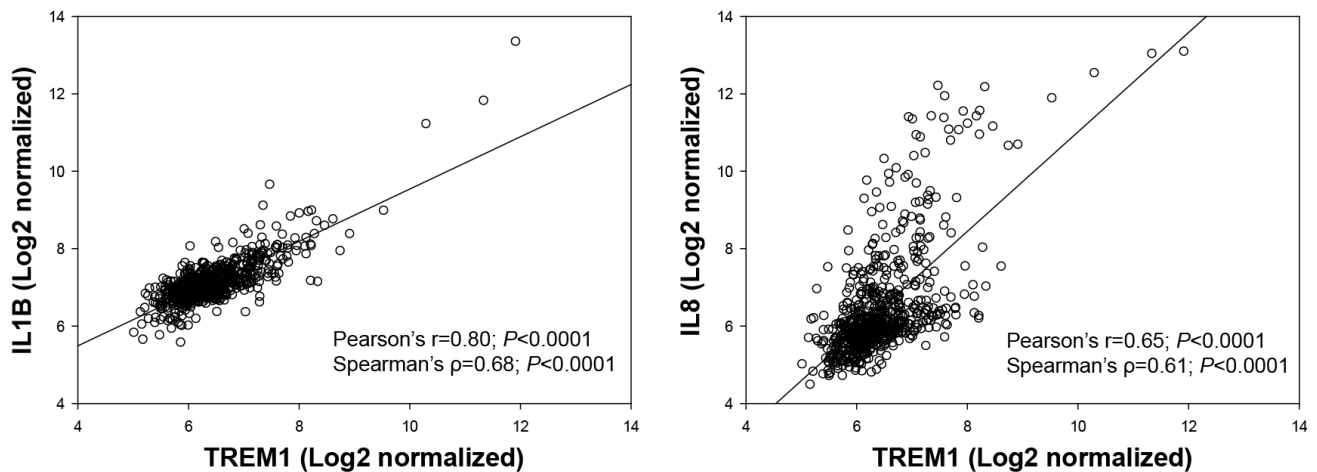


Supplementary Material

Supplementary Figures 1 – 7

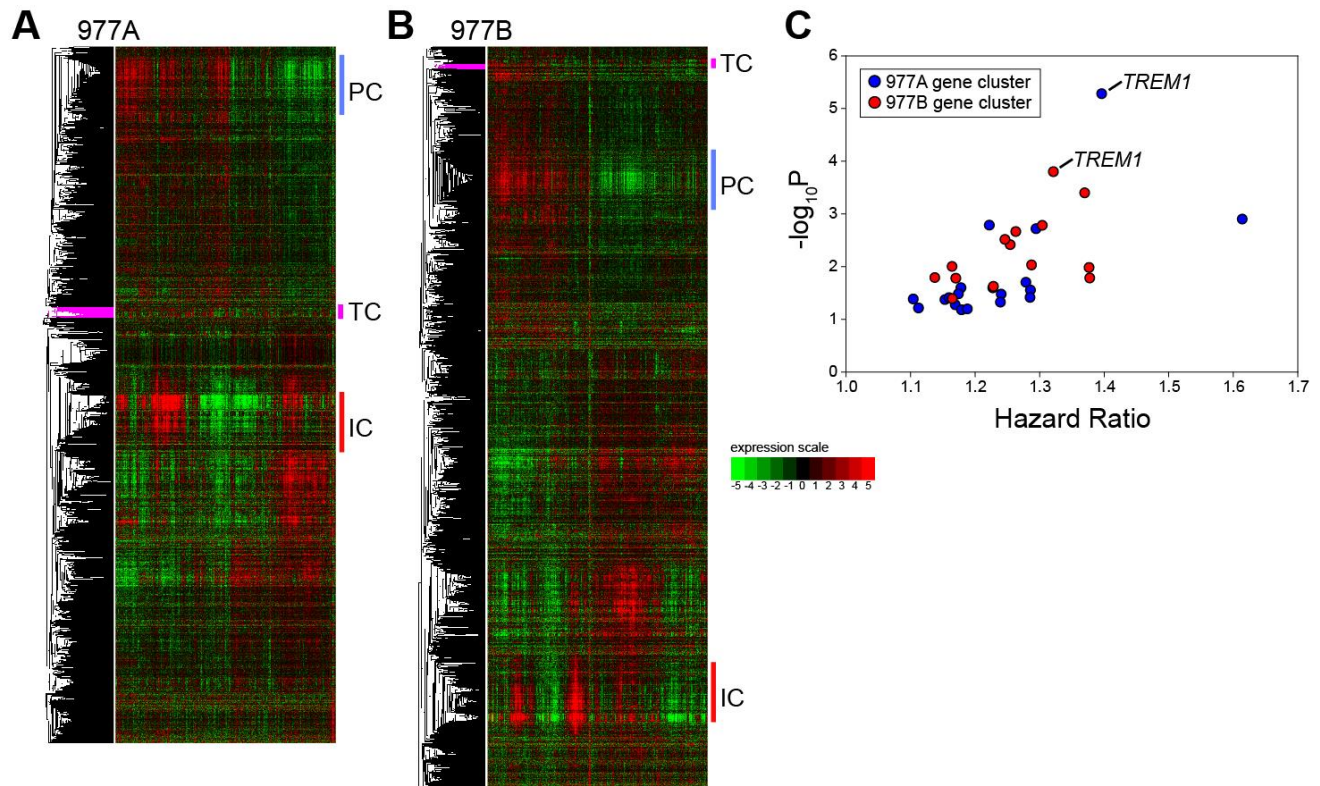
Supplementary Table 1

Supplementary Figure 1



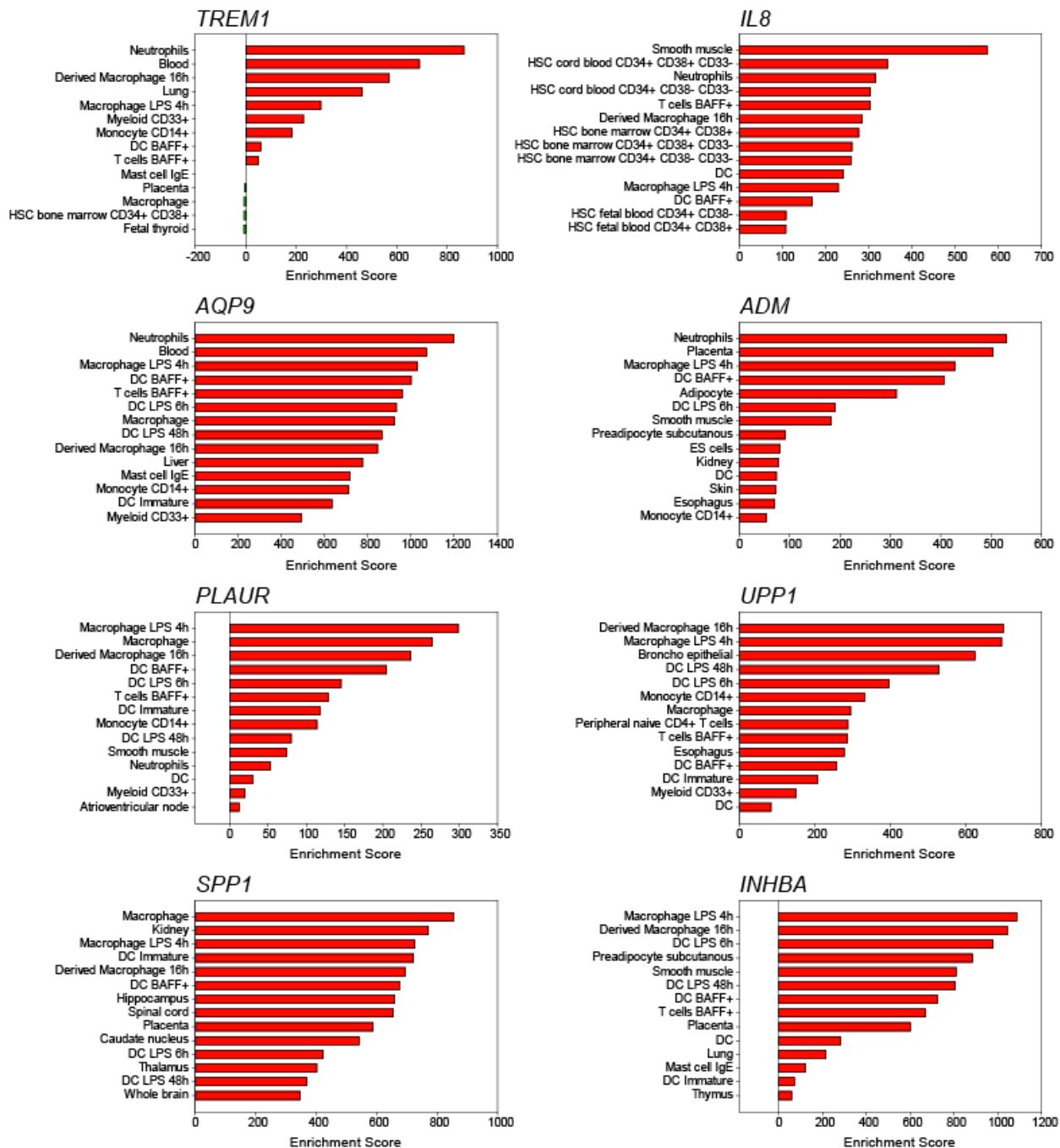
Supplemental Figure 1. *TREM1* expression correlates significantly with expression of *IL1B* (left panel) and *IL8/CXCL8* (right panel) in the MDACC-701 dataset. Pearson's and Spearman correlation coefficients and p-values are shown.

Supplementary Figure 2



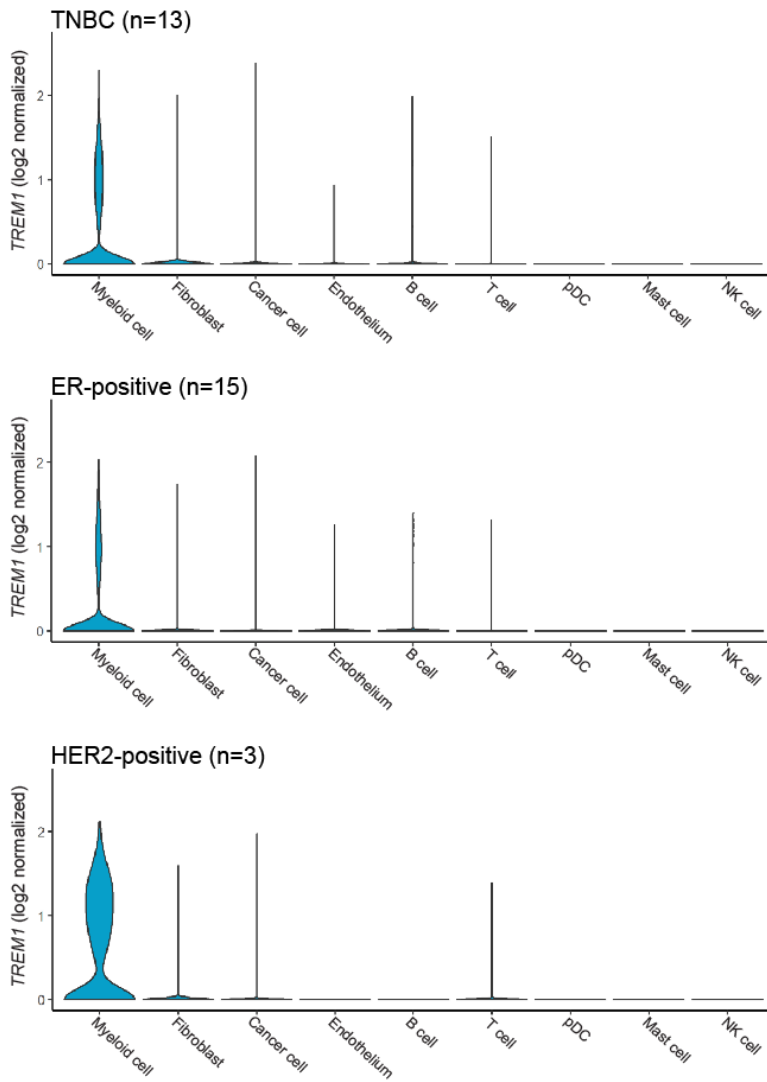
Supplemental Figure 2. Discovery of a *TREM1*-associated gene cluster associated with poor DMFS. Breast tumor expression profiles of the MC1 meta-cohort were randomized to two subgroups, comprising 977 tumors in each, for testing and validation purposes (977A and 977B, respectively). Within each subgroup, genes with significant associations with DMFS by Cox regression ($q < 0.1$, Benjamini-Hochberg) were selected for hierarchical clustering (977A, $n = 3,094$ genes; 977B, $n = 3,304$ genes). Genes corresponding to the proliferation gene cluster (PC, blue bar), immune gene clusters (IC, red bar) and the *TREM1*-associated gene cluster (TC, pink bar) are shown to the right of the heat maps for (A) 977A and (B) 977B. Tumors are represented by columns (dendrograms omitted); genes are represented by rows. Red indicates above-mean expression; green indicates below-mean expression. (C) Genes within the *TREM1*-associated gene clusters having average Pearson correlation of 0.28 in 977A (blue circles, $n = 19$ genes) and 977B (red circles, $n = 14$ genes) were compared by plotting Cox regression hazard ratios (X-axis) and false discovery rate-adjusted, negative \log_{10} p-values (Y-axis).

Supplementary Figure 3



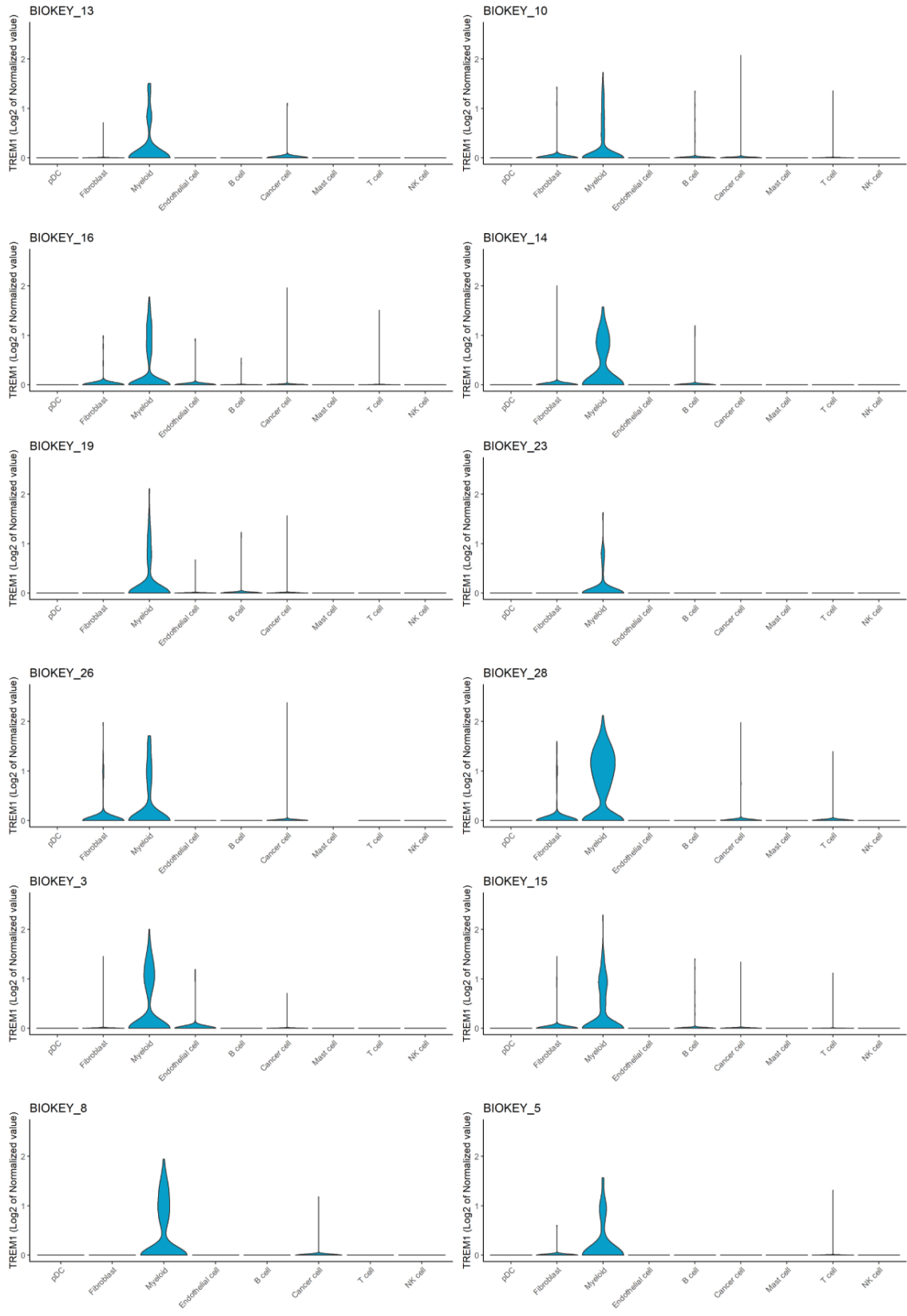
Supplemental Figure 3. *TREM1* and co-expressed genes in breast tumors exhibit enriched expression in cells of the myeloid lineage. Tissue-specific enrichment scores were generated according to methods described by Benita et. al. (Benita et al., 2010) for >14,000 genes comprising the Affymetrix U133A GeneChip and 126 human cell subsets and tissues. For each gene, the top 14 most enriched cell subsets and tissues are shown.

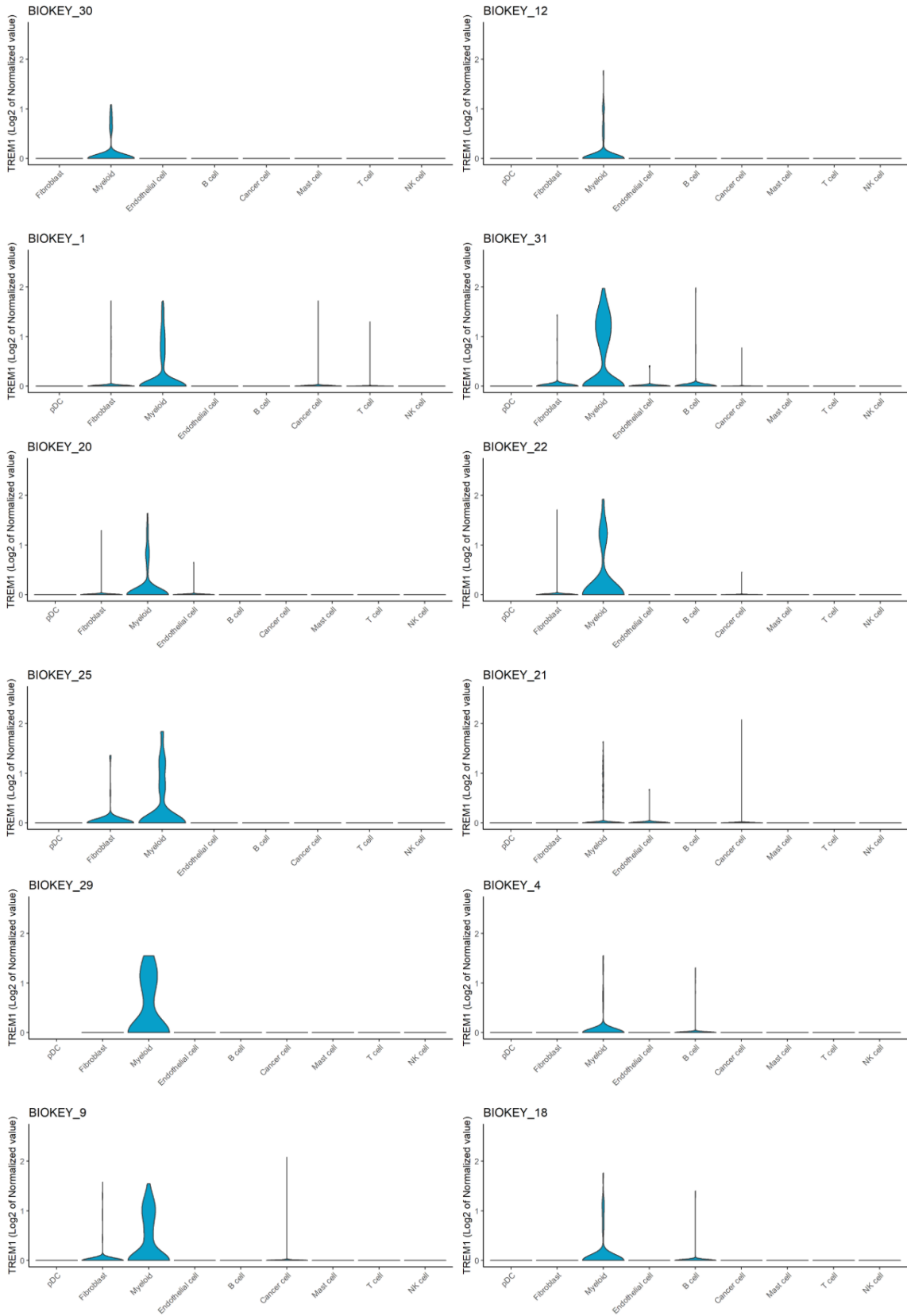
Supplementary Figure 4

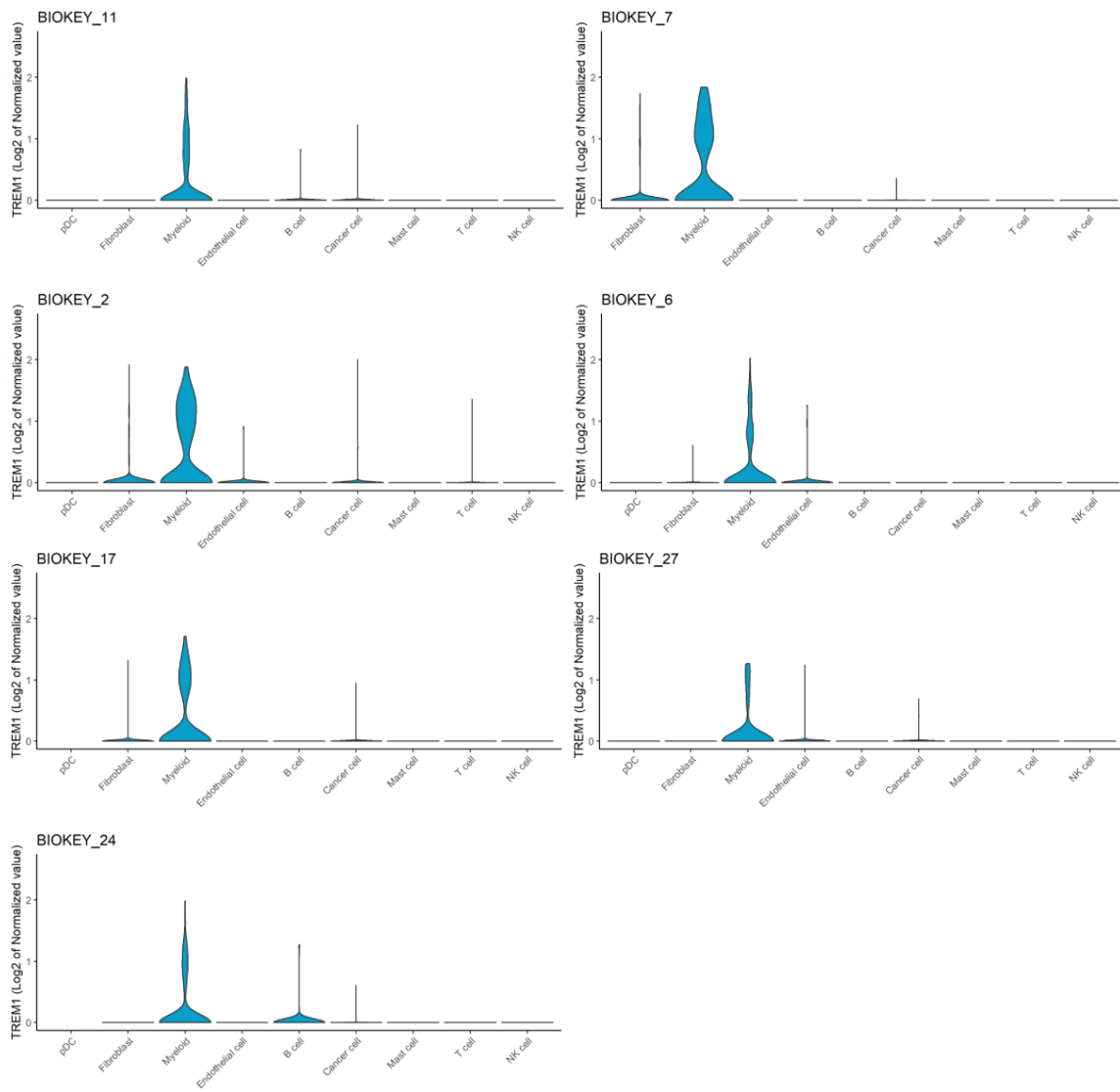


Supplemental Figure 4. *TREM1* expression in cellular subsets of the Bassez et. al. scRNAseq dataset grouped by breast cancer subtype.

Supplementary Figure 5

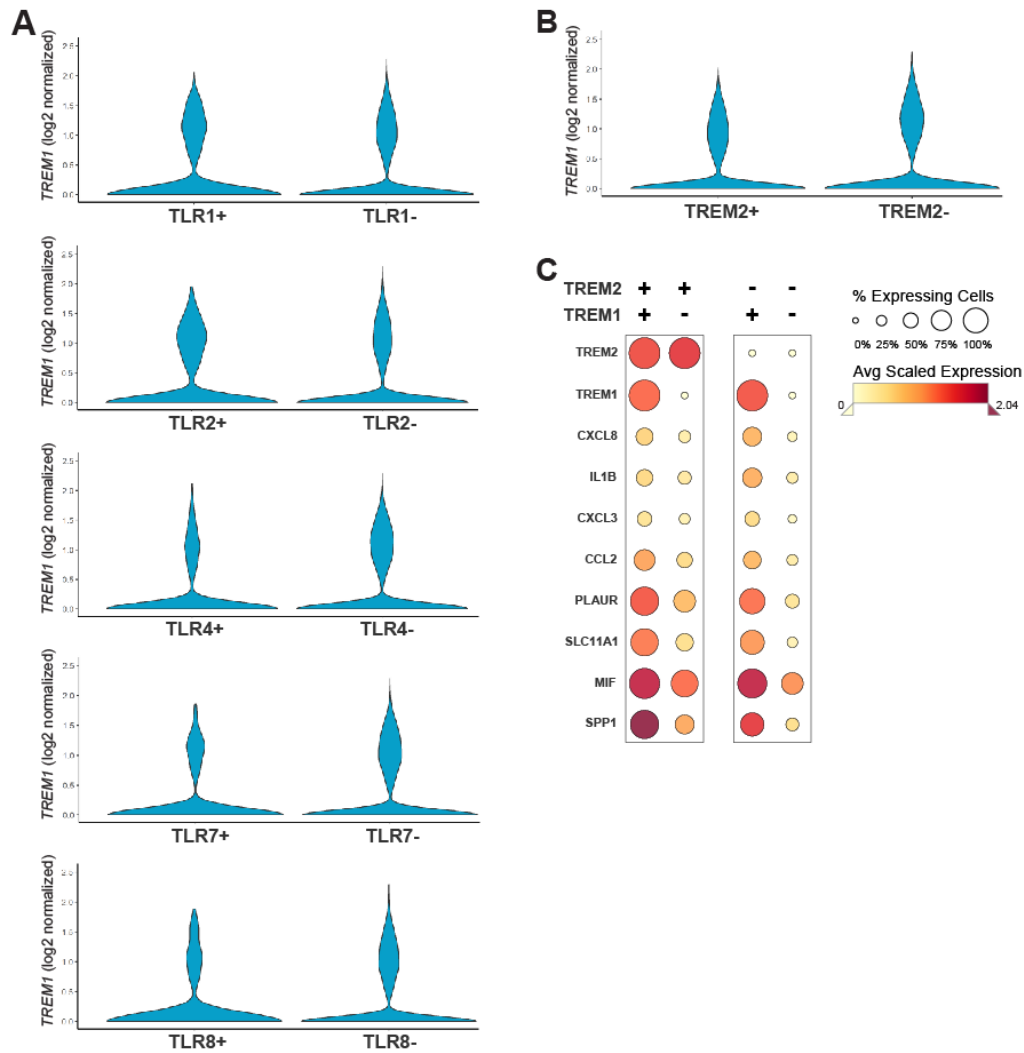






Supplemental Figure 5. *TREM1* expression in cellular subsets of the Bassez et. al. dataset grouped by individual breast cancer patients (n=31).

Supplementary Figure 7



Supplemental Figure 7. Analysis of *TREM1* myeloid expression and expression of myeloid receptors related to TREM-1 activation and function. In the myeloid compartment of the Bassez et. al. dataset, (A) *TREM1* expression distributions are shown compared between *TLR*+ and *TLR*- myeloid cell populations, and (B) *TREM2*+ and *TREM2*- myeloid cells. (C) *TREM1* target gene frequencies of expression (% expressing cells) and average expression levels (heatmap) are shown compared between *TREM2*+ and *TREM2*- myeloid cells.

Supplemental Table 1. Genes and corresponding Z values identified by logistic regression as having statistical negative interactions with immune-associated neoadjuvant chemo response in the MDACC-701 dataset.

Probest ID	Gene Symbol	Binary Z	Neutrophil Chemotaxis	Probest ID	Gene Symbol	Binary Z	Neutrophil Chemotaxis
202155_s_at	NUP214	-4.1		212777_at	SOS1	-2.3	
217336_at	RPS10	-3.9		207540_s_at	SYK	-2.3	X
218955_at	BRF2	-3.4		78495_at	ZNF783	-2.3	
44822_s_at	MIER2	-3.3		219623_at	ACTR5	-2.2	
37254_at	ZNF133	-3.3		219437_s_at	ANKRD11	-2.2	
219434_at	TREM1	-3.3	X	219521_at	B3GAT1	-2.2	
207205_at	CEACAM4	-3.1		210708_x_at	CASP10	-2.2	
209495_at	CEP250	-3.1		206728_at	ECE2	-2.2	
202573_at	CSNK1G2	-3.1		205321_at	EIF2S3	-2.2	
204594_s_at	MIEF1	-3.1		204817_at	ESPL1	-2.2	
204828_at	RAD9A	-3.1		205650_s_at	FGA	-2.2	
206019_at	RBM19	-3.1		220819_at	FRMD1	-2.2	
216036_x_at	WDC1	-3.1		204973_at	GJB1	-2.2	
209797_at	CNPY2	-3.0		211977_at	GPR107	-2.2	
32032_at	ESS2	-3.0		221287_at	GPRC5D	-2.2	
221983_at	RETREG2	-3.0		209558_s_at	HIP1R	-2.2	
216505_x_at	RPS10P5	-3.0		219281_at	MSRA	-2.2	
201320_at	SMARCC2	-3.0		217619_x_at	[unknown]	-2.2	
214783_s_at	ANXA11	-2.9		216551_x_at	PLCG1	-2.2	
205654_at	C4BPA	-2.9		201896_s_at	PSRC1	-2.2	
215568_x_at	[unknown]	-2.9		205228_at	RBMS2	-2.2	
203185_at	RASSF2	-2.9		211735_x_at	SFTPC	-2.2	
218141_at	UBE2O	-2.9		205592_at	SLC4A1	-2.2	
220730_at	ZNF778	-2.9		216271_x_at	SYDE1	-2.2	
218274_s_at	ANKZF1	-2.8		213107_at	TNIK	-2.2	
218457_s_at	DNMT3A	-2.8		218020_s_at	ZFAND3	-2.2	
214185_at	KHDRBS1	-2.8		205437_at	ZNF211	-2.2	
212844_at	RRP1B	-2.8		78330_at	ZNF335	-2.2	
41657_at	STK11	-2.8		220501_at	ACTL7A	-2.1	
217733_s_at	TMSB10	-2.8		90265_at	ADAP1	-2.1	
216409_at	ACSL6	-2.7		206105_at	AFB2	-2.1	
218950_at	ARAP3	-2.7		202206_at	ARL4C	-2.1	
219261_at	C7orf26	-2.7		212645_x_at	BRE	-2.1	
221357_at	CHRM4	-2.7		221764_at	C19orf22	-2.1	
202500_at	DNAJB2	-2.7		206327_s_at	CDH15	-2.1	
210736_x_at	DTNA	-2.7		219640_at	CLDN15	-2.1	
222113_s_at	EPS15L1	-2.7		206970_at	CNTN2	-2.1	
206885_x_at	GH1	-2.7		208159_x_at	DDX11	-2.1	
210079_x_at	KCNAB1	-2.7		208250_s_at	DMBT1	-2.1	
221792_at	RAB6A	-2.7		221812_at	FBXO42	-2.1	
37462_i_at	SF3A2	-2.7		208885_at	LCP1	-2.1	
221939_at	CARM1	-2.6		205036_at	LSM6	-2.1	
207556_s_at	DGKZ	-2.6		43544_at	MED16	-2.1	
218660_at	DYSF	-2.6		211599_x_at	MET	-2.1	
213242_x_at	KIAA0284	-2.6		215292_s_at	MKL1	-2.1	
213281_at	LOC100288387	-2.6		216554_s_at	[unknown]	-2.1	
35147_at	MCF2L	-2.6		217390_x_at	[unknown]	-2.1	
215484_at	[unknown]	-2.6		208360_s_at	[unknown]	-2.1	
205656_at	PCDH17	-2.6		209062_x_at	NCOA3	-2.1	
219459_at	POLR3B	-2.6		206402_s_at	NPFF	-2.1	
200869_at	RPL18A	-2.6		205261_at	PGC	-2.1	
207355_at	SLC1A7	-2.6		204613_at	PLCG2	-2.1	
214886_at	ZNF266	-2.6		208286_x_at	POUSF1	-2.1	
210662_s_at	ZNF589	-2.6		204044_at	QPRT	-2.1	
214322_at	CAMK2G	-2.5		216383_at	RPL18A	-2.1	
220015_at	CASZ1	-2.5		214370_at	S100A8	-2.1	X
214627_at	EPX	-2.5		203535_at	S100A9	-2.1	X
210981_s_at	GRK6	-2.5		209381_x_at	SF3A2	-2.1	
216309_x_at	JRK	-2.5		215926_x_at	SNAPC4	-2.1	
210058_at	MAPK13	-2.5		52169_at	STRADA	-2.1	
202463_s_at	MBD3	-2.5		203046_s_at	TIMELESS	-2.1	
221379_at	[unknown]	-2.5		215540_at	TRAC	-2.1	
219039_at	SEMA4C	-2.5		211432_s_at	TYRO3	-2.1	
218596_at	TBC1D13	-2.5		65521_at	UBE2D4	-2.1	
201768_s_at	CLINT1	-2.4		211682_x_at	UGT2B28	-2.1	
220522_at	CRB1	-2.4		212156_at	VPS39	-2.1	
214030_at	CRYBG3	-2.4		203892_at	WFDC2	-2.1	
219914_at	ECEL1	-2.4		207708_at	ALOXE3	-2.0	
221539_at	EIF4EBP1	-2.4		219141_s_at	AMBRA1	-2.0	
210043_at	FRMD8	-2.4		221077_at	ARMC4	-2.0	
214626_s_at	GANAB	-2.4		220659_s_at	C7orf43	-2.0	
220165_at	INO80D	-2.4		207861_at	CCL22	-2.0	X
49329_at	KLHL22	-2.4		213523_at	CCNE1	-2.0	
207908_at	KRT2	-2.4		216293_at	CLTA	-2.0	
210252_s_at	MADD	-2.4		214076_at	GFOD2	-2.0	
207473_at	MLN	-2.4		213180_s_at	GOSR2	-2.0	
215200_x_at	[unknown]	-2.4		204949_at	ICAM3	-2.0	
202924_s_at	PLAGL2	-2.4		213450_s_at	ICOSLG	-2.0	
220894_x_at	PRDM12	-2.4		205067_at	IL1B	-2.0	X
220758_s_at	ROBO4	-2.4		202859_x_at	IL8	-2.0	X
209980_s_at	SHMT1	-2.4		211806_s_at	KCNJ15	-2.0	
214096_s_at	SHMT2	-2.4		207339_s_at	LTB	-2.0	
209777_s_at	SLC19A1	-2.4		206571_s_at	MAP4K4	-2.0	
211123_at	SLC5A5	-2.4		210059_s_at	MAPK13	-2.0	
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219442_at	C16orf67	-2.3		215279_at	[unknown]	-2.0	
219161_s_at	CKLF	-2.3		217088_s_at	NCR1	-2.0	
204971_at	CSTA	-2.3		221526_x_at	PARD3	-2.0	
2028_s_at	E2F1	-2.3		215031_x_at	RNF126	-2.0	
206429_at	F2RL1	-2.3		201909_at	RPS4Y1	-2.0	
60528_at	JMJD7	-2.3		202917_s_at	S100A8	-2.0	X
210635_s_at	KLHL20	-2.3		221701_s_at	STRA6	-2.0	
37796_at	LRCH4	-2.3		216919_at	TP53I11	-2.0	
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				214795_at	ZMYND8	-2.0	