

BreastC	M1 Chr9q31-34: 138	138	0	4	7	0	0	15	0	1	9	0	0	0	0	29	33	0	0	1	0	0	1	2	5	2	1	0	7	1	3	0	1	3	0	1	0	2	0	4	49	1	3
	M2 chr19q13: 23	0	23	0	0	0	0	2	0	0	0	0	0	0	1	4	3	4	0	0	0	0	0	0	0	0	0	1	0	0	4	0	0	0	1	0	0	0	0	0	1	0	
	M3 Chr16p11-13: 77	4	0	77	3	1	3	12	0	1	5	0	0	0	0	9	11	0	0	0	2	0	0	0	3	0	1	2	8	3	1	1	3	3	0	0	0	5	0	0	3	2	2
	M4 Ribosome: 487	7	0	3	487	7	4	46	7	15	2	2	20	1	3	90	46	3	0	3	10	0	1	0	21	0	5	17	10	67	3	13	0	3	9	4	0	2	2	1	8	0	1
	M5 Chr17q11-12,21: 99	0	0	1	7	99	3	9	1	1	0	5	3	2	1	9	6	0	0	0	0	0	1	2	3	0	10	7	2	8	0	5	1	2	1	1	0	0	0	0	1	1	0
	M6 Luminal: 313	0	0	3	4	3	313	23	27	29	0	2	4	3	54	5	4	3	1	2	9	0	3	0	3	0	1	5	2	7	1	5	4	1	4	4	0	1	0	2	0	2	5
	M7 CellCycle: 1959	15	2	12	46	9	23	1959	24	55	4	3	7	2	25	275	87	2	5	15	31	0	10	13	41	0	9	71	14	108	10	33	0	16	21	4	1	5	11	8	37	2	1
	M8 Angiogenesis: 1181	0	0	0	7	1	27	24	1181	464	0	1	2	26	202	8	1	1	8	28	33	0	1	0	4	8	0	9	1	217	1	124	1	1	20	150	0	0	0	0	1	1	2
	M9 EMT: 1878	1	0	1	15	1	29	55	464	1878	2	1	3	38	311	19	4	0	7	38	55	0	1	1	6	9	0	15	4	284	1	187	3	1	16	73	2	1	1	1	2	0	0
	M10 Chr22q11-13: 51	9	0	5	2	0	0	4	0	2	51	1	2	0	1	10	16	0	0	0	2	0	1	1	0	0	0	0	0	2	1	2	0	1	1	0	0	1	0	3	7	1	1
	M11 chr17p11-13_TP53: 34	0	0	0	2	5	2	3	1	1	1	34	2	0	1	8	3	0	0	0	0	0	0	1	0	0	3	0	0	1	0	3	0	0	0	0	0	0	0	1	0	0	1
	M12 Chr3p: 218	0	0	0	20	3	4	7	2	3	2	2	218	0	0	48	16	13	0	0	5	0	2	1	0	0	2	2	2	29	3	6	1	0	1	0	0	2	5	2	0	2	2
	M13 HOXB: 140	0	0	0	1	2	3	2	26	38	0	0	0	140	30	1	0	0	2	0	5	0	0	0	2	0	2	6	0	13	0	8	0	0	0	14	1	0	0	0	0	0	0
	M14 Basal: 904	0	1	0	3	1	54	25	202	311	1	1	0	30	904	12	1	1	6	28	50	0	2	2	6	6	0	7	1	199	1	126	1	1	13	71	2	1	0	1	3	1	2
	M15 Haptotaxis SMARCA2-Correlated: 1455	29	4	9	90	9	5	275	8	19	10	8	48	1	12	1455	116	11	0	7	41	0	28	15	29	2	20	29	22	49	23	37	3	14	7	0	0	14	31	17	51	3	23
	M16 RNAprocessing Mitochondrion: 471	33	3	11	46	6	4	87	1	4	16	3	16	0	1	116	471	1	0	4	6	0	1	2	16	0	10	13	30	32	8	11	1	2	2	2	0	3	3	4	40	1	2
	M17 CommonCancerGenes: 76	0	4	0	3	0	3	2	1	0	0	0	13	0	1	11	1	76	0	0	3	0	1	0	1	0	0	0	0	9	0	3	0	1	1	0	0	0	2	0	2	1	1
	M18 MastCells: 32	0	0	0	0	0	1	5	8	7	0	0	0	2	6	0	0	0	32	1	3	0	0	0	0	0	0	1	0	17	0	0	0	0	1	4	0	0	0	0	0	0	0
	M19 Metalthiolate: 190	1	0	0	3	0	2	15	28	38	0	0	0	0	28	7	4	0	1	190	28	0	1	0	1	0	0	10	0	59	0	36	0	0	4	1	0	3	0	2	0	2	1
	M20 InterferonResponse: 858	0	0	2	10	0	9	31	33	55	2	0	5	5	50	41	6	3	3	28	858	0	4	3	4	0	0	12	6	524	2	50	0	1	24	10	0	0	2	0	1	0	0
	M21 Chr1q21: 1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
	M22 Chr6q: 84	1	0	0	1	1	3	10	1	1	1	0	2	0	2	28	1	1	0	1	4	0	84	3	2	0	1	4	1	9	2	2	1	3	1	0	0	1	7	4	0	3	1
	M23 Chr18 Chr19p Chr5p: 65	2	0	0	0	2	0	13	0	1	1	1	1	0	2	15	2	0	0	0	3	0	3	65	2	1	2	0	1	2	1	0	2	1	0	1	0	2	1	1	1	2	0
	M24 Chr8q: 128	5	0	3	21	3	3	41	4	6	0	0	0	2	6	29	16	1	0	1	4	0	2	2	128	0	4	6	2	3	7	0	6	8	1	0	0	5	4	0	13	4	1
	M25 Chr5q31: 27	2	0	0	0	0	0	0	8	9	0	0	0	0	6	2	0	0	0	0	0	0	0	1	0	27	0	0	1	4	0	1	0	0	0	12	5	0	0	0	0	0	0
	M26 Chr17q21-25: 67	1	0	1	5	10	1	9	0	0	0	3	2	2	0	20	10	0	0	0	0	0	1	2	4	0	67	0	0	0	1	0	2	3	0	1	0	0	1	0	2	0	1
	M27 Nucleosome: 281	0	0	2	17	7	5	71	9	15	0	0	2	6	7	29	13	0	1	10	12	0	4	0	6	0	0	281	3	43	1	24	0	2	9	14	0	0	1	0	1	0	1
	M28 Chr11p15,q12-14: 102	7	1	8	10	2	2	14	1	4	0	0	2	0	1	22	30	0	0																								

