Genes	P-value	Hazard ratio
APC	0.021	1.511(1.063−2.149)
TP53	<0.001	1.788(1.273−2.513)
TTN	0.065	1.378(0.981−1.937)
KRAS	<0.001	2.183(1.533−3.108)
SYNE1	0.544	1.124(0.771−1.641)
PIK3CA	0.388	1.184(0.807−1.738)
MUC16	0.941	0.986(0.675−1.440)
FAT4	0.790	0.946(0.631−1.420)
CSMD3	0.262	1.276(0.833−1.954)
DNAH5	0.268	1.277(0.828−1.968)
RYR2	0.959	0.989(0.648−1.508)
ZFHX4	0.446	1.185(0.765−1.836)
OBSCN	0.736	0.928(0.600−1.434)
RYR3	0.648	1.115(0.700−1.775)
LRP1B	0.745	0.927(0.587−1.463)
NEB	0.657	1.113(0.695−1.780)
ABCA13	0.926	1.022(0.640−1.633)
PCLO	0.715	0.917(0.576−1.460)
CSMD1	0.783	0.936(0.584−1.500)
FAT3	0.465	0.839(0.525−1.342)
DST	0.753	0.925(0.572−1.498)

High Risk vs Low Risk

Hazard ratio