

Table S7. Representative differentially expressed mRNA between *Fusobacterium nucleatum* treated and untreated HCT116 cells

Gene id	log2FoldChange	log2CPM	P value	FDR
NOS3	-0.42487	3.674845	0.009955	0.122415
VEGFB	-0.36375	3.260545	0.034425	0.21971
AKT1	-0.19661	7.452499	0.016224	0.157794
AKT2	-0.30243	6.152073	0.0022	0.054905
KEAP1	-0.32748	5.625378	0.00076	0.028799
AKT1S1	-0.36808	4.34958	0.005293	0.088878
PIK3R2	-0.25044	5.041885	0.043213	0.247248
HSP90B1	0.188804	10.63177	0.026438	0.197212
PIK3CB	0.257125	5.831532	0.047033	0.258099
CYP2J2	0.501714	2.107456	0.048475	0.262794
PRKCA	0.220682	5.986045	0.008934	0.115418
TLR6	0.543821	2.617278	0.023012	0.184886