Supplementary Table I. Genes whose low expression was associated with increased risk of recurrence and death in METABRIC TNBC patients.

						HR of low t	to high express	sion					
Gene	HR for death	95% CI of HR	p-value	HR for recurrence	95% Cl of HR	p-value	Gene	HR for death	95% Cl of HR	p-value	HR for recurrence	95% Cl of HR	p-value
RASSF5	3.024	1.951-4.796	<0.0001	2.652	1.747-4.099	<0.0001	PTPRO	2.167	1.412-3.385	0.0004	2.158	1.428-3.308	0.0002
DOCK2	2.853	1.852-4.466	<0.0001	2.428	1.609-3.706	<0.0001	CD52	2.167	1.426-3.323	0.0003	2.081	1.390-3.137	0.0004
HLA-DMB	2.618	1.718-4.034	<0.0001	2.667	1.773-4.056	<0.0001	HLA-DQA1	2.164	1.423-3.323	0.0003	2.115	1.413-3.190	0.0003
CLEC12A	2.612	1.665-4.229	<0.0001	2.309	1.507-3.628	<0.0001	LAT2	2.162	1.418-3.341	0.0003	1.952	1.301-2.958	0.0012
IFNG	2.588	1.640-4.223	<0.0001	2.448	1.582-3.899	<0.0001	ZNF831	2.148	1.388-3.394	0.0005	2.092	1.373-3.254	0.0005
TRAT1	2.576	1.667-4.064	<0.0001	2.542	1.669-3.943	<0.0001	PSMB10	2.145	1.401-3.345	0.0004	2.105	1.396-3.225	0.0003
CD3D	2.563	1.678-3.950	<0.0001	2.243	1.492-3.393	0.0001	HLA-DRA	2.139	1.406-3.292	0.0004	2.197	1.464-3.331	0.0001
STAT4	2.495	1.620-3.902	<0.0001	2.148	1.422-3.281	0.0003	IL10RA	2.138	1.401-3.306	0.0004	2.047	1.363-3.113	0.0005
S1PR4	2.416	1.565-3.801	<0.0001	2.421	1.591-3.752	<0.0001	LAPTM5	2.127	1.394-3.293	0.0004	1.954	1.304-2.965	0.0011
PREX1	2.408	1.567-3.768	<0.0001	2.214	1.467-3.389	0.0001	PRAMEF17	2.124	1.396-3.269	0.0004	1.910	1.278-2.879	0.0016
ZBED2	2.402	1.553-3.799	<0.0001	2.320	1.525-3.602	<0.0001	P2RY13	2.116	1.380-3.303	0.0005	1.899	1.261-2.901	0.0020
HLA-DMA	2.398	1.575-3.693	<0.0001	2.667	1.773-4.056	<0.0001	IL2RB	2.106	1.376-3.269	0.0006	2.024	1.345-3.085	0.0007
CD53	2.396	1.564-3.730	<0.0001	2.387	1.582-3.655	<0.0001	CXCL9	2.095	1.375-3.225	0.0006	2.075	1.380-3.149	0.0004
GBP5	2.389	1.558-3.728	<0.0001		1.539-3.571	<0.0001	VCAM1	2.087	1.342-3.334	0.0009		1.372-3.276	0.0005
CD3G	2.361	1.544-3.670	<0.0001		1.453-3.333	0.0001	TBXAS1	2.075	1.358-3.217	0.0007	1.969	1.311-2.991	0.0010
CTSS	2.355	1.540-3.660	<0.0001		1.419-3.243	0.0002	CYSLTR1	2.060	1.339-3.237	0.0009		1.277-2.965	0.0017
CXCR6	2.316	1.502-3.647	0.0001	2.211	1.456-3.419	0.0002	AKNA	2.048	1.340-3.182	0.0009		1.290-2.977	0.0013
KIR2DL3	2.313	1.473-3.746	0.0002	2.214	1.438-3.501	0.0002	HLA-DOA	2.025	1.329-3.124	0.0010		1.249-2.831	0.0023
CD226	2.306	1.491-3.650	0.0001	2.154	1.420-3.327	0.0003	CD300LF	2.010	1.303-3.166	0.0014		1.258-2.942	0.0022
KIR3DL3	2.303	1.506-3.567	0.0001	2.279	1.512-3.509	<0.0001	PARVG	2.006	1.317-3.093	0.0012		1.265-2.871	0.0019
LPXN	2.294	1.495-3.577	0.0001	2.084	1.382-3.185	0.0004	SPCS3	1.941	1.275-2.992	0.0019		1.175-2.641	0.0059
DOCK10	2.293	1.499-3.559	0.0001	2.288	1.516-3.497	<0.0001	SLC52A1	1.940	1.264-3.036	0.0023		1.308-3.032	0.0011
SLCO2B1	2.285	1.496-3.540	0.0001	2.117	1.411-3.210	0.0003	GZMH	1.929	1.256-3.022	0.0025		1.262-2.931	0.0020
SLAMF7	2.285	1.482-3.597	0.0002	2.062	1.363-3.173	0.0005	PITPNM2	1.928	1.259-3.000	0.0024		1.179-2.701	0.0058
ARHGAP30	2.279	1.486-3.557	0.0001	2.315	1.531-3.559	<0.0001	KLRD1	1.928	1.263-2.986	0.0022		1.281-2.934	0.0015
APOBEC3G		1.484-3.562	0.0001	2.319	1.532-3.577	<0.0001	C8orf31	1.919	1.263-2.949	0.0022		1.179-2.660	0.0016
KIR3DL1	2.275	1.461-3.629	0.0001	2.404	1.562-3.790	<0.0001	PLCL2	1.887	1.238-2.914	0.0022		1.260-2.888	0.0030
CD6	2.274	1.487-3.524	0.0002	2.122	1.409-3.233	0.0003	CCDC28A	1.883	1.226-2.937	0.0030	1.993	1.318-3.059	0.0021
SASH3	2.260	1.476-3.518	0.0001	2.160	1.432-3.307	0.0003	WNT2	1.879	1.230-2.917	0.0037	1.847	1.231-2.808	0.0010
PVRIG	2.255	1.461-3.531	0.0002	2.100	1.391-3.240	0.0002	AGAP2	1.861	1.222-2.866	0.0034	1.868	1.247-2.824	0.0023
PCED1B	2.252	1.464-3.506	0.0002	2.061	1.364-3.145	0.0004	INSRR	1.854		0.0037			0.0024
									1.216-2.866			1.112-2.518	
PRKCB PLEK	2.251 2.251	1.466-3.512 1.472-3.485	0.0002 0.0002	2.129	1.430-3.261	0.0003	RCBTB2 PRKCH	1.853 1.850	1.205-2.900	0.0048	1.715 1.764	1.137-2.618	0.0099
	2.251			2.186	1.450-3.333 1.405-3.298	0.0002			1.213-2.861			1.175-2.683	0.0036
SEL1L		1.453-3.541	0.0003	2.137			ADARB2-AS		1.205-2.877	0.0046		1.217-2.803	
HCLS1	2.247	1.473-3.473	0.0002	2.065	1.377-3.130	0.0004	PPM1M	1.801	1.184-2.772	0.0058		1.140-2.578	0.0093
VAV1	2.245	1.457-3.534	0.0002	2.276	1.498-3.529	<0.0001	PLBD1	1.788	1.171-2.767	0.0070		1.292-2.965	0.0014
ITK	2.239	1.458-3.499	0.0002	2.157	1.425-3.317	0.0002	REM1	1.784	1.166-2.773	0.0073		1.204-2.765	0.0042
GZMA	2.232	1.456-3.477	0.0002	2.183	1.444-3.348	0.0002	RFTN1	1.776	1.163-2.751	0.0077	1.755	1.168-2.672	0.0066
EVI2B	2.218	1.450-3.434	0.0002	2.033	1.350-3.088	0.0007	KCNE1	1.774	1.165-2.724	0.0074		1.206-2.725	0.0041
CD83	2.214	1.441-3.467	0.0002	2.079	1.380-3.183	0.0004	LCP1	1.762	1.163-2.683	0.0076		1.200-2.689	0.0044
DEF6	2.191	1.431-3.403	0.0003	1.963	1.304-2.990	0.0012	TPK1	1.749	1.150-2.692	0.0088		1.158-2.628	0.0074
BTN3A2	2.181	1.422-3.404	0.0003	2.077	1.379-3.171	0.0004	RASGEF1A	1.734	1.136-2.683	0.0106		1.308-3.014	0.0011
IGFALS	2.175	1.426-3.362	0.0003	1.976	1.318-2.990	0.0009	HHEX	1.733	1.134-2.677	0.0109		1.156-2.642	0.0078
FLT3LG	2.172	1.417-3.382	0.0003	1.975	1.312-3.010	0.0010	FBXO47	1.689	1.096-2.648	0.0171	1.671	1.102-2.579	0.0154

HR, hazard ratio; Cl. confidence interval. Hazard ratio was adjusted for age at diagnosis (continuous) and NPI (continuous). The corresponding 95% Cl and p-value was based on likelihood ratio test. Sample size n = 233. Genes were listed in descending order of HR values for death.

Supplementary Table II. Genes whose high expression was associated with increased risk of recurrence and death in METABRIC TNBC patients.

HR of high to low expression										
Gene	HR for death	95% Cl of HR	p-value	HR for recurrence	95% Cl of HR	p-value				
LINC00943	2.106	1.380-3.256	0.0005	1.967	1.311-2.981	0.0010				
EXD2	2.096	1.370-3.264	0.0006	1.947	1.296-2.967	0.0013				
USP30	2.066	1.349-3.212	0.0008	1.986	1.318-3.031	0.0010				
TMEM59L	2.046	1.333-3.200	0.0009	2.002	1.330-3.062	0.0008				
SNORA65	2.044	1.339-3.164	0.0009	1.836	1.225-2.777	0.0032				
PHOX2A	2.036	1.326-3.186	0.0010	1.863	1.239-2.859	0.0027				
CAPN14	2.007	1.309-3.131	0.0013	1.919	1.273-2.938	0.0017				
NCBP2	2.006	1.313-3.104	0.0012	1.815	1.210-2.747	0.0039				
DCAF4	2.004	1.315-3.083	0.0012	1.857	1.240-2.799	0.0026				
FFAR1	1.974	1.292-3.058	0.0016	2.007	1.336-3.055	0.0008				
CYTH2	1.964	1.287-3.048	0.0017	1.896	2.264-2.882	0.0019				
LBX2	1.963	1.291-3.002	0.0016	2.012	1.343-3.032	0.0007				
DNAAF3	1.954	1.282-3.002	0.0018	1.878	1.253-2.837	0.0023				
DNAJB7	1.941	1.280-2.965	0.0018	1.848	1.239-2.774	0.0026				
FAM204A	1.920	1.255-2.991	0.0025	2.054	1.359-3.162	0.0006				
ZNF271P	1.904	1.256-2.906	0.0024	1.781	1.193-2.669	0.0048				
PTRHD1	1.894	1.244-2.920	0.0028	1.861	1.241-2.823	0.0026				
DNM3	1.889	1.246-2.890	0.0027	2.173	1.452-3.283	0.0002				
SCAF4	1.850	1.212-2.865	0.0042	1.713	1.144-2.593	0.0088				
MAP3K9	1.848	1.205-2.870	0.0048	1.941	1.287-2.960	0.0015				
DOLK	1.848	1.208-2.869	0.0045	1.839	1.221-2.807	0.0034				
TSEN34	1.825	1.197-2.816	0.0050	1.715	1.147-2.585	0.0086				
CA9	1.802	1.185-2.729	0.0062	1.702	1.136-2.539	0.0102				
NCAPG	1.793	1.170-2.793	0.0071	1.645	1.094-2.501	0.0165				
GRWD1	1.757	1.153-2.713	0.0085	1.742	1.161-2.644	0.0072				
RCAN3	1.755	1.146-2.716	0.0095	1.795	1.193-2.728	0.0049				
H2BC9	1.749	1.143-2.705	0.0099	1.876	1.245-2.855	0.0026				
TMEM190	1.734	1.135-2.687	0.0108	1.738	1.155-2.649	0.0078				
MRGBP	1.703	1.118-2.627	0.0129	1.592	1.064-2.408	0.0237				
ENO1	1.664	1.099-2.529	0.0163	1.582	1.061-2.365	0.0246				
PLIN3	1.633	1.074-2.494	0.0218	1.619	1.081-2.438	0.0195				
DONSON	1.619	1.060-2.508	0.0258	1.526	1.016-2.316	0.0417				
PDCD6IPP2	1.609	1.058-2.466	0.0262	1.651	1.103-2.489	0.0147				
PRADC1	1.605	1.044-2.494	0.0311	1.686	1.113-2.584	0.0136				
SKA1	1.601	1.045-2.486	0.0304	1.482	0.986-2.249	0.0586				

HR, hazard ratio; CI, confidence interval. Hazard ratio was adjusted for age at diagnosis (continuous) and NPI (continuous). The corresponding 95% CI and p-value was based on likelihood ratio test. Sample size n = 233. Genes were listed in descending order of HR values for death.