Additional file 11: The comparision of the top ten up- and down-regulated genes of African American and others with ESCC

GSE77861		GSE17351		GSE33810		GSE20347		GSE29001		GSE33426		GSE45670		GSE23400	
Gene	logFC	Gene	logFC	Gene	logFC	Gene	logFC	Gene	logFC	Gene	logFC	Gene	logFC	Gene	logFC
KRT17*	17.6	**CRISP3**	8.9	SFTA2	7.5	MMP1	5.1	MMP1	7.4	LHX2*	9.5	S100A7	8.3	MMP1	4.4
IGHG1	12.1	**TMPRSS11B**	7.5	SCGB2A2	7.0	SPP1	4.6	LHX2*	6.5	MAGEA3/6*	8.7	MMP1	8.2	SPP1	3.2
ODC1	10.1	CRNN	7.1	LGI1	6.9	COL11A1*	4.4	MAGEA3/6*	5.7	SNX10	7.9	MAGEA3/6*	7.3	POSTN*	3.1
IGKC*	9.9	**KRT78***	6.5	MUC5B*	6.8	NTS	3.9	HOXA9*	5.3	ZIC1	7.6	MMP10	6.6	COL1A1*	3.1
IGLC1*	9.7	EDN3	6.5	GREM2*	6.2	POSTN*	3.8	CXCL11*	5.3	LGALS1	7.6	MMP3	6.5	**KRT17***	2.8
PFN2	6.3	SPINK7	6.3	GYS2	6.2	ANO1	3.7	APOE*	5.1	MAGEA12	7.5	SPRR2G	5.8	COL1A2*	2.7
GPNMB	4.7	CEACAM7*	6.2	TFF3	6.1	MMP12	3.7	HOXD10	5.0	COL5A2*	7.4	**KRT17***	5.8	COL11A1*	2.4
EPCAM	4.7	**SCEL***	6.0	**CRISP3**	5.7	VCAN*	3.4	MMP13	5.0	MET*	7.3	S100A7A*	5.7	CDH11*	2.4
HAS3	4.1	**SLURP1**	6.0	MAB21L2*	5.5	CXCL8*	3.4	INHBA*	4.9	HOXA9*	7.2	DEFB4A/B	5.6	MMP12	2.2
ALCAM	4.1	SPINK5	5.9	B3GALT2*	5.4	COL1A2*	3.2	CCL18*	4.8	FBXO5	6.9	ABCA13*	5.3	MAGEA3/6*	2.2
CRISP3	-9	MMP1	-7.0	GAGE1*	-7.0	**CRISP3**	-7.2	**CRISP3**	-10.7	CRNN	-7.9	C7*	-7.9	**CRISP3**	-4.2
TMPRSS11B	-8.6	CST1	-6.4	COL11A1*	-7.0	CRNN	-6.4	CRNN	-8.8	KRT4*	-6.7	ADH1B*	-7.1	MAL	-4.0
CLCA4	-8.4	COL11A1*	-6.2	GAGE3	-6.5	MAL	-5.85	SYNPO2L	-8.5	**CRISP3**	-6.4	MFAP5*	-6.3	CRNN	-3.7
SCEL*	-8.0	MMP12	-5.8	TUSC7*	-6.0	**CRCT1**	-5.63	**CRCT1**	-7.3	**CRCT1**	-5.7	C2orf40	-6.3	**SCEL**	-3.5
SLURP1	-7.8	MMP13	-5.7	SKIL*	-5.7	TGM3	-5.57	GYS2	-7.2	SYNPO2L	-5.4	MYH11*	-6.2	**CLCA4**	-3.4
KRT78*	-6.2	MMP3	-5.4	CCNYL2	-5.6	**SCEL**	-5.36	FLG	-7.1	SPRR3	-5.2	RBPMS2	-6.2	TGM3	-3.3
GBP6	-6.0	HOXD10	-5.3	LOC283485*	-5.6	**CLCA4**	-5.34	SPRR2C	-7.0	ADGRF1	-5.1	CHRDL1	-6.1	**CRCT1**	-3.2
A2ML1	-5.6	NELL2	-5.1	TNNC1	-5.6	CLIC3	-5.33	ENDOU	-7.0	MAL	-4.9	PGM5-AS1*	-6.1	TMPRSS11E	-3.1
CRCT1	-5.6	ADAM12*	-5.0	KRTAP19-1	-5.5	KRT4*	-5.09	TGM3	-6.8	SPRR1A	-4.7	ATP1A2*	-6.1	**SLURP1**	-3.0
PPP1R3C	-5.5	DNMT3B	-5.0	CTAG1A/B*	-5.5	SPINK5	-5.02	CLIC3	-6.8	ENDOU	-4.5	PLN*	-6.1	CLIC3	-2.9