

Figure S1. OS curves. OS of patients with lung cancer with high expression levels of the identified 10 genes was compared with that of patients with low expression levels of the identified genes. For ABLIM3, GNG2, SLCO2B1, HTR1D, ROR1, TBXAS1, THBS1 and HSPA1A, two-stage P-values were calculated and these are presented for Kaplan-Meier curves showing crossover. For IL31RA and BCAT1, the log-rank P-value is presented for the Kaplan-Meier curve. OS, overall survival; HR, hazard ratio.

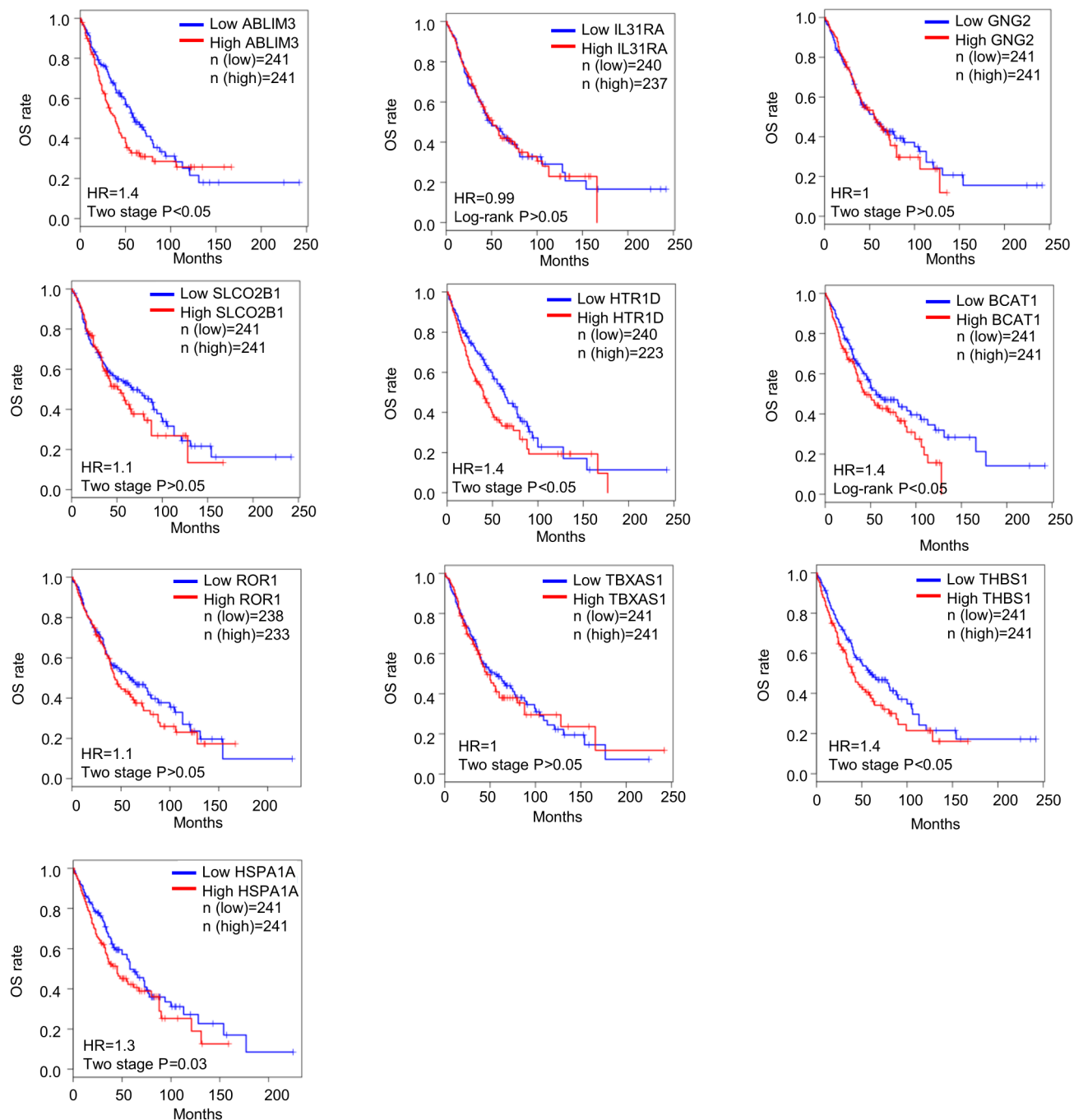


Figure S2. DFS curves. DFS of patients with lung cancer with high expression levels of the identified 10 genes was compared with that of patients with low expression levels of the identified genes. For TBXAS1 and HSPA1A, two-stage P-values were calculated and these are presented for the Kaplan-Meier curves showing crossover. For the other genes, the log-rank P-value is presented for the Kaplan-Meier curves. DFS, disease-free survival; HR, hazard ratio.

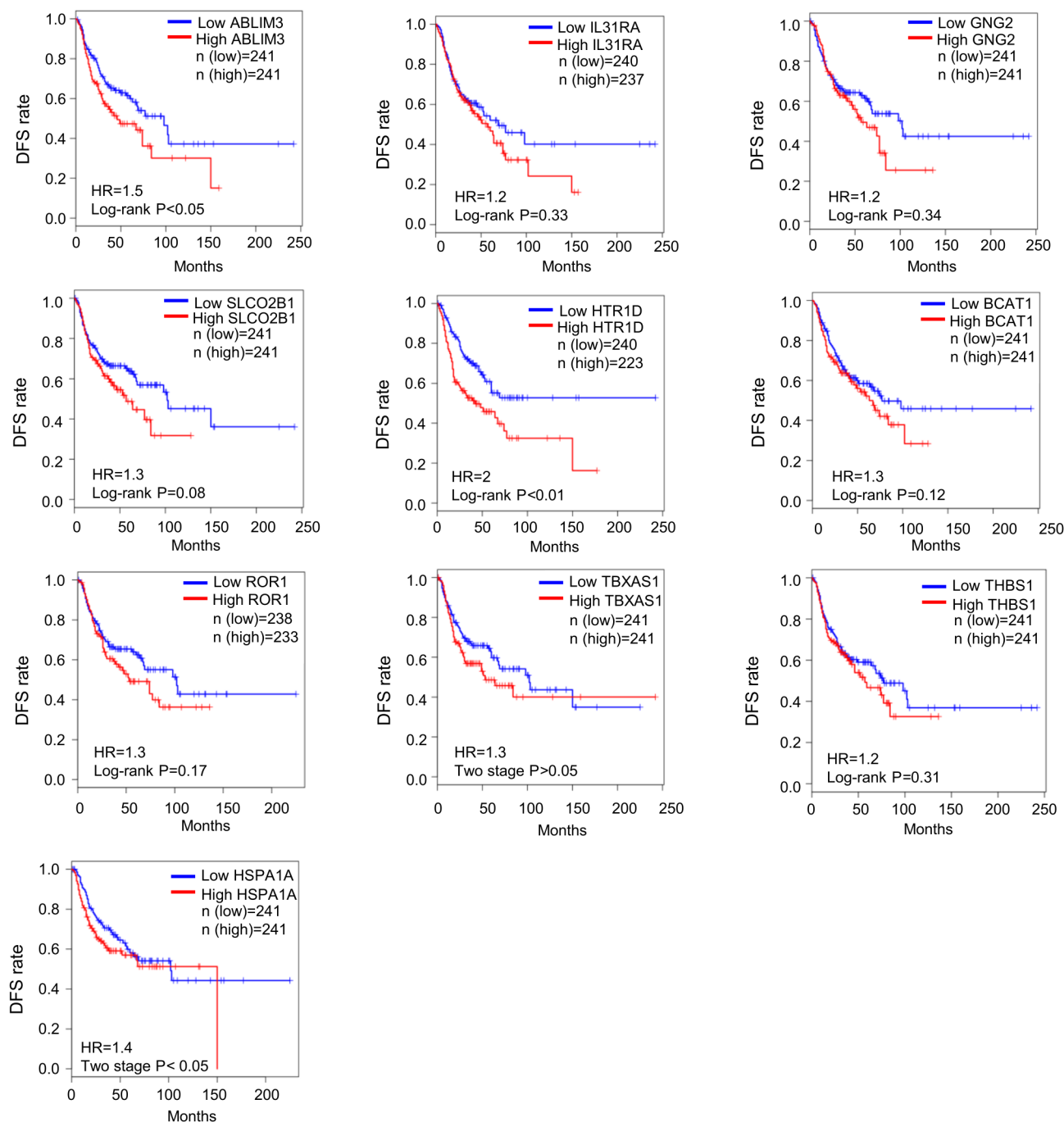


Figure S3. DFS curves for different cancer types. DFS of patients with (A) colon, (B) gastric, (C) lung, (D) liver and (E) pancreatic cancer was compared between groups with high and low expression of the three-gene signature. For liver and pancreatic cancer, two-stage P-values were calculated and these are presented for the Kaplan-Meier curves showing crossover. For colon, gastric and lung cancer, the log-rank P-value is presented for the Kaplan-Meier curves. DFS, disease-free survival; HR, hazard ratio.

