SUPPLEMENTARY MATERIALS

Upregulation of MHC I Antigen Processing Machinery Gene Expression in Breast Cancer Cells by Trichostatin A

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Table S1. Primer sequences for real-time PCR

Gene	Accession Number	Forward Primer Sequence (5'-3')	Reverse Primer Sequence (5'-3')
PSMB9	[1]	ATGCTGACTCGACAGCCTTT	GCAATAGCGTCTGTGGTGAA
PSMB8	NM_148919.3	TGCTGTCCAACATGATGTGC	AGTCCCATGTTCATCCACGTAG
PSMB10	NM_002801.4	GTGAAGCCACTAACCCTGGA	CTGGGTTTATTCCCCCTTGT
TAP1	NM_001292022.1	AGTTCGAAGCTTTGCCAACG	AGAGGATTCCCACTTTCAGCAG
TAP2	NM_001290043.1	TTTTGACCCCCATGCCTTTG	TCCGCAAGTTGATTCGAGAC
B2M	NM_004048.4	TCGCGCTACTCTCTCTTTCTG	ATTCTCTGCTGGATGACGTGAG
MHC I	[1]	GCGGCTACTACAACCAGAGC	GATGTAATCCTTGCCGTCGT
CD274	[2]	GTGGCATCCAAGATACAAACTCAA	TCCTTCCTCTTGTCACGCTCA
nNav1.5	NM_001160160.2	TGATTATCATGGCGTATGTATCAGA	TGAGGGCAAAGACGCTGAG
MMP1	[3]	ACGAATTTGCCGACAGAGAT	GGAAGCCAAAGGAGCTGTAG
FN1	[4]	GACCTATCCAAGCTCAAGTGGT	ATCCAAGGTTTCTGGGTGGG

β-actin	[5]	ATTGCCGACAGGATGCAGAAG	AGAAGCATTTGCGGTGGACG

Table S2. Effects of TSA treatment on MCF-7 spheroid culture against the monolayer culture

Gene	**Average Fold Change of Treatment Effects**
PSMB9	4.35
PSMB8	19.38
PSMB10	1.26
TAP1	13.33
TAP2	107.54
B2M	2.33
MHC I	7.68
CD274	21.9
nNav1.5	3.8
MMP1	18.25
FN1	–4.43

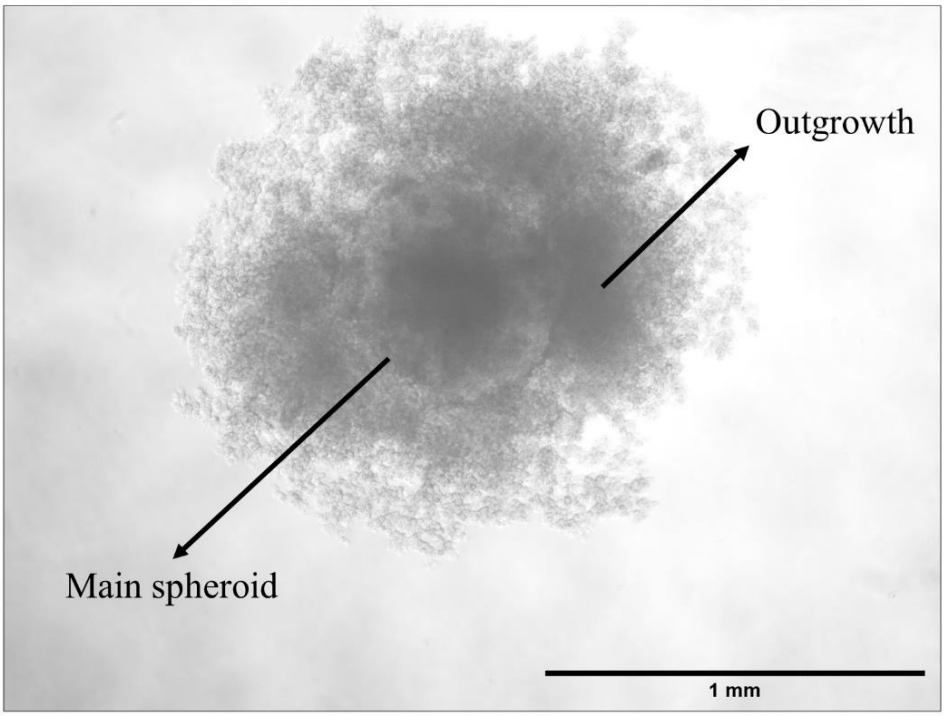

Fig. S1. The definition of main spheroid and outgrowth. The perimeter is utilized for volume analysis. Black error bars represent 1mm.

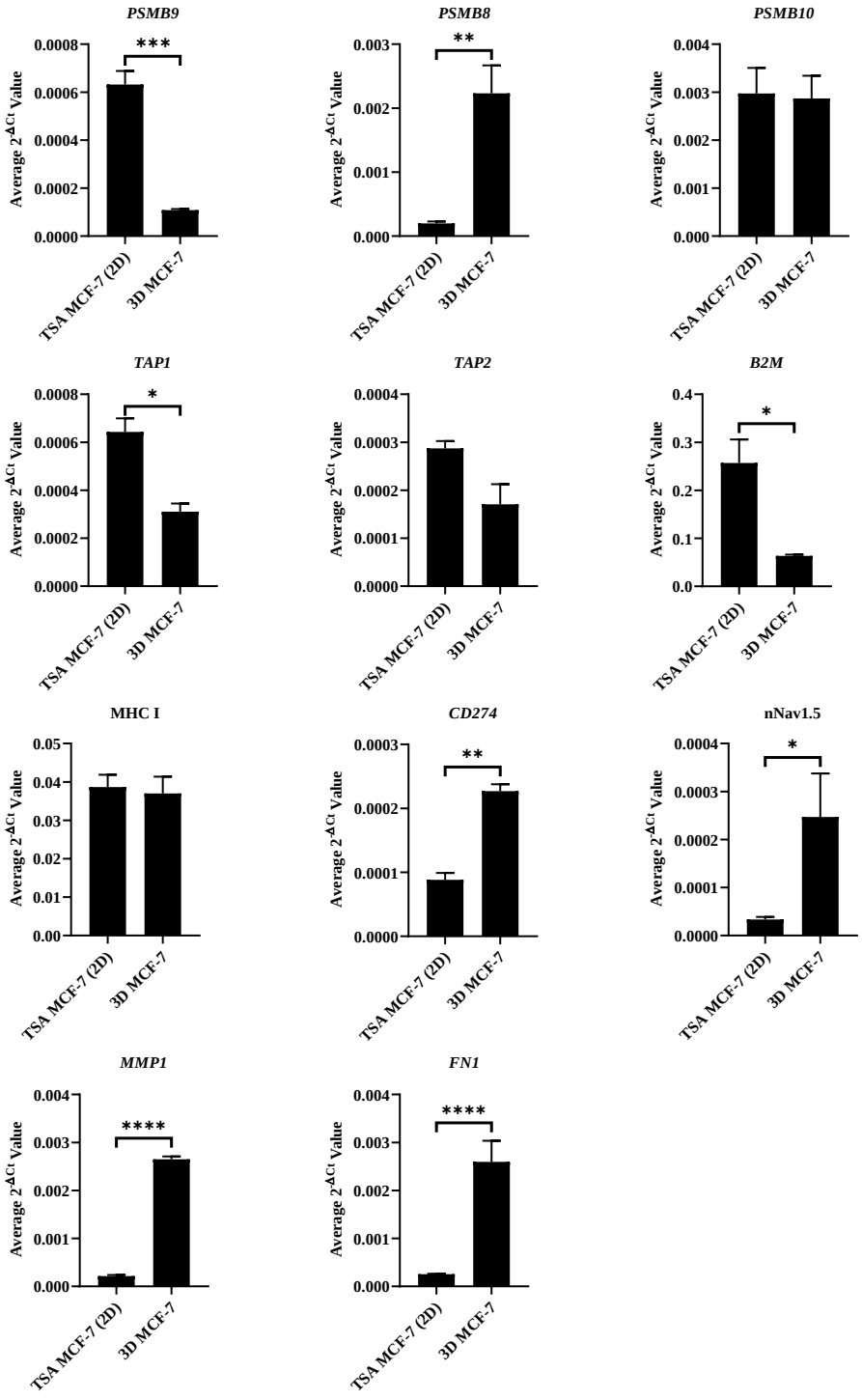

Fig. S2. mRNA expression comparison of MHC I APM, *CD274* and metastatic markers in 1000 ng/mL TSA-treated monolayer MCF-7 compared to the untreated MCF-7 spheroids. The untreated MCF-7 spheroids were developed via the liquid overlay method and grown for 15 days. Analysis was done using the $2^{-\Delta Ct}$ method.

REFERENCES (TABLE S1)

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