Table S7: DAVID Gene ontology for retrogenes in the Wrangel Island Mammoth

Function	EASE score
Ribosome	8.3
Ubl conjugation	6.8
Spliceosome	4.3
Translation initiation	2.8
Lipoprotein	2.6
Nuclear body	2.3
Cytoskeleton	2.0
Aminoacylation	1.8
HEAT elongation	1.6
RNA splicing	1.6