
97	Genetic and epigenetic regulation of human aging and longevity. <b>2019</b> , 1865, 1718-1744	37
96	The rs2516839 variation of USF1 gene is associated with 4-year mortality of nonagenarian women: The Vitality 90+ study. <b>2019</b> , 83, 34-45	2
95	Longevity defined as top 10% survivors and beyond is transmitted as a quantitative genetic trait. <b>2019</b> , 10, 35	35
94	The ApoE $\epsilon$ Isoform: Can the Risk of Diseases be Reduced by Environmental Factors?. <b>2019</b> , 74, 99-107	13
93	Roles of tau pathology in the locus coeruleus (LC) in age-associated pathophysiology and Alzheimer's disease pathogenesis: Potential strategies to protect the LC against aging. <b>2019</b> , 1702, 17-28	34
92	The genetics of human ageing. <b>2020</b> , 21, 88-101	86
91	The Impact of Apolipoprotein E Genetic Variability in Health and Life Span. <b>2020</b> , 75, 1855-1857	0



90	Activity recognition using wearable sensors for tracking the elderly. <b>2020</b> , 30, 567-605	13
89	Genome-wide identification of genes regulating DNA methylation using genetic anchors for causal inference. <b>2020</b> , 21, 220	10
88	A Long-Read Sequencing Approach for Direct Haplotype Phasing in Clinical Settings. <b>2020</b> , 21,	3
87	APOE2: protective mechanism and therapeutic implications for Alzheimer's disease. <b>2020</b> , 15, 63	35
86	Longevity Relatives Count score identifies heritable longevity carriers and suggests case improvement in genetic studies. <i>Aging Cell</i> , <b>2020</b> , 19, e13139	9.9 5
85	Vascular ageing in hypertension: Focus on mitochondria. <b>2020</b> , 189, 111267	6
84	Deconvolution of bulk blood eQTL effects into immune cell subpopulations. <b>2020</b> , 21, 243	15
83	GWAS for Lifespan and Decline in Climbing Ability in Flies upon Dietary Restriction Reveal decima as a Mediator of Insulin-like Peptide Production. <b>2020</b> , 30, 2749-2760.e3	13
82	Aging Biomarkers: From Functional Tests to Multi-Omics Approaches. <b>2020</b> , 20, e1900408	15
81	The Medical Genome Reference Bank contains whole genome and phenotype data of 2570 healthy elderly. <b>2020</b> , 11, 435	20
80	Cohort Differences in the Associations of Selected Candidate Genes With Risk of All-Cause Mortality at Advanced Ages. <b>2020</b> , 189, 708-716	0
79	Genetic Basis of Increased Lifespan and Postponed Senescence in. <b>2020</b> , 10, 1087-1098	0
78	Decline in biological resilience as key manifestation of aging: Potential mechanisms and role in health and longevity. <b>2021</b> , 194, 111418	13
77	Exome-Wide Association Study Identifies FN3KRP and PGP as New Candidate Longevity Genes. <b>2021</b> , 76, 786-795	4
76	A joint analysis of longevity and age-related disease variants for gene expression association.	
75	Cardiovascular Aging and Longevity: JACC State-of-the-Art Review. <b>2021</b> , 77, 189-204	14
74	Longevity as a complex genetic trait. <b>2021</b> , 3-42	
73	Genetic factors influencing a neurobiological substrate for psychiatric disorders. <b>2021</b> , 11, 192	0

72	Integrated genetic analyses revealed novel human longevity loci and reduced risks of multiple diseases in a cohort study of 15,651 Chinese individuals. <i>Aging Cell</i> , <b>2021</b> , 20, e13323	9.9	7
71	The genetic predisposition to longevity acts through behavioral phenotypes in females. <b>2021</b> , 45, 1-14		1
70	Intelligence, health and death. <b>2021</b> , 5, 416-430		10
69	Omics in a Digital World: The Role of Bioinformatics in Providing New Insights Into Human Aging. <b>2021</b> , 12, 689824		2
68	Metabolic regulation in mitochondria as a prospective way of body rejuvenation. <b>2021</b> , 78-82		
67	Genetic signature of human longevity in PKC and NF- $\kappa$ B signaling. <i>Aging Cell</i> , <b>2021</b> , 20, e13362	9.9	2
66	Extending human healthspan and longevity: a symposium report. <b>2021</b> ,		0
65	Centenarians as models of healthy aging: Example of REST. <b>2021</b> , 70, 101392		0
64	Improved selection of participants in genetic longevity studies: family scores revisited. <b>2021</b> , 21, 7		0
63	Cytokine Expression and Production Changes in Very Old Age. <b>2018</b> , 1-24		1
62	Quantitative analysis of population-scale family trees using millions of relatives.		8
61	Genome-wide analysis reveals distinct genetic mechanisms of diet-dependent lifespan and healthspan in <i>D. melanogaster</i> .		4
60	The burden of rare protein-truncating genetic variants on human lifespan.		2
59	Identification of 12 genetic loci associated with human healthspan.		3
58	Genomic underpinnings of lifespan allow prediction and reveal basis in modern risks.		1
57	Deconvolution of bulk blood eQTL effects into immune cell subpopulations.		3
56	Biases in GWAS [the dog that did not bark.		4
55	Genetic factors influencing a neurobiological substrate for psychiatric disorders.		2

54	Genome-wide identification of genes regulating DNA methylation using genetic anchors for causal inference.	1
53	Molecular therapies delaying cardiovascular aging: disease- or health-oriented approaches. <b>2020</b> , 2, R45-R58	2
52	Decoupling of DNA Methylation Status and Gene Expression Levels in Aging Individuals. <b>2018</b> , 4, 100040	1
51	Decreased epigenetic age of PBMCs from Italian semi-supercentenarians and their offspring. <b>2015</b> , 7, 1159-70	211
50	Human longevity is influenced by many genetic variants: evidence from 75,000 UK Biobank participants. <b>2016</b> , 8, 547-60	84
49	Assessment of the contribution of APOE gene variants to metabolic phenotypes associated with familial longevity at middle age. <b>2016</b> , 8, 1790-801	4
48	Investigation of the 5q33.3 longevity locus and age-related phenotypes. <b>2017</b> , 9, 247-255	7
47	Identification of new genetic variants of HLA-DQB1 associated with human longevity and lipid homeostasis-a cross-sectional study in a Chinese population. <b>2017</b> , 9, 2316-2333	11
46	The correlation of copy number variations with longevity in a genome-wide association study of Han Chinese. <b>2018</b> , 10, 1206-1222	12
45	Positive association of familial longevity with the moderate-high HDL-C concentration in Bama Aging Study. <b>2018</b> , 10, 3528-3540	9
44	Identification of novel genes associated with longevity in - a computational approach. <b>2019</b> , 11, 11244-11267	4
43	Epigenetics: Linking Nutrition to Molecular Mechanisms in Aging. <b>2017</b> , 22, 81-89	19
42	Genomics of 1 million parent lifespans implicates novel pathways and common diseases and distinguishes survival chances. <b>2019</b> , 8,	82
41	Biomedical Research. <b>2015</b> , 27-38	
40	The Aging Superorganism. <b>2016</b> , 265-290	
39	TECPR2 a positive regulator of autophagy is implicated in healthy brain ageing.	0
38	Cytokine Polymorphisms, Immunosenescence, and Neurodegeneration. <b>2018</b> , 1-34	
37	Genome-wide identification of directed gene networks using large-scale population genomics data.	1

- 36 Uncovering Natural Longevity Alleles from Intercrossed Pools of Aging Fission Yeast Cells. 0
- 35 Longevity defined as top 10% survivors is transmitted as a quantitative genetic trait: results from large three-generation datasets.
- 34 The Medical Genome Reference Bank: Whole genomes and phenotype of 2,570 healthy elderly. 1
- 33 Genetic Markers of Extreme Human Longevity. **2019**, 137-153
- 32 Homeostasis, Homeodynamics and Aging. **2019**, 2
- 31 Cytokine Polymorphisms, Immunosenescence, and Neurodegeneration. **2019**, 1057-1090
- 30 Encyclopedia of Gerontology and Population Aging. **2019**, 1-5
- 29 Cytokine Expression and Production Changes in Very Old Age. **2019**, 1335-1358
- 28 Longevity Relatives Count score identifies heritable longevity carriers and suggests case improvement in genetic studies.
- 27 Encyclopedia of Gerontology and Population Aging. **2020**, 1-6
- 26 Exome-wide association studies in general and long-lived populations identify genetic variants related to human age.
- 25 De biologie van veroudering. **2020**, 11-16
- 24 Biological Health and Homeodynamic Space. **2020**, 43-51 1
- 23 Identification of cardiovascular health gene variants related to longevity in a Chinese population. **2020**, 12, 16775-16802 0
- 22 Encyclopedia of Gerontology and Population Aging. **2021**, 2479-2484
- 21 Encyclopedia of Gerontology and Population Aging. **2021**, 2060-2064
- 20 Heterogeneity in Alzheimer's Disease Diagnosis and Progression Rates: Implications for Therapeutic Trials.. **2022**, 1 2
- 19 Comprehensive Statistical and Bioinformatics Analysis in the Deciphering of Putative Mechanisms by Which Lipid-Associated GWAS Loci Contribute to Coronary Artery Disease.. **2022**, 10, 1

18	The burden of rare protein-truncating genetic variants on human lifespan.		0
17	No Evidence That Genetic Variation At The Klotho Locus Is Associated With Longevity In Caucasians From The Newcastle 85 Plus Study And The Uk Biobank. <b>2021</b> ,		0
16	Association of rs3027178 polymorphism in the circadian clock gene PER1 with susceptibility to Alzheimer's disease and longevity in an Italian population.. <b>2021</b> , 1		0
15	Associations of APOE Gene Variants rs429358 and rs7412 with Parameters of the Blood Lipid Profile and the Risk of Myocardial Infarction and Death in a White Population of Western Siberia. <i>Current Issues in Molecular Biology</i> , <b>2022</b> , 44, 1713-1724	2.9	
14	DataSheet_1.pdf. <b>2019</b> ,		
13	Intermediate alleles of HTT: A new pathway in longevity.. <i>Journal of the Neurological Sciences</i> , <b>2022</b> , 438, 120274	3.2	
12	Genetic associations with healthy ageing among Chinese adults. <b>2022</b> , 8,		
11	Insights Into Sibling Relationships and Longevity From Genetics of Healthy Ageing Nonagenarians: The Importance of Optimisation, Resilience and Social Networks. <i>Frontiers in Psychology</i> , <b>2022</b> , 13,	3.4	0
10	Genetic trade-offs between complex diseases and longevity. <i>Aging Cell</i> , <b>2022</b> , 21,	9.9	0
9	High polygenic risk score for exceptional longevity is associated with a healthy metabolic profile.	2	
8	TOMM40 genetic variants associated with healthy aging and longevity: a systematic review. <b>2022</b> , 22,		
7	Long-Lived Individuals Show a Lower Burden of Variants Predisposing to Age-Related Diseases and a Higher Polygenic Longevity Score. <b>2022</b> , 23, 10949		0
6	The aging cardiovascular system. <b>2023</b> , 109-119		0
5	Current Trends and Approaches to the Search for Genetic Determinants of Aging and Longevity. <b>2022</b> , 58, 1427-1443		0
4	Lipid-induced transcriptomic changes in blood link to lipid metabolism and allergic response. <b>2023</b> , 14,		0
3	Genome-wide association study reveals BET1L associated with survival time in the 137,693 Japanese individuals. <b>2023</b> , 6,		0
2	Genetic scores for predicting longevity in the Croatian oldest-old population. <b>2023</b> , 18, e0279971		0
1	Aging Hallmarks and the Role of Oxidative Stress. <b>2023</b> , 12, 651		0

