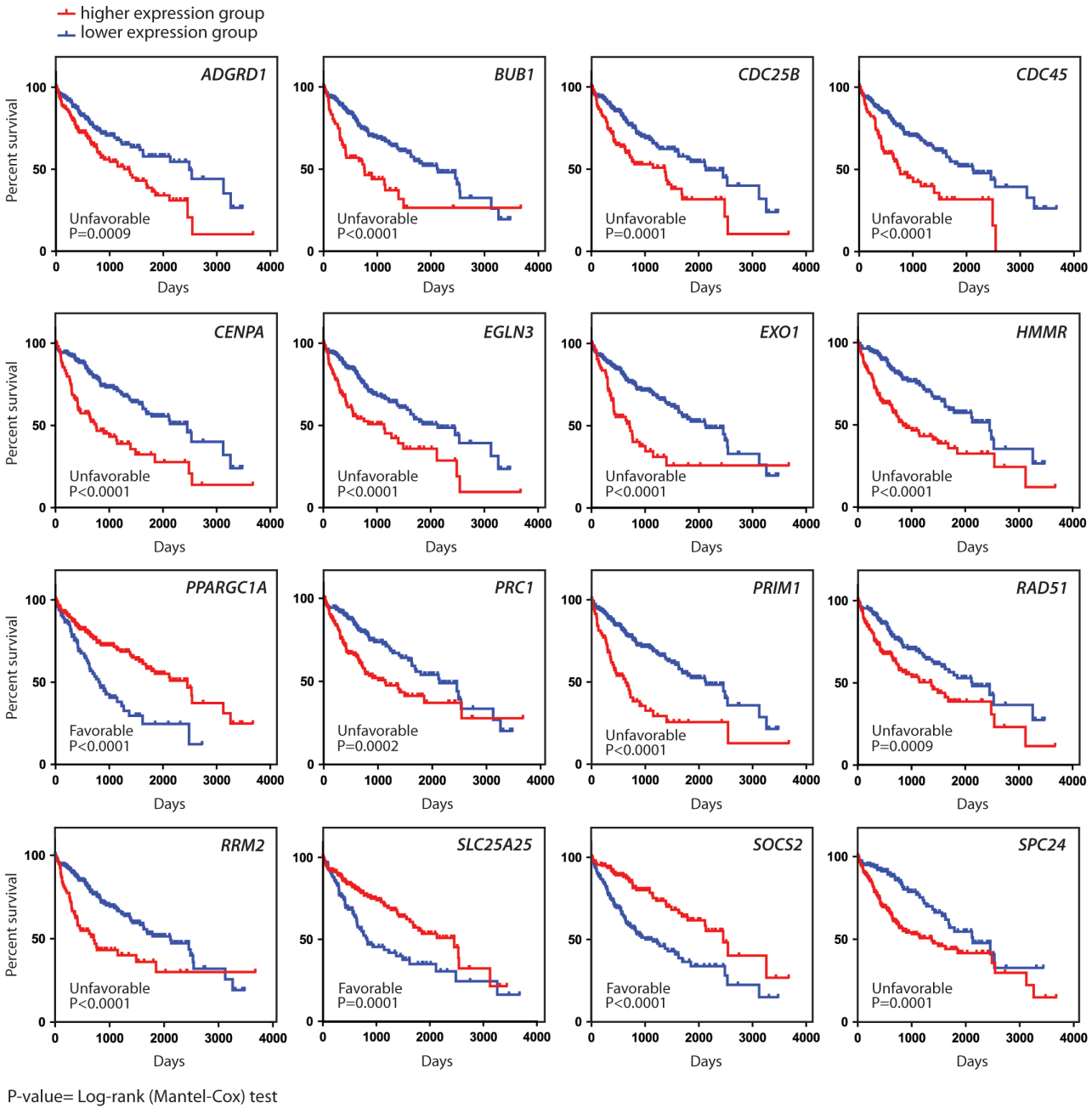

Figure S2

Survival rates of the high-expresser patients (red lines) and the low-expresser patients (blue lines) for the 16 TSPY downstream genes. Classification of the patients was based on the Human Protein Atlas data-portal (https://www.proteinatlas.org/). Log-rank test P-values calculated by using Prism6 are indicated. High expression levels of PPARGC1A, SLC25A25 and SOCS2 are associated with favorable survival rates while those of the remaining 13 genes are associated with unfavorable survival rates for HCC patients.