Supplementary Table S5. Genes enriched in cancer cell compartment, identified using the limma package in R. The magnitude of enrichment is indicated by the the difference in log2 expression (logFC).

	Symbol	logFC	adj.P.value		Symbol	logFC	adj.P.value		Symbol	logFC	adj.P.value
1	DSG2	2.45	0.0124	17	ELF3	1.78	0.0307	33	CLDN7	1.61	0.0474
2	RAB25	2.36	0.0136	18	MPPED2	2.21	0.0307	34	AP1M2	1.71	0.0477
3	MAL2	2.21	0.0136	19	MUC16	2.03	0.0327	35	MOCOS	1.61	0.0477
4	ELF5	2.26	0.0136	20	BAIAP2L1	1.77	0.0328	36	CLDN4	1.73	0.0477
5	VTCN1	2.34	0.0187	21	PFN2	1.77	0.0344	37	CKS1B	1.81	0.0484
6	DSP	2.15	0.0215	22	ANLN	1.85	0.0344	38	CKS1B	1.77	0.0489
7	IRF6	1.95	0.0215	23	DSC2	1.67	0.0377	39	SERPINA3	1.78	0.0493
8	LOC151009	1.98	0.0219	24	TM4SF1	1.96	0.0390	40	HIST1H2BK	1.67	0.0493
9	PEG10	1.93	0.0237	25	FRMD3	1.79	0.0390	41	MYO6	1.74	0.0493
10	CD24	2.12	0.0246	26	SORBS2	1.94	0.0409	42	CHPT1	1.61	0.0494
11	PROM1	2.73	0.0280	27	ESRP1	2.50	0.0410	43	UBE2T	1.60	0.0494
12	ITGB8	2.39	0.0288	28	PHGDH	1.70	0.0414	44	SLC7A5	1.55	0.0494
13	EHF	2.28	0.0295	29	PRSS8	2.10	0.0425	45	PTPLB	1.69	0.0494
14	KRT7	1.93	0.0295	30	MYO5B	1.87	0.0443	46	GABRP	2.15	0.0494
15	MUC5B	1.92	0.0300	31	TRPS1	1.94	0.0456	47	GPR56	1.73	0.0499
16	GLYATL2	2.27	0.0303	32	SERTAD4	1.84	0.0462	48	MPZL2	1.95	0.0499