

Comprehensive immunogenomic landscape analysis of prognosis-related genes in head and neck cancer

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Gene	Normal Mean	Tumor Mean	logFC	P value	FDR
IL12A	0.555976095	0.223027363	-1.317802	0.000136	0.00026
TEC	1.46982345	0.60201631	-1.287768	6.31E-13	4.69E-12
SFTPA2	0.766671931	0.336438314	-1.188267	1.59E-10	7.76E-10
CXCL2	9.723854423	4.565688464	-1.090696	0.017695	0.024321
INHA	0.076166441	0.214585107	1.4943226	1.13E-06	2.98E-06
CSPG5	0.109208815	0.314350353	1.5252841	0.000161	0.000305
IL22RA2	0.1338516	0.453174929	1.7594337	6.36E-07	1.74E-06
IL1B	3.710741148	13.10607739	1.8204567	1.08E-06	2.85E-06
SSTR2	0.105505064	0.406339649	1.9453739	4.01E-08	1.32E-07
IGLV3-21	45.49312311	177.2342444	1.961937	0.005405	0.0081
BIRC5	5.051176243	20.54924905	2.0243944	2.61E-24	1.53E-21
CD70	0.308391571	2.87821674	3.22234	1.26E-15	1.65E-14
PDIA2	0.059697766	1.199043433	4.3280632	2.80E-05	5.92E-05