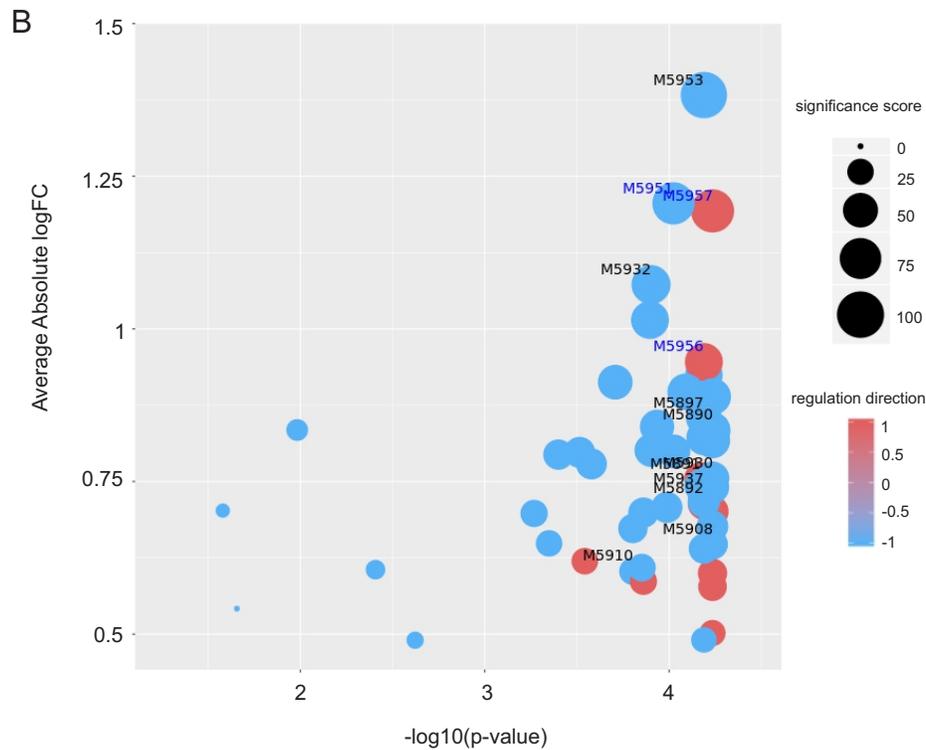
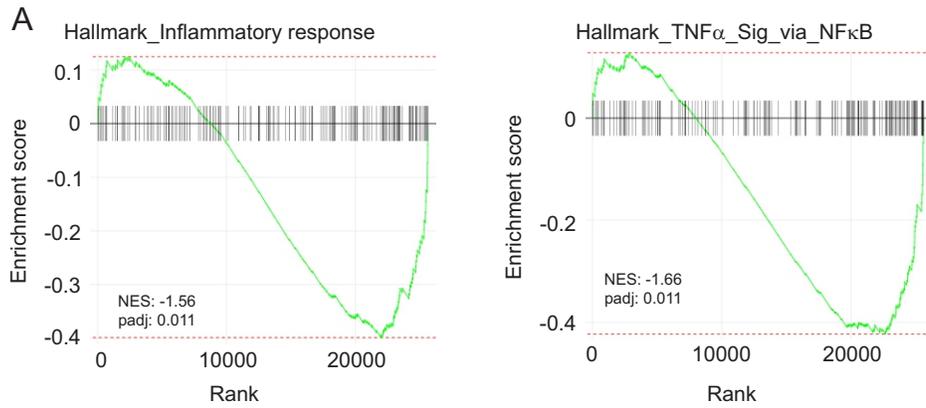


Figure S5



C

	ID	size	rank	p.adj	avg.logFC
Fatty acid metabolism	M5935	151/158	12	5.82e-05	0.60
Hypoxia	M5891	197/200	6	7.74e-05	0.754
Inflammatory response	M5932	185/200	8	1.25e-04	-1.072
TNF α _Sig_via_NF κ B	M5890	192/200	7	5.82e-05	-0.835

Figure S5. Enrichment of h-Hallmarks Gene Set collection in ACHN OIP5 tumors. **(A)** Enrichment was analyzed using FGSEA within the Galaxy platform. **(B, C)** Enrichment analyses were performed using EGSEA by UseGalaxy.org. The summary plot of mean absolute fold change ($\log_2\text{FC}$) and p.adj value (B). The details of the indicated gene sets, which were also derived from FGSEA analysis [panel A, Figure 3C], are presented (C).