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Table S6: DAVID Gene ontology for retrogenes in the Oimyakon Mammoth

Function	EASE score
Ribosome	6.3
Post translational modification	4.4
Lipoproteins	3.4
Spliceosome	3.1
RNA binding	2.6
Lipoprotein metabolism	2.2
Nucleolus	2.0
Glutamine metabolism	1.9
Aspartate metabolism	1.8
Starch and drug metabolism	1.7
Proteasome	1.6
Translation initiation	1.6