

Supplemental Figures

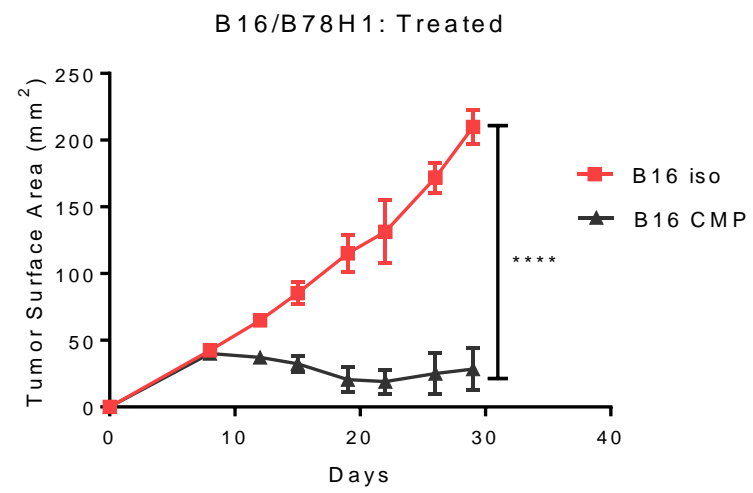


Figure S1: Average tumor growth curve of treated B16F10 tumors in animals bearing distant B78H1 tumors.

HGNCSymbol	Gene Symbol	P.Value	adj.P.Val	logFC	ENTREZID	
GZMK	Gzmk	6.62E-07	0.019109	3.067493	3003	
ADAM8	Adam8	5.06E-05	0.063651	3.123511	101	
ANPEP	Anpep	3.65E-05	0.063651	2.428938	290	
ASS1	Ass1	3.45E-05	0.063651	1.581637	445	
ATP1A3	Atp1a3	4.13E-05	0.063651	2.016082	478	
ATP6V0D2	Atp6v0d2	7.24E-06	0.063651	3.389887	245972	
C3	C3	3.12E-05	0.063651	4.104003	718	
CCL23	Ccl6	1.84E-05	0.063651	2.385635	6368	
CCL26	Ccl8	1.39E-05	0.063651	3.429348	10344	
CCL5	Ccl5	3.65E-05	0.063651	2.877596	6352	
CFP	Cfp	6.86E-05	0.063651	2.432155	5199	
CSF2RB	Csf2rb2	2.64E-05	0.063651	2.922136	1439	
FOLR2	Folr2	5.77E-05	0.063651	2.144152	2350	
IL1RN	Il1rn	7.66E-05	0.063651	3.278842	3557	
ITGB1BP2	Irgb1bp2	5.70E-05	0.063651	1.929512	26548	
LGALS3	Lgals3	1.77E-05	0.063651	2.459773	3958	
MAFB	Mafb	6.61E-05	0.063651	2.270724	9935	
MFHAS1	Mfhas1	2.88E-05	0.063651	2.084097	9258	
MMP19	Mmp19	2.30E-05	0.063651	2.293279	4327	
NFATC1	Nfatc1	5.33E-05	0.063651	1.840636	4772	
NOS2	Nos2	1.66E-05	0.063651	4.900492	4843	
PTAFR	Ptafr	7.72E-05	0.063651	3.035468	5724	
SH2B2	Sh2b2	1.00E-05	0.063651	2.141262	10603	
TARM1	Tarm1	5.05E-05	0.063651	3.125284	441864	
TFEC	Tfec	6.46E-05	0.063651	1.900903	22797	
TREM2	Trem2	7.53E-05	0.063651	2.849572	54209	
XDH	Xdh	7.03E-05	0.063651	2.342691	7498	
CLEC6A	Clec4n	9.85E-05	0.075831	4.011309	93978	
LILRB4	Gm15922	9.48E-05	0.075831	2.599271	11006	
CCL11	Ccl11	0.000114	0.078015	2.558672	6356	
FLRT2	Flrt2	0.000107	0.078015	1.907153	23768	
C1orf162	I830077J02Rik	0.000152	0.078749	1.318517	128346	
CCL23	Ccl9	0.00012	0.078749	2.580178	6368	
CD52	Cd52	0.00014	0.078749	2.221747	1043	
CLEC4D	Clec4d	0.000164	0.078749	5.18802	338339	
CYBA	Cyba	0.000153	0.078749	1.69344	1535	
FMNL1	Fmnl1	0.000161	0.078749	1.739132	752	
GLRX	Glrx	0.000151	0.078749	1.850243	2745	
GSTK1	Gstk1	0.00015	0.078749	1.505521	373156	
IL21R	Il21r	0.000154	0.078749	2.60111	50615	
MARCO	Marco	0.000128	0.078749	2.569466	8685	
PI16	Pi16	0.000139	0.078749	2.568487	221476	
S1PR4	S1pr4	0.000165	0.078749	1.686857	8698	
CD3D	Cd3d	0.000177	0.079788	2.277825	915	
FGR	Fgr	0.000181	0.080455	2.310856	2268	
CLEC10A	Mgl2	0.000196	0.082172	1.335802	10462	
IGSF6	Igsf6	0.000199	0.082172	2.37001	10261	
SLAMF6	Slamf6	0.000199	0.082172	2.697009	114836	
C1QC	C1qc	0.000203	0.082475	1.959845	714	
CLEC4E	Clec4e	0.00021	0.083096	4.952239	26253	
SLC7A2	Slc7a2	0.00021	0.083096	2.784737	6542	
LILRA4	Pirb	0.000218	0.085055	2.728993	23547	
B4GALNT1	B4galnt1	0.00023	0.088485	1.895877	2583	
CD84	Cd84	0.000238	0.090502	2.365296	8832	
FERMT3	Fermt3	0.000264	0.094703	1.919157	83706	
CSTB	Cstb	0.000305	0.096298	1.203798	1476	
NFAM1	Nfam1	0.000287	0.096298	2.014159	150372	
SFRP4	Sfrp4	0.000291	0.096298	2.659627	6424	
THY1	Thy1	0.000307	0.096298	2.500864	7070	
TNFAIP2	Tnfaip2	0.000307	0.096298	2.069053	7127	
COX6A2	Cox6a2	0.000314	0.097269	2.151099	1339	
CTSD	Ctsd	0.000323	0.097269	1.187588	1509	
IL2RG	Il2rg	0.000324	0.097269	2.877191	3561	
CACNB1	Cacnb1	0.000359	0.098224	2.259612	782	
CD300LF	Cd300ld	0.000335	0.098224	2.398153	146722	
DPEP2	Dpep2	0.000375	0.098224	1.954277	64174	
EMX2	Emx2	0.000365	0.098224	1.107609	2018	
IL7R	Il7r	0.000341	0.098224	3.09894	3575	
LPAR1	Lpar1	0.000353	0.098224	2.026258	1902	
PLA2G7	Pla2g7	0.00035	0.098224	4.154612	7941	
TNFRSF1B	Tnfrsf1b	0.000345	0.098224	2.779606	7133	
WAS	Was	0.000337	0.098224	1.537643	7454	
CD7	Cd7	0.000379	0.098422	2.085454	924	

Table S1: The significantly differentially expressed genes (N = 73) that are increased upon CMP treatment as compared to isotype control (logFC > 0) evaluated in Figure 6A and as described in the Methods section.