

A

Gene sets enriched in phenotype

	GS follow link to MSigDB	GS DETAILS	SIZE	ES	NES	NOM p-val	FDR q-val	FWER p-val	RANK AT MAX	LEADING EDGE
1	HALLMARK_TNFA_SIGNALING_VIA_NFKB	Details...	183	-0.52	-1.90	0.000	0.002	0.004	1506	tags=46%, list=9%, signal=51%
2	HALLMARK_EPITHELIAL_MESENCHYMAL_TRANSITION	Details...	125	-0.31	-1.38	0.045	0.228	0.471	1008	tags=20%, list=6%, signal=21%
3	HALLMARK_IL6_JAK_STAT3_SIGNALING	Details...	71	-0.37	-1.29	0.151	0.262	0.586	1139	tags=35%, list=7%, signal=38%
4	HALLMARK_KRAS_SIGNALING_DN	Details...	83	-0.30	-1.25	0.114	0.244	0.631	1156	tags=17%, list=7%, signal=18%
5	HALLMARK_INFLAMMATORY_RESPONSE	Details...	147	-0.31	-1.21	0.178	0.232	0.677	941	tags=27%, list=6%, signal=29%
6	HALLMARK_IL2_STAT5_SIGNALING	Details...	185	-0.29	-1.07	0.351	0.360	0.806	954	tags=26%, list=6%, signal=27%
7	HALLMARK_INTERFERON_GAMMA_RESPONSE	Details...	182	-0.29	-0.99	0.484	0.418	0.862	1139	tags=32%, list=7%, signal=34%
8	HALLMARK_KRAS_SIGNALING_UP	Details...	142	-0.24	-0.98	0.470	0.378	0.869	979	tags=20%, list=6%, signal=21%

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