

Table S3: Heterozygosity for Deletions Identified in Elephantid Genomes

		Oimyakon	Wrangel	Maya
All	Homozygous ¹	5035	6147	5141
	Hemizygous	16223	21081	22954
Exon Deletions	Homozygous	136	173	165
	Hemizygous	1347	1985	4248

¹ $\leq 10\%$ coverage