

Population genomics of Bronze Age Eurasia

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Citation Report

#	ARTICLE	IF	CITATIONS
3	Human Trichuriasis: Whipworm Genetics, Phylogeny, Transmission and Future Research Directions. <i>Current Tropical Medicine Reports</i> , 2015, 2, 209-217.	1.6	26
4	Inferring the genetic history of lactase persistence along the Italian peninsula from a large genomic interval surrounding the <i>LCT</i> gene. <i>American Journal of Physical Anthropology</i> , 2015, 158, 708-718.	2.1	7
5	A method for positive forensic identification of samples from extremely low-coverage sequence data. <i>BMC Genomics</i> , 2015, 16, 1034.	1.2	18
7	Distribution of two OCA2 polymorphisms associated with pigmentation in East-Asian populations. <i>Human Genome Variation</i> , 2015, 2, 15058.	0.4	11
8	Regional Differences in the Accumulation of SNPs on the Male-Specific Portion of the Human Y Chromosome Replicate Autosomal Patterns: Implications for Genetic Dating. <i>PLoS ONE</i> , 2015, 10, e0134646.	1.1	12
9	Fine Dissection of Human Mitochondrial DNA Haplogroup HV Lineages Reveals Paleolithic Signatures from European Glacial Refugia. <i>PLoS ONE</i> , 2015, 10, e0144391.	1.1	23
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19	The Role of Recent Admixture in Forming the Contemporary West Eurasian Genomic Landscape. <i>Current Biology</i> , 2015, 25, 2518-2526.	1.8	68
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21	Genome-wide patterns of selection in 230 ancient Eurasians. <i>Nature</i> , 2015, 528, 499-503.	13.7	1,160

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