

Table S7. Levels and patterns of nucleotide variation at synonymous sites for highly (top 5%) and lowly (bottom 5%) expressed genes. Significant comparisons are highlighted in red.

Subspecies	Expression category	FPKM ^a	$\pi(\%)^b$	D_t^c
<i>O. c. algirus</i>	Highly	552.4	0.912	-0.447
	Lowly	8.0	0.863	-0.318
<i>O. c. cuniculus</i>	Highly	582.7	0.626	-0.445
	Lowly	7.8	0.870	-0.217

Unweighted average values among genes. Genes with less than 50 bp of synonymous sites were excluded.

^aMedian fragments per kilobase of gene model and million reads values among all *O. c. algirus* and *O. c. cuniculus* individuals and averaged across genes for each expression category.

^bAverage number of pairwise differences in a sample [68].

^cTajima's D [69].