

Supplementary table 4 KEGG enrichment analysis of the thirteen prognostic genes derived from the predictive signature in COAD.

ID	Description	Count	p-value	q-value	Gene ID
hsa00983	Drug metabolism - other enzymes	2	0.002	0.014	NAT1/NAT2
hsa05204	Chemical carcinogenesis	2	0.002	0.014	NAT1/NAT2
hsa00020	Citrate cycle (TCA cycle)	1	0.026	0.087	SUCLG2
hsa00640	Propanoate metabolism	1	0.029	0.087	SUCLG2
hsa04960	Aldosterone-regulated sodium reabsorption	1	0.032	0.087	NR3C2