

BreastC

M15 Haptotaxis SMARCA2-Correlated: 1731 1455	M1 high-CpG-densityPromoters: 1192 145	10 0	1 9	11 0	27 49	3 4	0 2	10 1	8 2	1 27	0 18	4 0	3 0	89 27	15 0	4 7	0 1	9 82	4 10	1 6	3 1	0 0	11 1	4 7	7 6	0 2	2 1	18 19	0 1	0 9	0 0	0 0	0 1	3 6	0 0	0 0	2 0	0 0	
	M2 chr19q13: 110 23	2 0	0 0	2 1	0 1	0 0	0 0	0 0	11 2	0 1	0 0	0 0	72 10	0 0	3 1	0 0	0 0	0 5	0 0	2 1	0 0	0 4	9 0	0 0	1 0	0 0	0 0	0 0	0 0	1 0	0 0	0 0	0 0	0 0	0 0	0 0	0 0	0 0	
	M3 Chr16p11-13: 321 77	31 0	1 3	16 3	19 14	2 2	3 1	16 3	8 1	0 4	0 3	6 0	12 1	0 1	18 0	1 1	0 2	125 40	1 4	7 2	0 1	0 2	20 0	5 2	17 4	0 0	3 1	2 2	0 0	1 6	1 0	0 0	0 2	2 2	0 1	0 1	1 0	0 0	
	M4 Ribosome: 104 487	2 1	0 22	4 2	4 108	5 10	0 19	0 7	1 20	2 12	0 13	1 0	4 11	1 7	1 0	2 9	0 0	22 160	9 82	3 8	0 9	0 21	1 0	1 7	0 6	0 2	0 58	3 37	1 2	4 69	0 10	0 0	0 23	0 10	0 0	0 11	2 4	0 2	
	M5 Chr17q11-12,21: 204 99	6 0	117 42	8 2	2 11	0 2	0 1	5 3	5 5	1 0	0 0	3 0	6 1	0 3	11 0	3 3	0 1	7 13	0 1	2 0	1 2	0 3	15 1	3 2	1 7	0 1	0 3	0 0	0 0	0 0	0 3	0 1	0 0	0 4	0 2	0 0	0 3	1 0	0 0
	M6 Luminal: 1860 313	131 6	28 4	189 66	69 10	22 0	3 0	219 120	153 78	7 0	18 4	61 2	105 8	32 1	79 0	31 3	3 4	94 3	18 3	35 1	58 5	7 14	215 4	18 3	43 8	8 1	26 8	13 1	13 2	12 0	1 2	0 0	2 17	40 9	0 3	0 4	11 2	0 0	
	M7 CellCycle: 1119 1959	38 0	15 89	61 18	247 934	34 108	13 22	31 24	28 92	13 38	35 103	15 0	41 15	35 33	37 0	26 42	6 6	109 290	11 33	35 22	28 90	0 48	63 1	40 54	11 31	8 10	11 110	2 71	6 7	6 47	2 29	0 0	1 45	89 92	0 5	0 52	15 31	0 1	
	M8 Angiogenesis: 661 1181	30 25	0 2	51 59	7 5	6 3	0 0	31 125	306 851	0 2	1 3	19 1	12 2	1 0	26 7	1 1	0 0	3 2	2 1	9 1	2 1	0 125	74 59	1 1	8 15	0 2	39 224	0 2	1 6	3 4	4 35	0 0	0 53	5 1	1 32	0 14	9 1	0 3	
	M9 EMT: 1140 1878	39 7	1 2	72 75	13 28	10 2	1 0	47 200	680 1623	1 2	2 4	10 3	33 5	4 1	23 6	6 2	2 4	25 18	0 1	9 3	1 1	2 209	58 21	5 1	12 21	2 3	22 283	5 4	1 7	1 8	1 27	0 0	2 94	9 2	1 66	0 18	8 1	0 4	
	M10 Chr22q11-13: 205 51	5 0	2 5	6 0	2 13	1 0	1 4	10 2	4 2	106 30	0 7	3 0	9 1	1 1	8 1	1 1	0 1	21 19	0 6	0 3	0 0	0 4	14 0	2 0	1 1	0 0	1 2	0 4	0 1	1 2	0 1	0 0	1 3	3 4	0 0	0 0	0 0	0 0	
	M11 chr17p11-13_TP53: 162 34	12 0	3 6	8 1	2 5	2 0	0 0	2 1	3 2	0 1	0 0	5 0	2 1	7 1	5 0	0 2	0 0	1 3	1 0	2 1	0 0	0 4	3 0	7 1	87 12	0 0	0 0	0 0	0 0	0 0	0 0	0 0	0 0	1 2	0 0	0 0	1 0	0 0	
	M12 Chr3p: 588 218	25 0	2 4	25 4	15 39	6 1	0 6	24 0	36 7	8 1	0 1	6 0	157 71	6 0	13 2	22 3	0 1	39 42	75 15	7 5	2 5	0 12	45 0	4 0	15 9	0 2	3 19	3 9	1 0	5 2	0 2	0 0	3 10	24 18	0 0	0 3	7 8	0 0	
	M13 HOXB: 13 140	0 6	0 2	10 15	0 2	0 0	0 0	1 5	1 68	0 0	0 0	0 0	0 1	0 0	0 3	0 0	0 0	0 0	0 0	0 0	0 1	0 7	0 0	0 0	0 0	0 0	1 11	0 0	0 0	0 1	0 2	0 0	0 6	0 0	0 68	0 12	0 0	0 0	
	M14 Basal: 1178 904	64 9	4 1	126 86	36 16	11 1	2 0	151 304	205 466	5 1	2 3	47 3	37 1	10 3	39 3	7 2	3 3	36 4	0 1	15 3	15 3	3 139	155 34	22 3	11 8	8 2	51 223	1 0	9 9	14 6	3 20	0 0	4 79	13 6	9 54	0 11	13 2	1 3	
	M16 RNAprocessing Mitochondrion: 1086 471	36 0	5 71	82 8	424 635	126 61	12 25	61 16	48 55	9 27	7 92	9 0	178 81	37 32	12 0	34 64	0 5	24 240	50 78	78 46	105 80	1 53	32 0	30 37	23 35	5 6	16 42	124 145	14 19	12 63	0 8	0 0	2 51	104 102	0 2	0 23	13 42	0 0	
	M17 CommonCancerGenes: 608 76	71 1	6 36	28 2	55 131	6 8	66 27	33 6	34 9	1 33	6 33	19 0	15 8	9 8	49 2	7 14	1 1	318 265	19 37	45 18	2 11	2 25	100 0	43 20	8 15	5 1	3 18	7 39	0 0	3 26	0 3	0 0	1 12	95 56	0 2	0 8	2 4	0 0	
	M18 MastCells: 16 32	36 1	4 0	23 9	25 0	8 0	2 0	6 2	16 3	2 0	3 1	10 0	287 55	10 0	19 1	5 2	0 1	8 2	1 0	5 1	5 1	1 6	37 0	27 2	7 1	2 0	9 7	4 5	5 1	0 1	0 0	2 3	5 0	0 1	0 0	7 5	1 2		
	M19 Metallothiolate: 8 190	1 0	0 0	0 3	0 1	0 1	0 0	1 5	0 10	0 1	0 1	0 0	0 0	0 0	1 0	0 0	0 0	0 0	0 0	0 0	0 0	0 2	4 1	0 0	0 0	0 0	0 15	0 0	0 0	0 0	8 14	0 3	0 1	0 4	0 1	0 0	0 0	0 1	
	M20 InterferonResponse: 92 858	0 0	0 0	0 4	0 8	0 1	0 0	8 60	0 65	0 0	0 1	0 0	0 2	0 2	0 0	0 3	0 2	0 4	0 0	0 0	0 2	44 1	0 0	0 0	0 0	0 0	0 58	0 0	0 4	0 1	0 5	0 0	0 35	0 3	0 4	0 9	0 0	0 0	0 1
	M21 Chr1q21: 9 1	2 2	0 2	2 10	3 21	0 2	1 0	4 47	2 97	3 9	0 2	0 0	0 4	2 0	4 1	0 1	1 1	3 20	0 0	0 3	1 10	0 58	2 3	1 2	0 7	0 1	3 379	0 3	0 5	0 4	0 35	0 0	56 712	1 5	0 6	0 21	0 2	0 1	
	M22 Chr6q: 182 84	0 0	0 0	0 0	0 0	0 0	0 0	0 0	2 0	0 0	0 1	0 0	4 0	0 0	0 0</																								