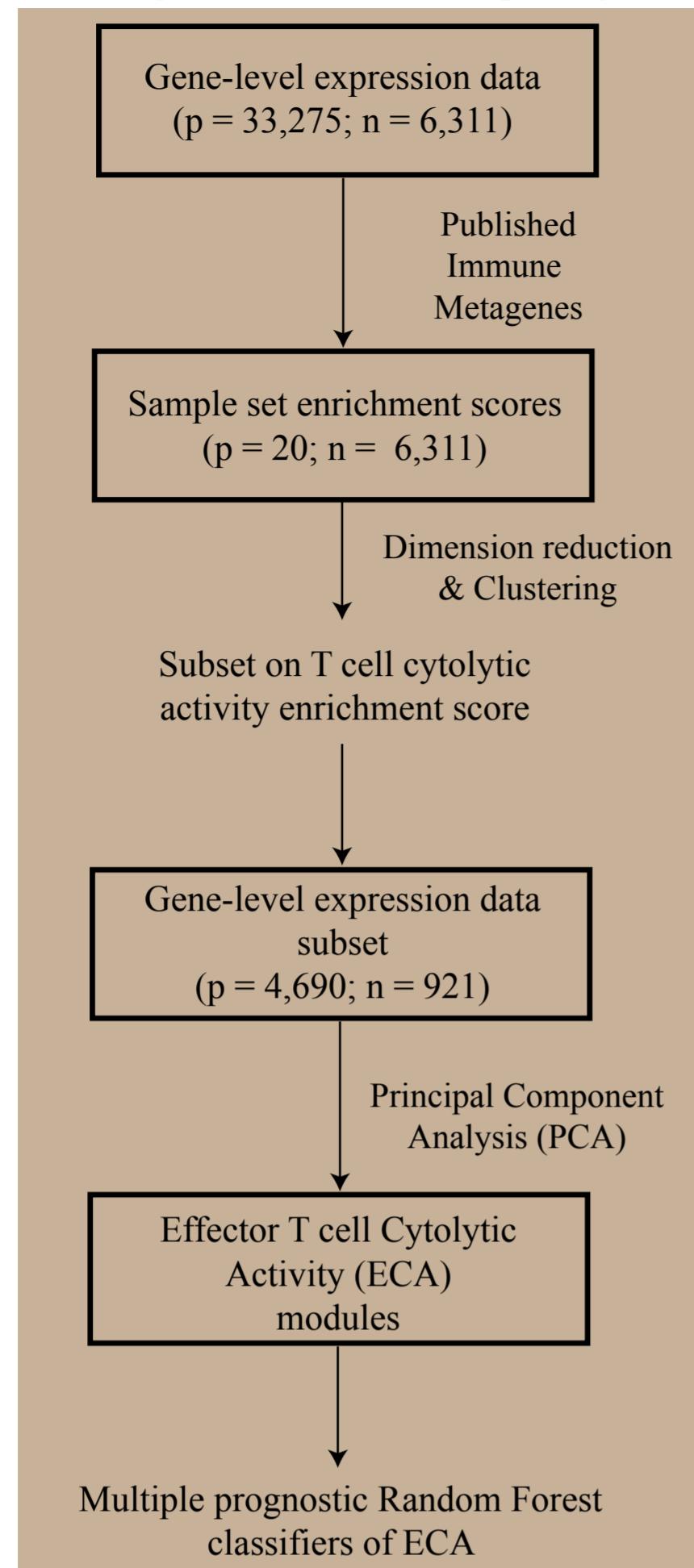
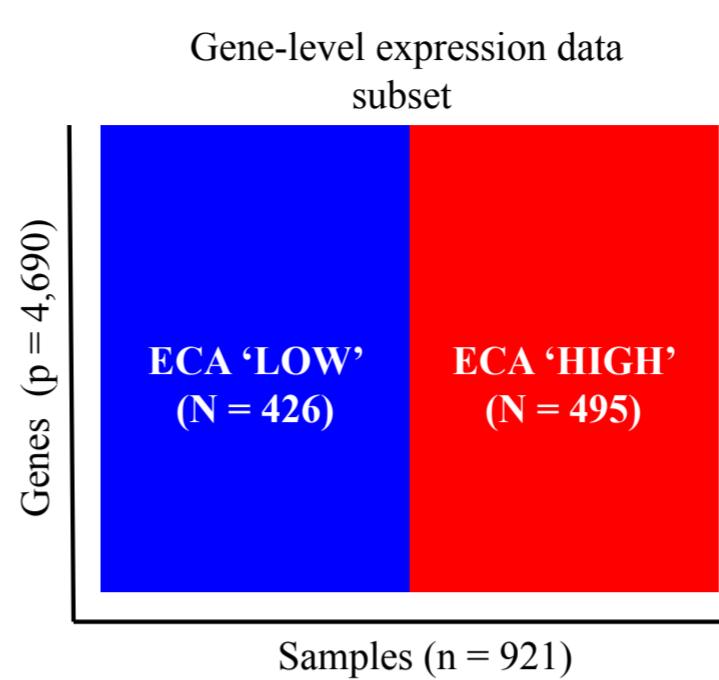


a.

CD3+ single T cells from human primary TNBC



b.



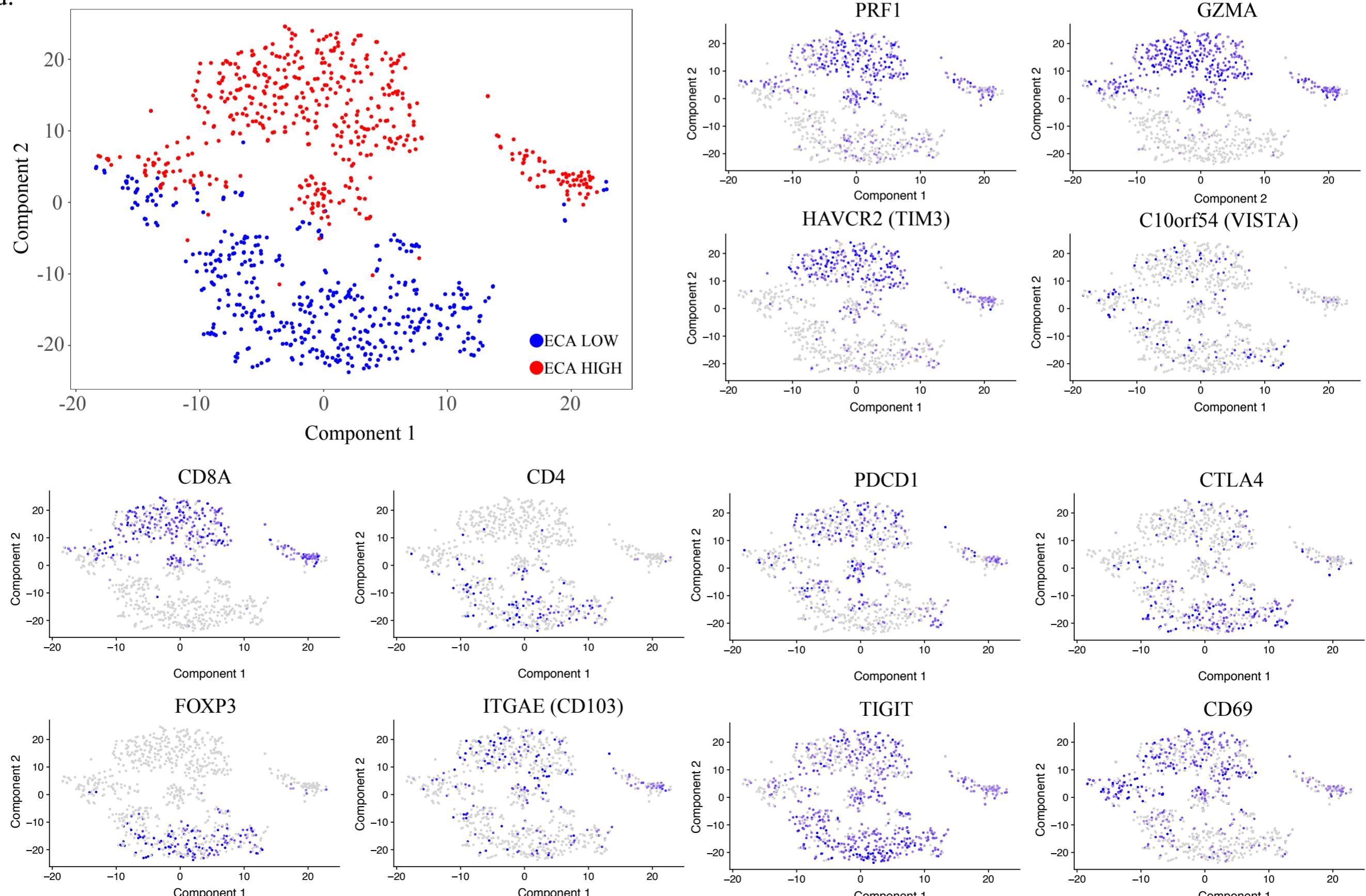
MODULES

MOD1	(1,438 genes)
MOD2	(1,018 genes)
MOD3	(382 genes)
MOD4	(374 genes)
MOD5	(301 genes)
MOD6	(339 genes)
MOD7	(226 genes)
MOD8	(150 genes)

c.

	MOD1	MOD2	MOD3	MOD4	MOD5	MOD6	MOD7	MOD8
MOD1	1438	334	161	155	101	85	73	43
MOD2		1018	182	212	162	159	102	78
MOD3			382	140	132	117	55	43
MOD4				374	123	103	69	51
MOD5					301	99	43	41
MOD6						339	52	54
MOD7							226	21
MOD8								150

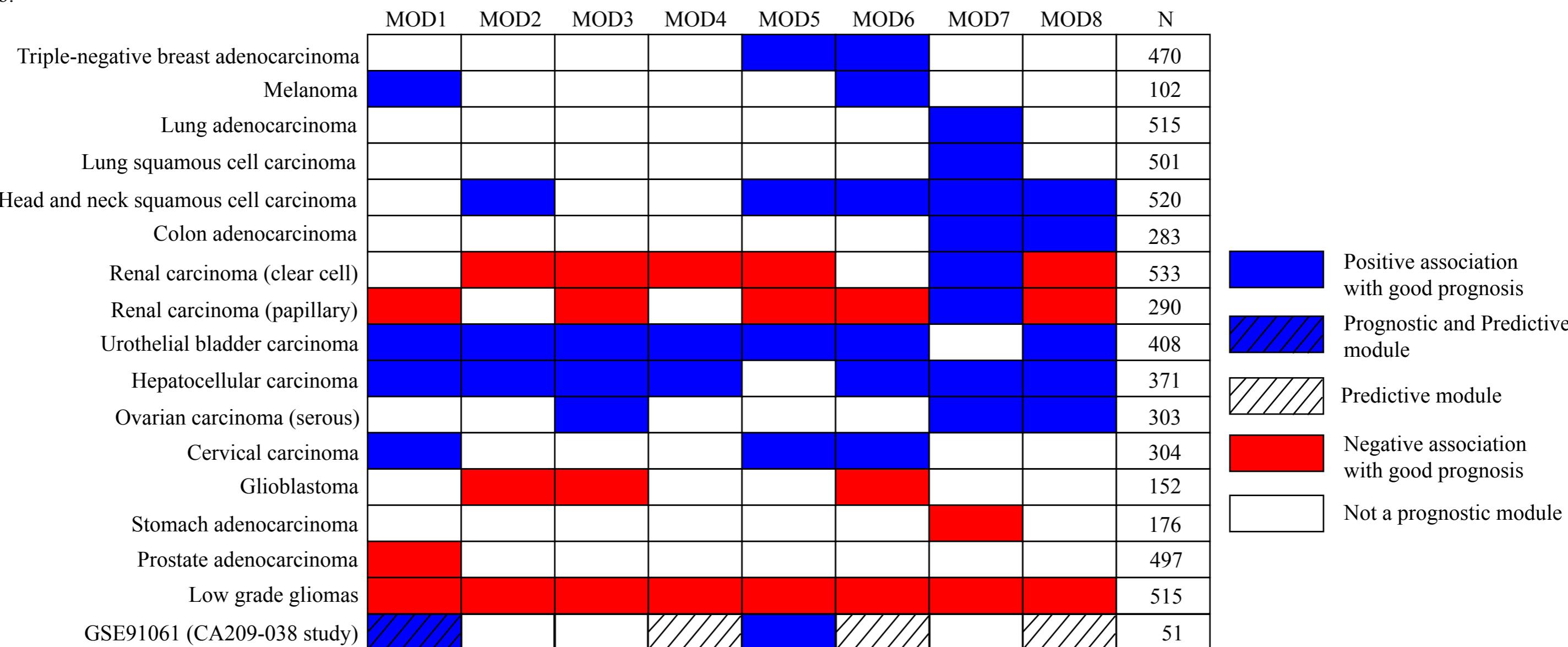
d.



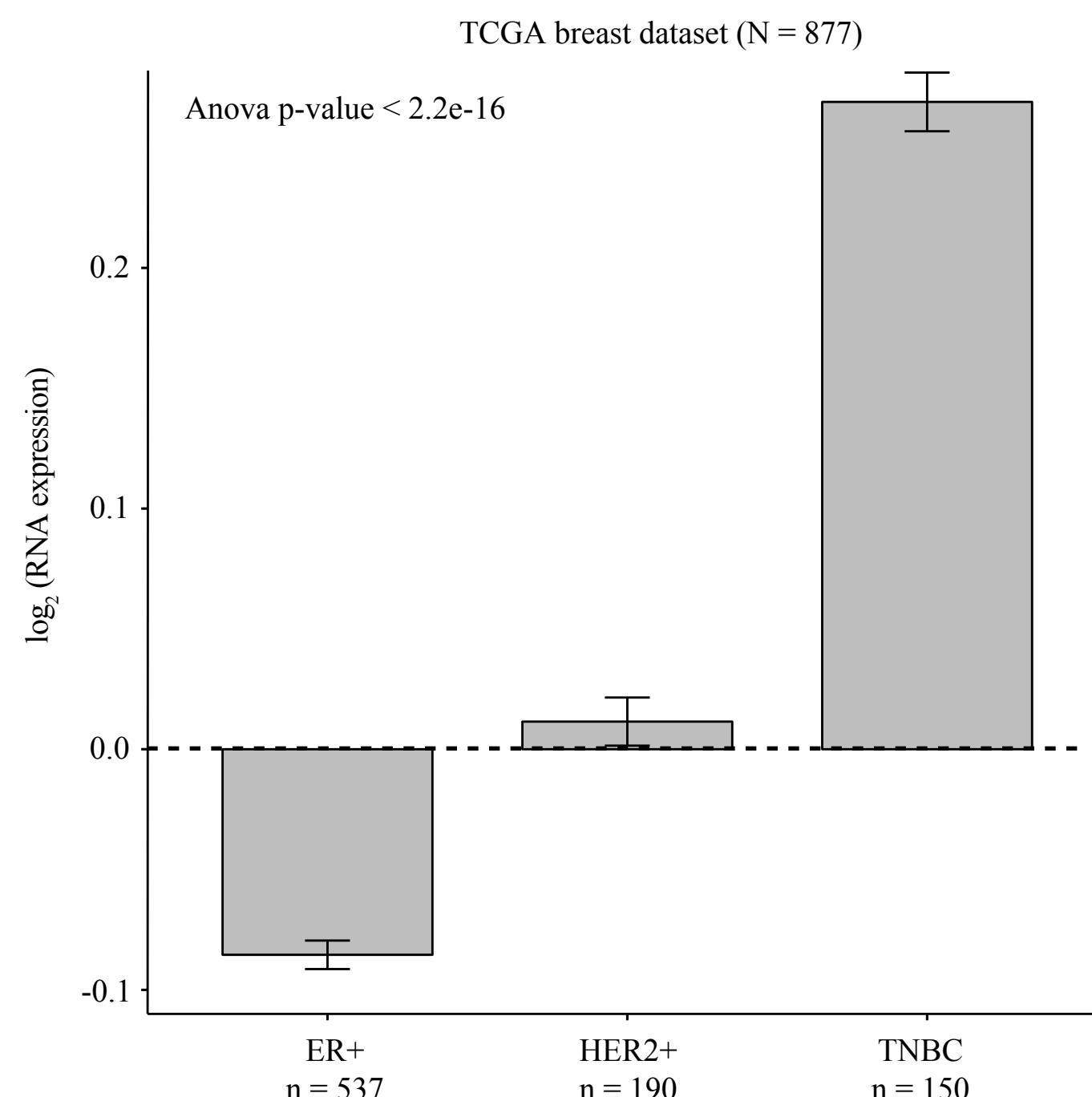
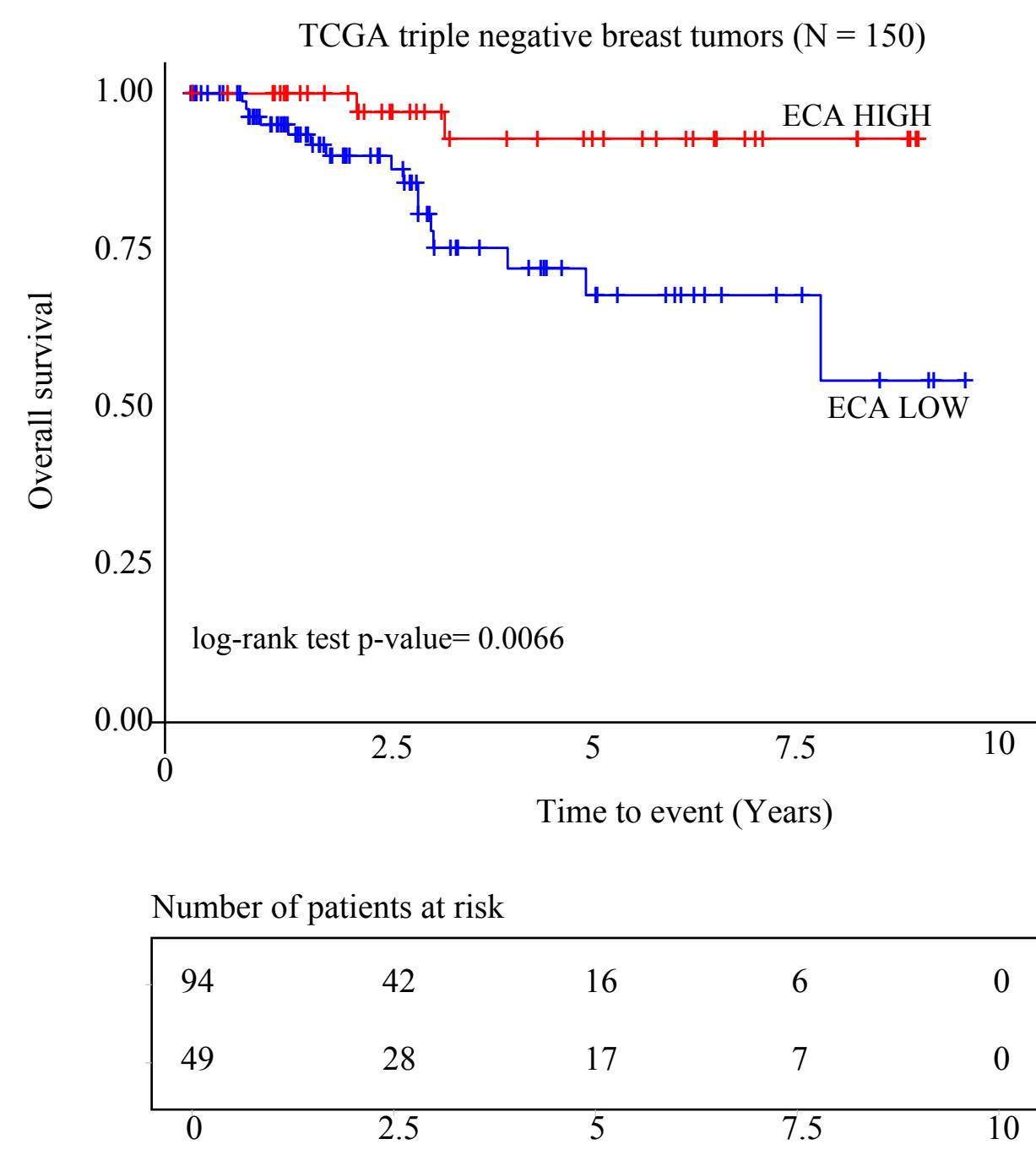
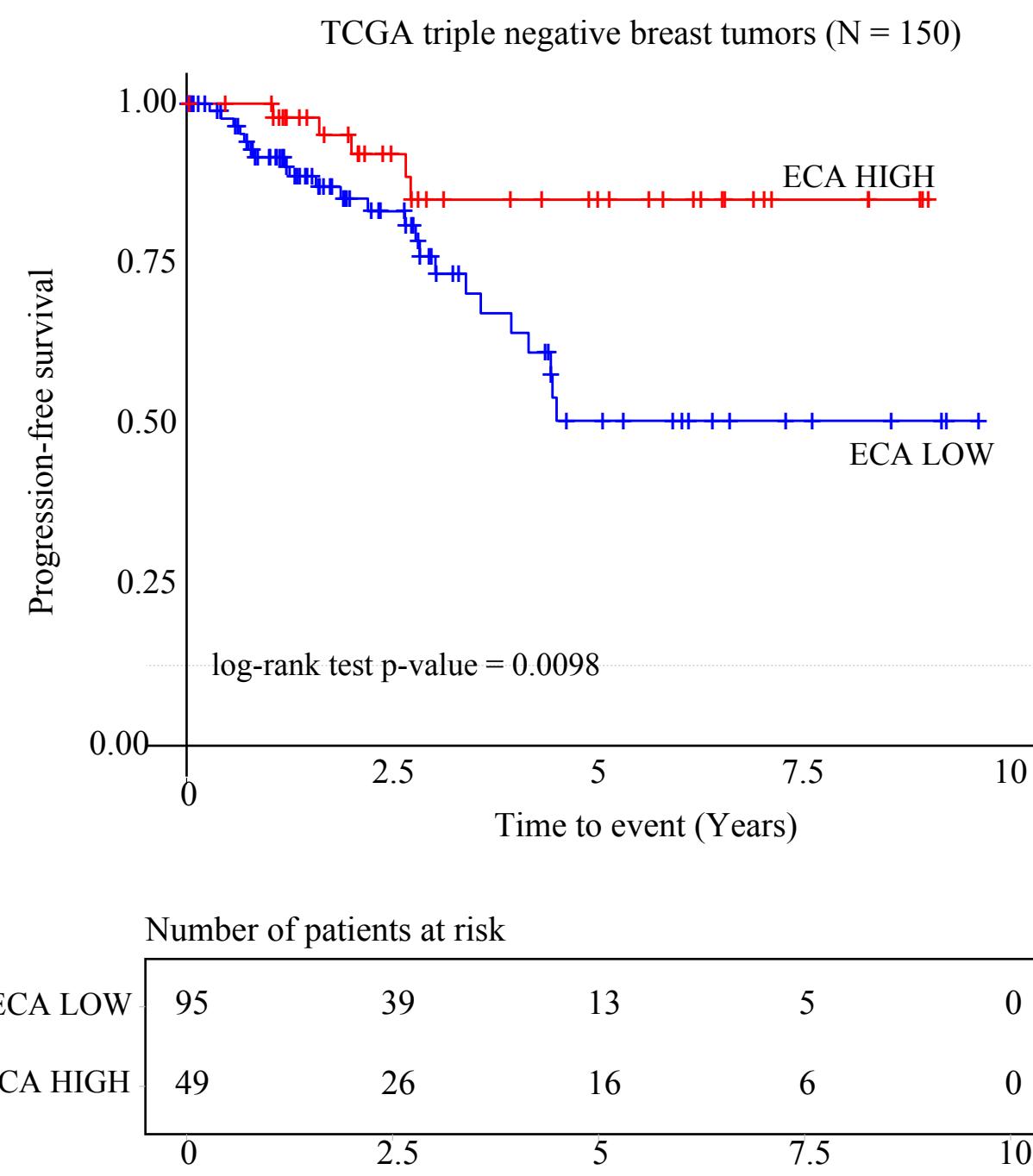
a.

	Tfh signature	Th1 signature	Treg signature	STAT1 signature	Immune1 signature	Immune2 signature	Interferon- γ signature	MOD1	MOD2	MOD3	MOD4	MOD5	MOD6	MOD7	MOD8
Tfh signature	8	0	0	0	1	2	1	4	2	1	2	1	0	2	
Th1 signature		12	1	1	0	2	3	2	5	5	6	3	3	2	0
Treg signature			7	0	0	0	0	2	0	2	1	1	0	0	
STAT1 signature				4	0	3	3	1	1	3	2	3	2	1	0
Immune1 signature					7	0	0	0	1	0	1	0	1	0	0
Immune2 signature						95	7	10	22	26	21	25	20	8	3
Interferon- γ signature							27	6	10	10	8	6	5	2	3

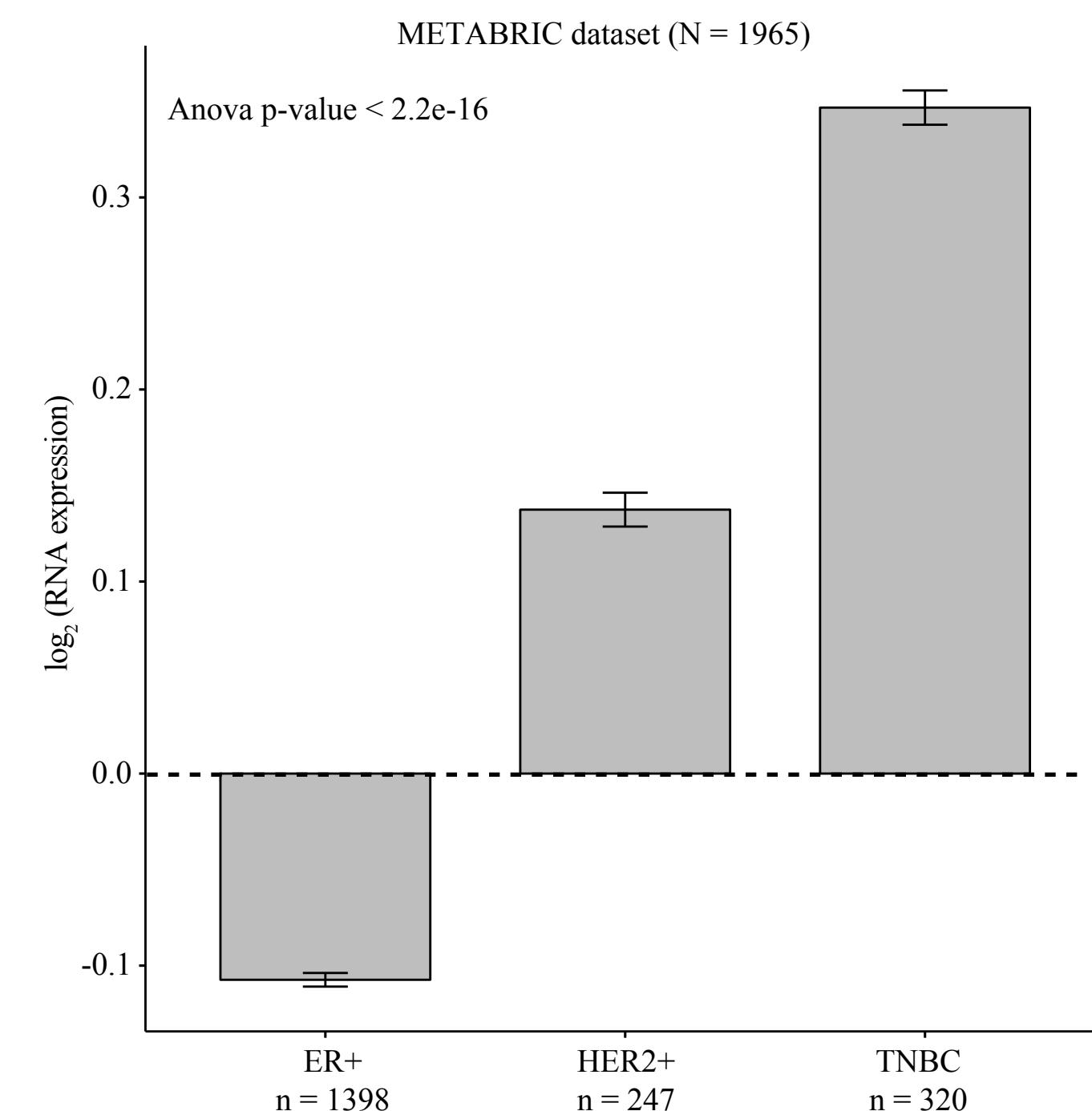
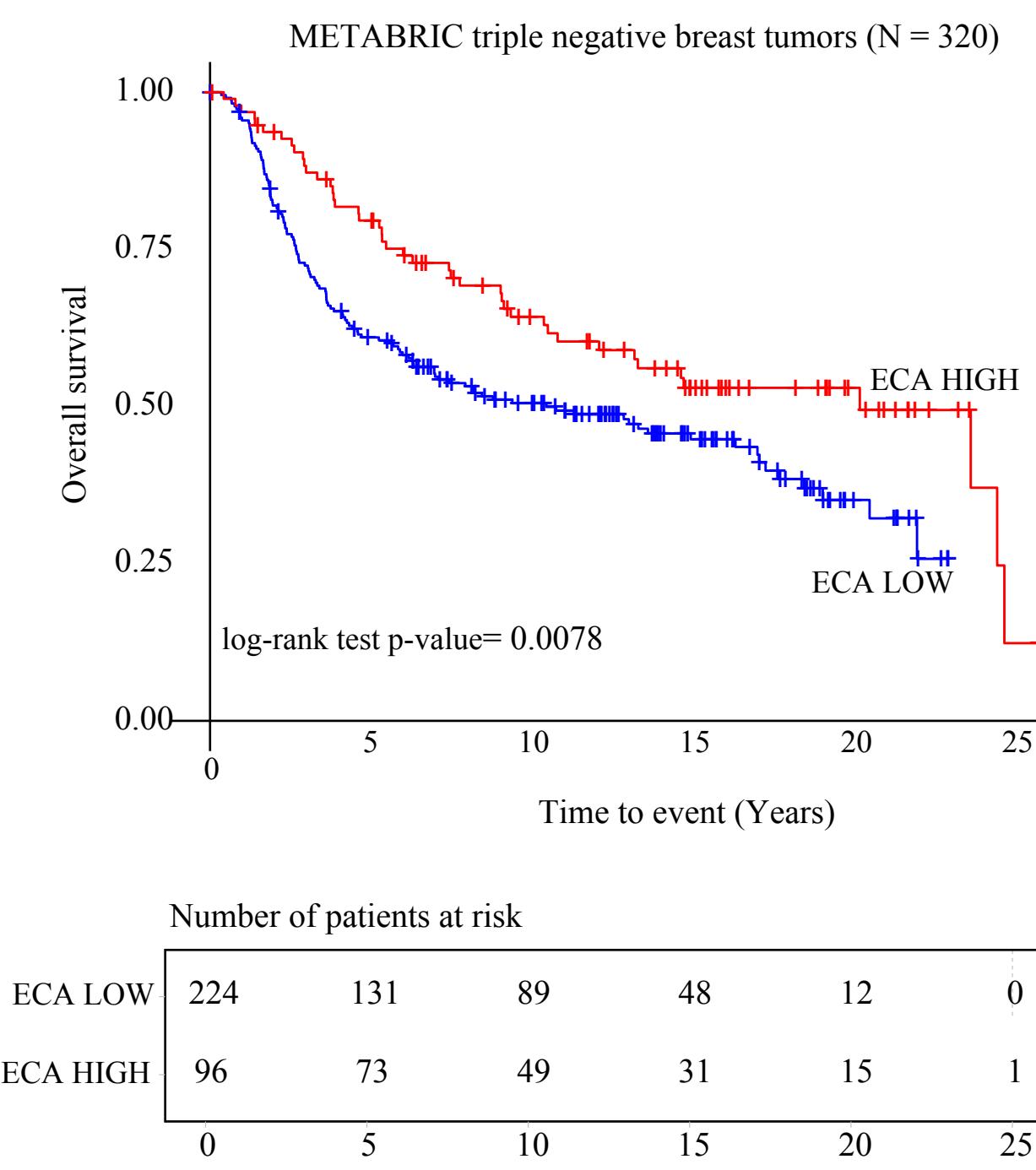
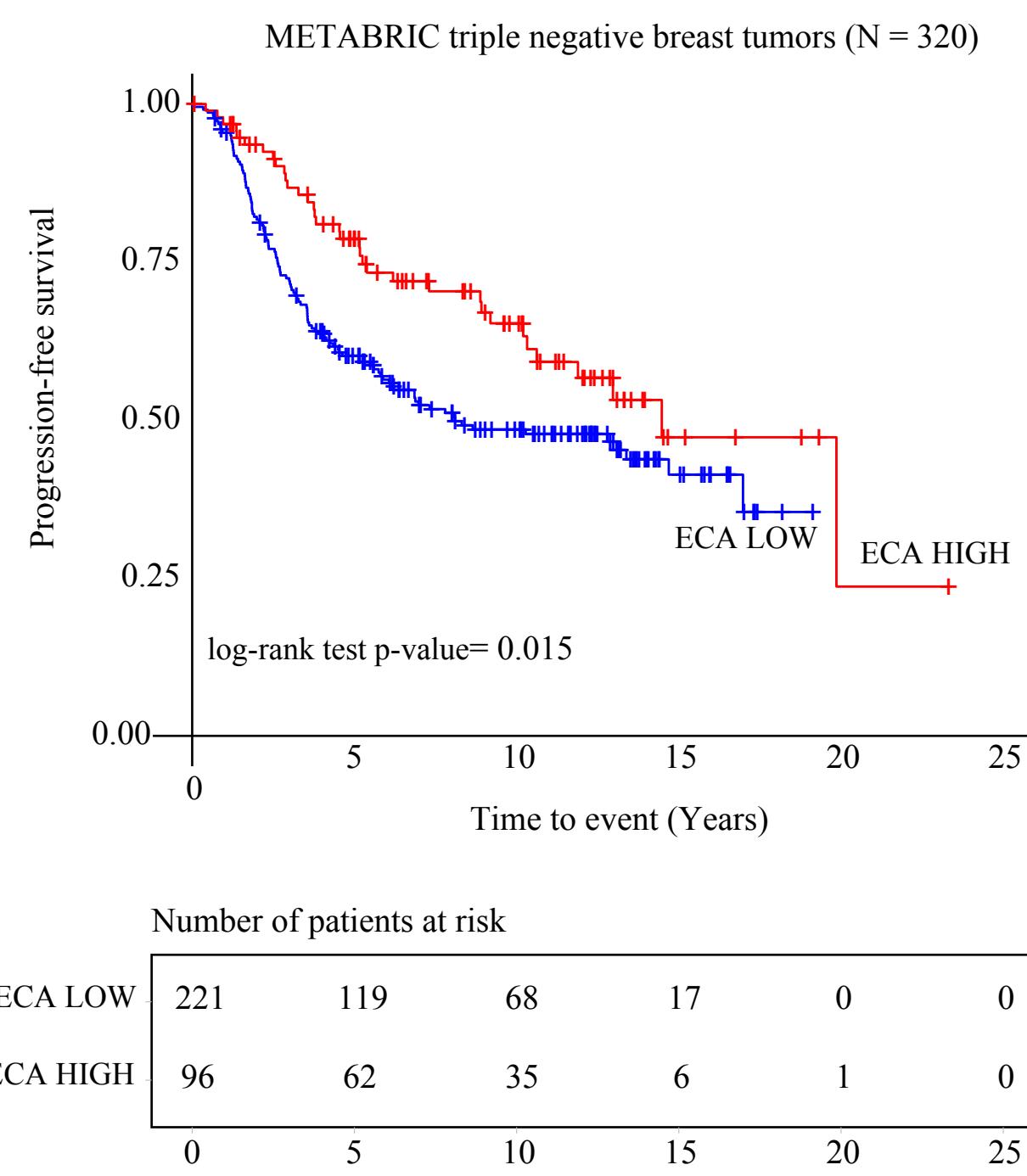
b.



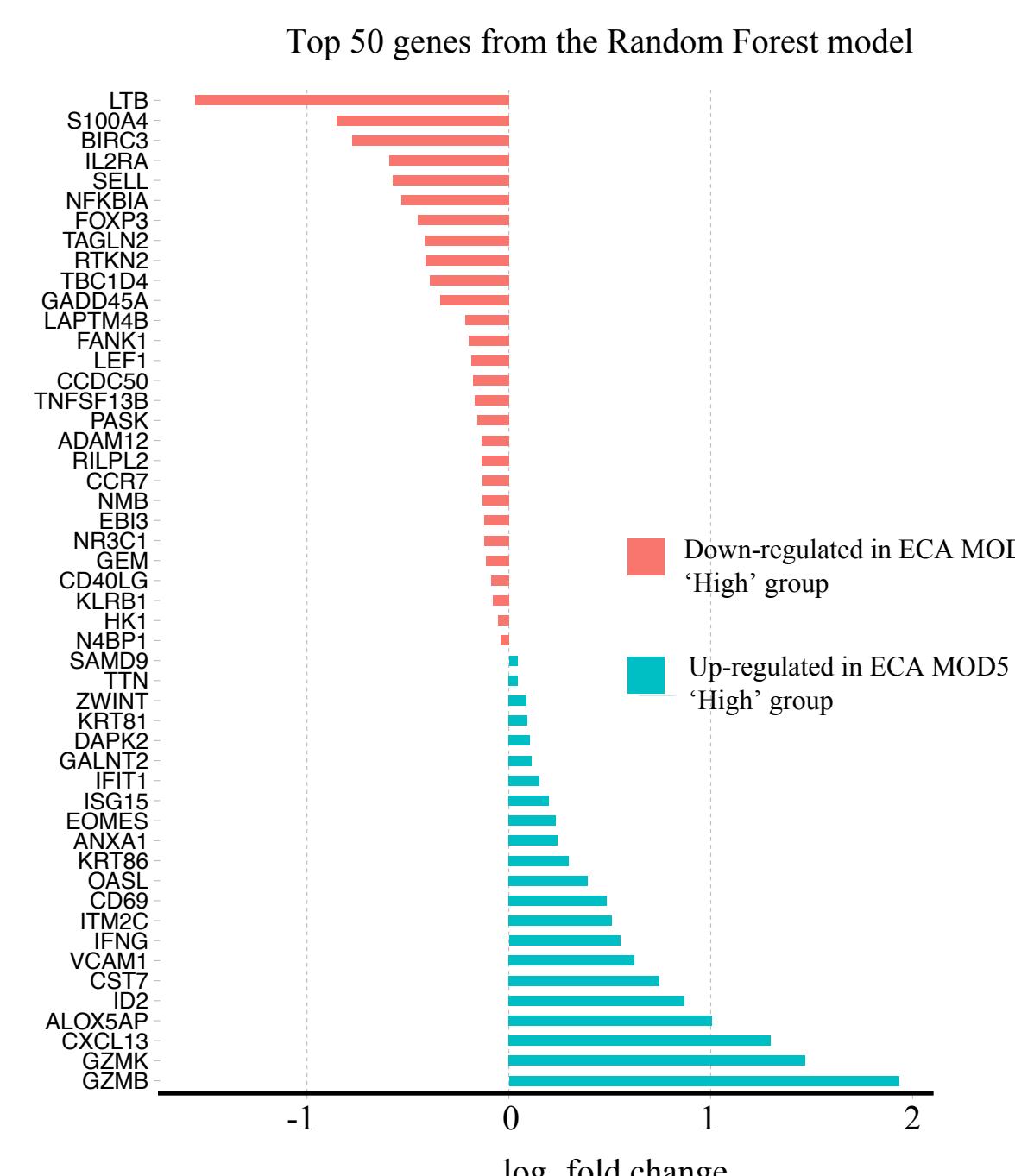
a.



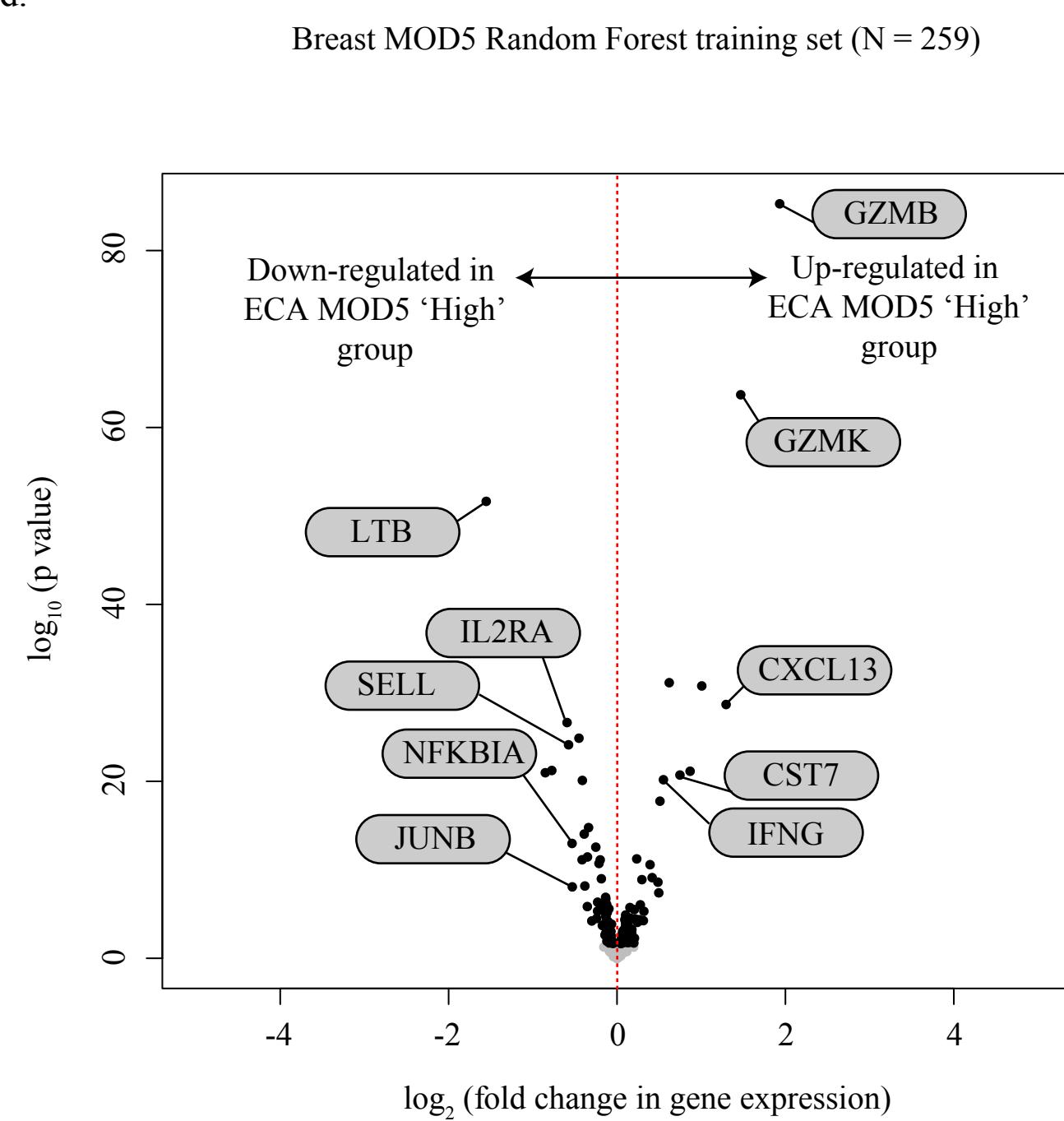
b.



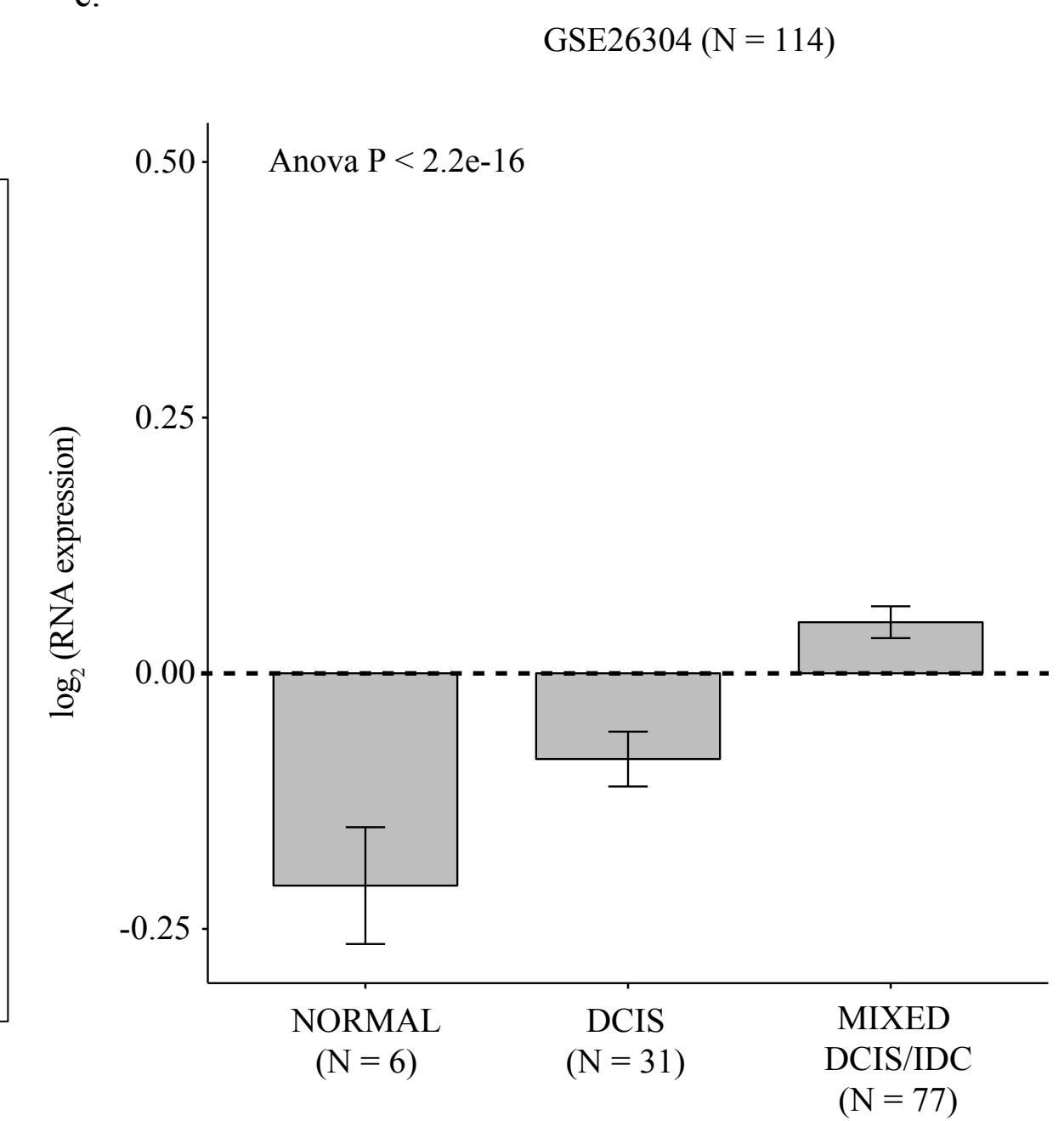
c.



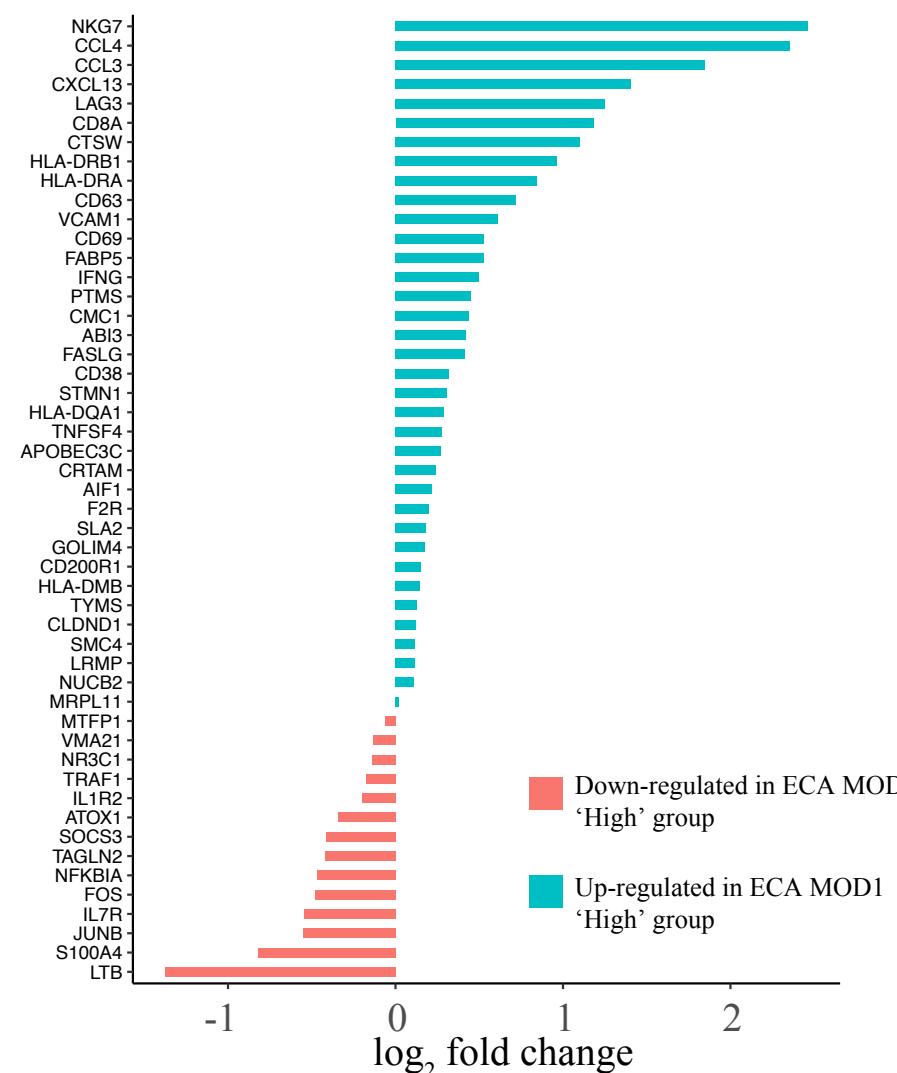
d.



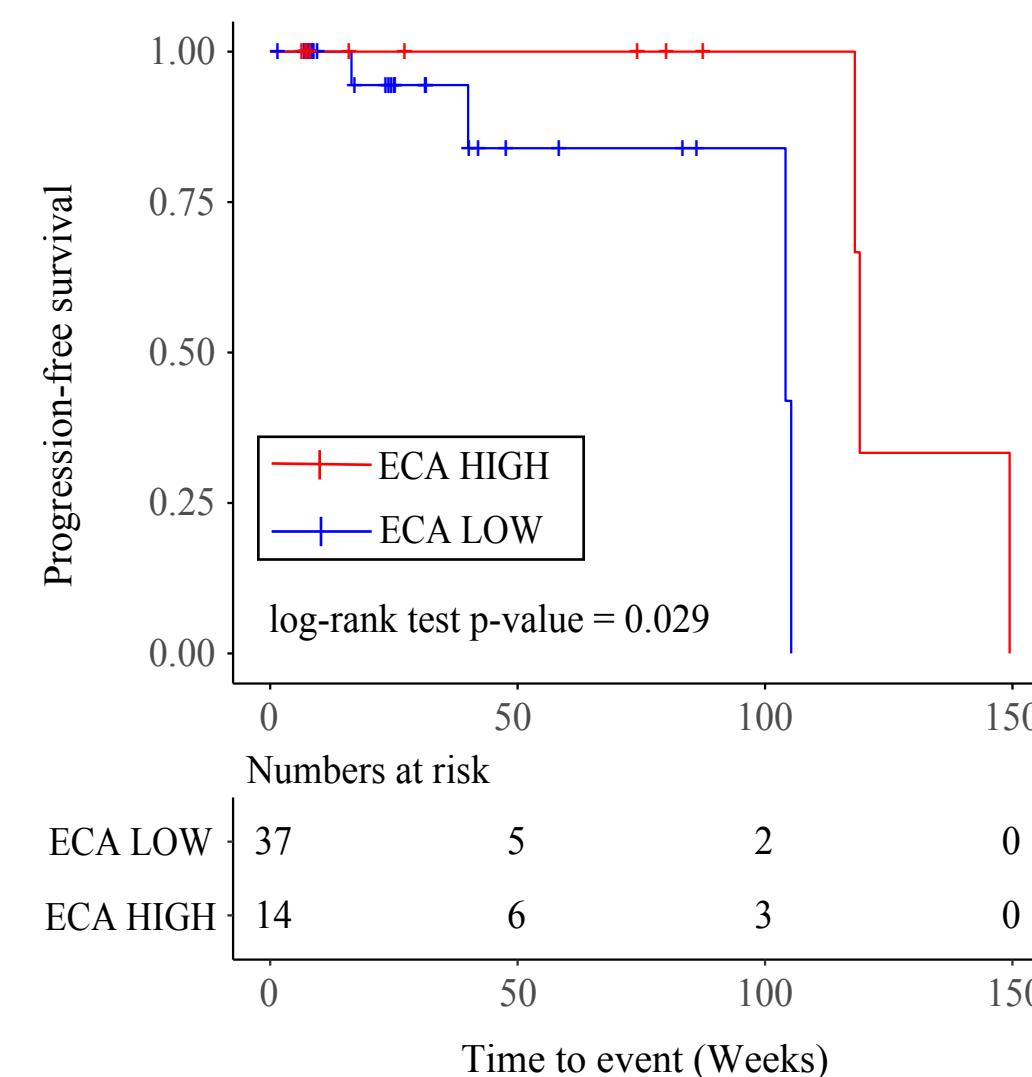
e.



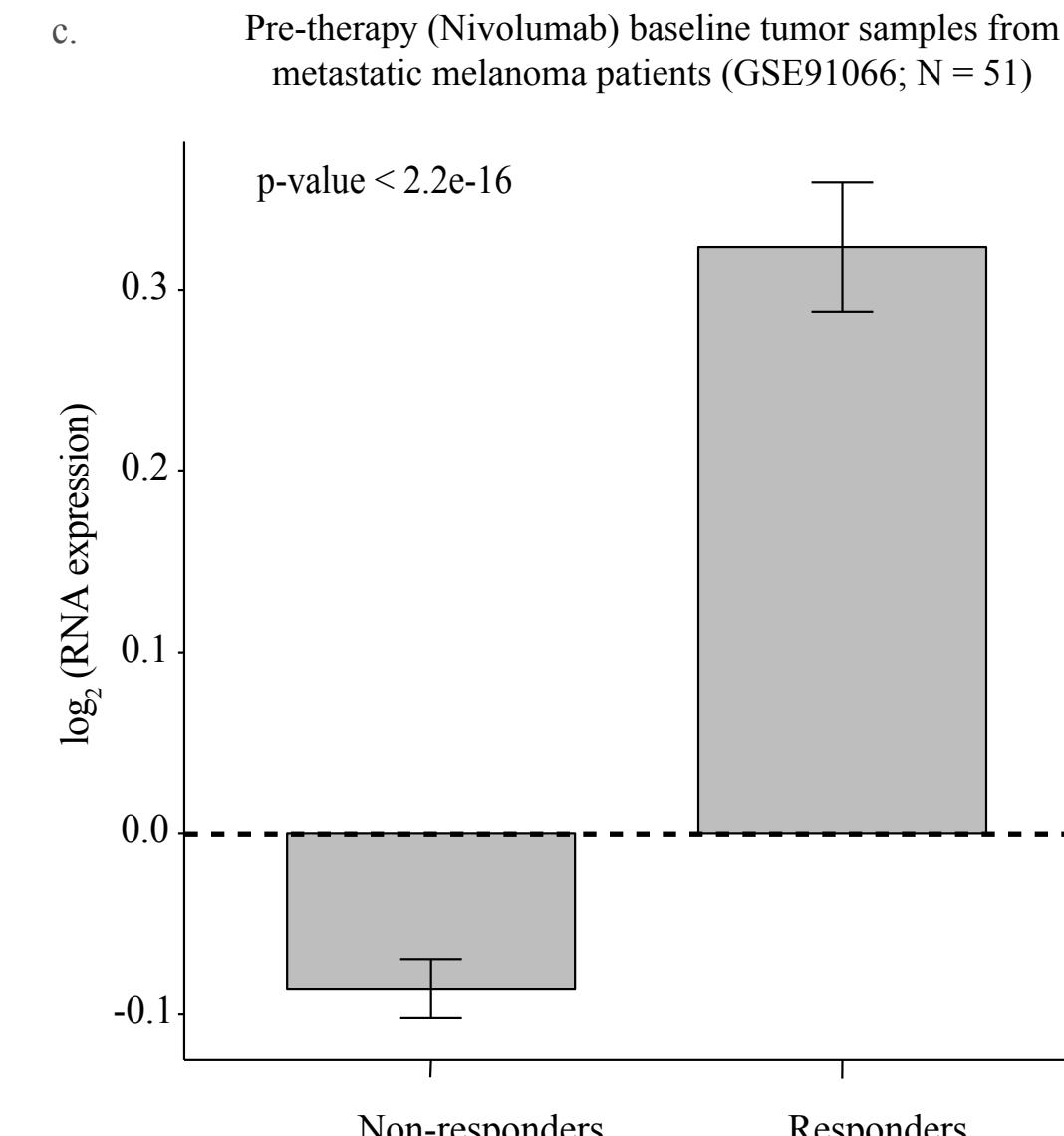
a. Top 50 genes from the Random Forest model



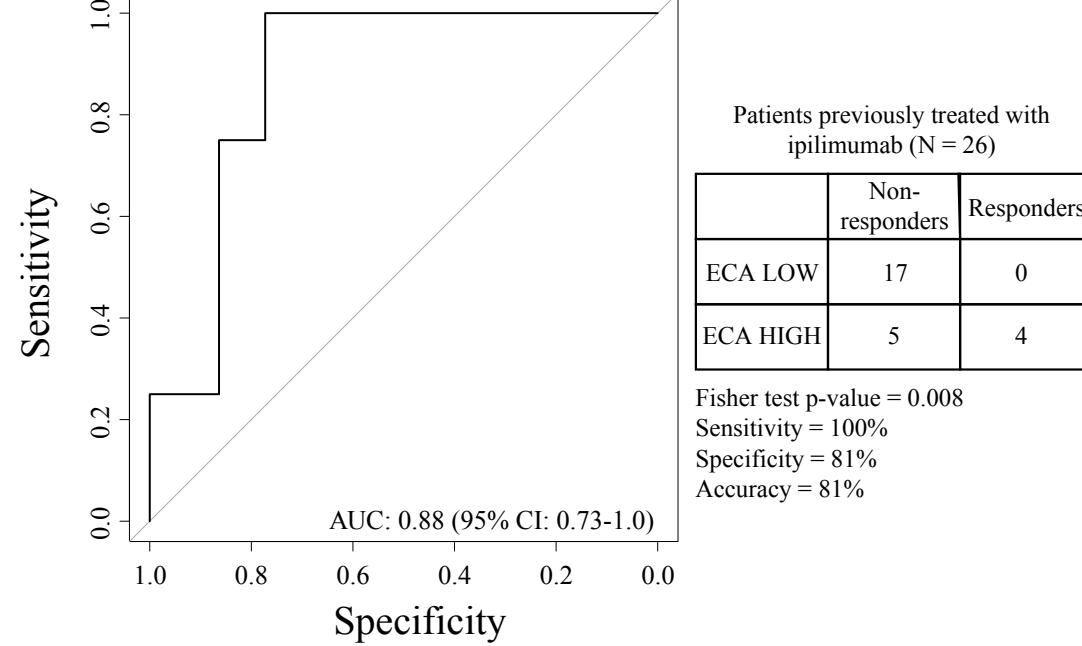
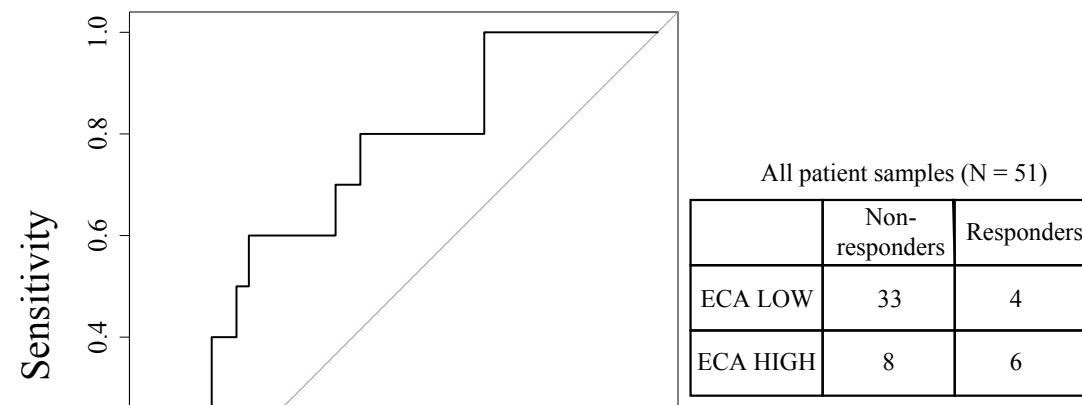
b.



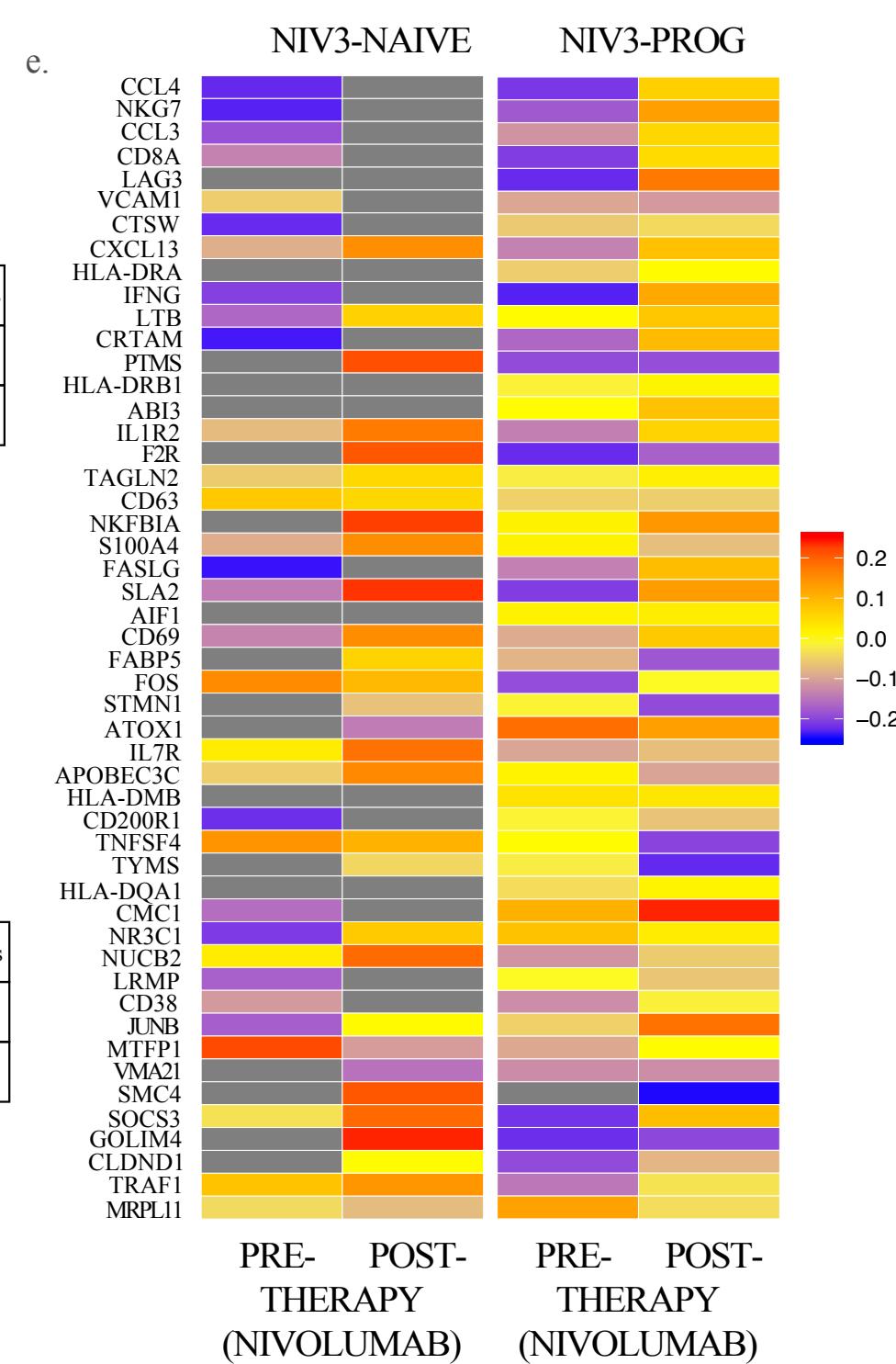
c.



d.



e.



f.

