

Supplementary Table 2 - Gene counts per prognostic signature

The number of genes with data in each annotation and platform for the signatures.

Signature	# of genes	Default Annotation		Alternative Annotation	
		U133A	U133 Plus 2.0	U133A	U133 Plus 2.0
Buffa metagene	51	42	46	44	48
Chi	145	113	129	118	137
Elvidge	178	167	170	169	170
Hu	13	12	12	12	12
Seigneuric 0% early	68	54	67	52	65
Seigneuric 0% early	34	28	30	27	31
Sorensen	28	19	22	21	27
Winter metagene	101	79	90	77	92
Cluster 1	69	48	60	52	68
Cluster 2	246	205	231	211	246
Cluster 3	157	114	144	115	155
Cluster 4	95	61	86	64	92
Cluster 5	162	119	142	123	159
Cluster 6	14	8	11	10	14
Cluster 7	28	19	25	15	26