

Table S7. Univariable Cox regression analysis of the differential gene expression profile between high-risk and low-risk groups in overall survival for CRC patients

Gene	HR	HR.95L	HR.95H	pvalue
CLCA4	0.876134	0.769175	0.997966	0.046523
WNT10A	1.237252	1.008591	1.517754	0.04115
CDKN2A	1.281775	1.074598	1.528895	0.005784
VGF	1.193416	1.007586	1.413518	0.040611
ISYNA1	1.352037	1.125498	1.624173	0.001266
SFRP2	1.110648	1.015988	1.214127	0.020947
CLCA1	0.92715	0.864377	0.994482	0.03446
SERPINE1	1.202128	1.038627	1.391369	0.013585
CALB2	1.323927	1.133031	1.546985	0.000412
CSAG1	1.272312	1.115939	1.450596	0.000319
UCHL1	1.266232	1.045012	1.534282	0.01598
AC012354.2	1.295856	1.087289	1.54443	0.003796
TNNT1	1.273113	1.10939	1.460999	0.000586
MIR210	1.357849	1.082363	1.703451	0.008192
KREMEN2	1.477684	1.098216	1.988271	0.009918
FABP4	1.185065	1.049732	1.337846	0.006062
UGT2B7	0.804338	0.647403	0.999315	0.049282
AMH	1.310129	1.07154	1.601842	0.008448