

## SUPPLEMENTARY TABLE

Supplementary Table 5. Comparing the expression of genes in inflammation-activities signature between KRAS-mutated and wild-type groups.

Gene	WildMean	MutationMean	logFC	pValue
<b>IL12A</b>	0.282451	0.430636	0.608472	<b>3.36E-06</b>
<b>GZMB</b>	7.518245	13.26506	0.819164	<b>4.63E-06</b>
<b>CXCL9</b>	12.71954	17.27174	0.441366	<b>0.001911</b>
<b>CXCL10</b>	24.05325	29.22686	0.281063	<b>0.006641</b>
<b>PRF1</b>	3.191711	4.06926	0.350436	<b>0.009838</b>
<b>CXCL13</b>	4.056069	6.041906	0.574921	<b>0.020125</b>
<b>GNLY</b>	2.000802	2.894216	0.532595	<b>0.021077</b>
<b>IFNG</b>	0.27372	0.389159	0.507657	<b>0.028949</b>
<b>STAT1</b>	37.46403	44.23563	0.239703	<b>0.049644</b>
<b>TBX21</b>	0.247669	0.293948	0.247147	0.061036
<b>IRF1</b>	15.09839	17.20898	0.188767	0.064267
<b>CCL5</b>	12.90831	15.46207	0.260433	0.186282
<b>CD8B</b>	1.457315	1.674802	0.200678	0.279428
<b>CD19</b>	0.511267	0.742265	0.537859	0.283726
<b>IL12B</b>	0.033961	0.038595	0.184569	0.507014