

Positive selection on the H3 hemagglutinin gene of hum

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Predicting the Evolution of Human Influenza A. <i>Science</i> , 1999, 286, 1921-1925.	6.0	444
2	Factors Influencing Phylogenetic Inference: A Case Study Using the Mammalian Carnivores. <i>Molecular Phylogenetics and Evolution</i> , 2000, 16, 113-126.	1.2	23
3	Maximum Likelihood Estimation on Large Phylogenies and Analysis of Adaptive Evolution in Human Influenza Virus A. <i>Journal of Molecular Evolution</i> , 2000, 51, 423-432.	0.8	202
4	Effects of passage history and sampling bias on phylogenetic reconstruction of human influenza A evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 6974-6980.	3.3	88
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20	Assessment of Hemagglutinin Sequence Heterogeneity during Influenza Virus Transmission in Families. <i>Journal of Infectious Diseases</i> , 2002, 186, 1575-1581.	1.9	37
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