



Figure S1 High presence/absence polymorphism in mitochondrial-encoded GC42 in comparison of five nuclear-encoded transposons (Ty1-Ty5) in *S. cerevisiae*. The presence/absence pattern of Ty transposons is obtained from Carr et al. (2012). The pairwise difference is calculated as $\frac{\# \text{ different}}{\# \text{ identical} + \# \text{ different}}$, sites with missing information are excluded from each pairwise comparison.