Table S3. dN/dS ratios determined for glycolytic genes in different *Saccharomyces* species comparisons.

Gene	dN/dS Spar/Scer	dN/dS Suva/Scer	dN/dS Skud/Scer	dN/dS Smik/Scer
HXK1	0.060	0.038	0.037	0.031
HXK2	0.009	**0.052**	0.040	0.021
GLK1	0.015	0.006	0.009	0.009
PGI1	0.003	0.018	0.028	0.006
PFK1	0.009	0.013	0.006	0.019
PFK2	0.021	0.025	0.020	0.006
FBA1	0.270	**0.278**	0.222	0.205
TPI1	0.045	–	**0.115**	–
TDH1	0.069	0.079	0.065	0.046
TDH2	0.115	–	0.099	0.053
TDH3	0.037	–	–	0.069
PGK1	0.100	**0.121**	0.066	0.068
GPM1	0.020	**0.226**	**0.100**	0.030
ENO1	0.016	**0.107**	**0.067**	0.021
ENO2	0.039	**0.168**	**0.148**	**0.125**
CDC19	0.065	**0.098**	–	0.076
PYK2	0.055	0.028	–	0.048

Spar: *S. paradoxus*; **Scer:***S. cerevisiae*; **Skud**: *S kudriavzevii*; **Smik:** *S. mikatae*
–: no ortholog identified
Values shown in red are outliers identified in each species comparison as shown in Figure 3 for *S. cerevisiae/S. uvarum.*