



Figure S1 The prognostic value of HNRNPM in our cohort.

Table S1 Gene set enrichment analysis of HNRNPM in KEGG genesets

KEGG gene sets name	Size	NES	P value
KEGG_SPLICEOSOME	123	2.509394	0
KEGG_CELL_CYCLE	124	2.404002	0
KEGG_RNA_DEGRADATION	55	2.37708	0
KEGG_PYRIMIDINE_METABOLISM	94	2.312553	0
KEGG_NUCLEOTIDE_EXCISION_REPAIR	44	2.276061	0
KEGG_PURINE_METABOLISM	150	2.273012	0
KEGG_HOMOLOGOUS_RECOMBINATION	27	2.263864	0
KEGG_BASE_EXCISION_REPAIR	34	2.205231	0
KEGG_MISMATCH_REPAIR	23	2.17423	0
KEGG_DNA_REPLICATION	36	2.127939	0
KEGG_UBIQUITIN_MEDIATED_PROTEOLYSIS	133	2.100534	0.001984
KEGG_OOCYTE_MEIOSIS	108	2.093884	0
KEGG_BASAL_TRANSCRIPTION_FACTORS	32	2.081933	0
KEGG_RNA_POLYMERASE	28	2.031649	0
KEGG_THYROID_CANCER	29	1.942894	0
KEGG_AMINOACYL_TRNA_BIOSYNTHESIS	41	1.933486	0.00616
KEGG_PROGESTERONE_MEDIATED_OOCYTE_MATURATION	84	1.906034	0
KEGG_PATHOGENIC_ESCHERICHIA_COLI_INFECTION	55	1.819869	0.011811
KEGG_RIBOSOME	87	1.749618	0.015123
KEGG_CYTOSOLIC_DNA_SENSING_PATHWAY	40	1.721275	0.015625