

Table S1 Multivariate Cox regression analysis of the eight methylated genes signature and survival c

gene	exp(coef)	exp(-coef)	lower .95	upper .95	P
NBEA	4.87E+03	2.05E-04	2.34E+01	1.01E+06	0.0018
DDX28	1.19E-17	8.41E+16	6.82E-25	2.07E-10	4.63E-06
TMED8	4.03E+04	2.48E-05	1.99E-01	8.16E+09	0.08896
LOC151174	3.35E-03	2.98E+02	9.28E-06	1.21E+00	0.05798
EFNB2	1.81E+04	5.51E-05	1.18E+00	2.79E+08	0.0462
GHRHR	2.15E-02	4.65E+01	4.94E-04	9.39E-01	0.04627
MIMT1	1.08E+02	9.23E-03	3.09E+00	3.80E+03	0.00984
SLC29A3	2.26E+05	4.43E-06	1.14E-02	4.47E+12	0.15047

of neuroblastomas patients in the training grou