

**Supplementary Table 4. The result of differential analysis of gene expression between normal tissue and tumor tissue by R package “limma”.**

<b>Gene</b>	<b>logFC</b>	<b>AveExpr</b>	<b>t</b>	<b>P.Value</b>	<b>adj.P.Val</b>	<b>B</b>
CCL5	2.761844	4.760074	16.69318	8.34E-52	1.82E-50	106.7216
CD2	2.28705	3.233462	15.32089	4.60E-45	7.75E-44	91.25291
CXCL9	2.625227	3.859447	13.8642	3.42E-38	4.27E-37	75.49697
CD3D	2.063852	2.945952	13.82764	5.04E-38	6.23E-37	75.11169
CCR5	1.667775	2.207914	13.2961	1.32E-35	1.44E-34	69.57093
CD8A	2.076911	2.644136	12.42316	9.51E-32	8.71E-31	60.73685
CD3G	1.022994	1.294262	10.84224	3.62E-25	2.40E-24	45.68733
IRF1	0.899948	3.35192	8.877243	7.60E-18	3.42E-17	28.98526
LCK	0.966669	2.169819	8.614633	6.00E-17	2.55E-16	26.94321
IFNG	0.555027	0.53599	6.865693	1.63E-11	5.02E-11	14.62135