

Table S2 *IRA1* and *IRA2* have high nucleotide diversity in *S. paradoxus* populations

Population ^a	Proportion CDS with $\pi > IRA1^b$	Proportion CDS with $\pi > IRA2^c$
European	.307	.004
American	.049	.071
Far East	.052	.037

^a *S. paradoxus* population as in (Liti et al. 2009).

^b The proportion of gene-coding regions (CDS) with higher nucleotide diversity (π) than *IRA1* across the indicated population.

^c The proportion of gene-coding regions with higher nucleotide diversity than *IRA2* across the indicated population.