

Supplem. Table 1. Differentially expressed (DE) genes identified in IP wash and tumor samples, by NanoString

1A. List of DE genes (n=397) identified in IP wash samples (C2D4 v C2D1), q<0.1.

	log2Fold			log2Fold			
	Change*	pvalue	padj	Change	pvalue	padj	
SERPING1	4.13	3.02E-12	1.20E-09	CD3G	-2.80	7.43E-07	1.35E-05
IFI27	3.89	3.30E-12	1.20E-09	TNFSF10	2.89	8.54E-07	1.48E-05
SIGLEC1	3.72	6.51E-11	1.58E-08	CD86	2.59	1.23E-06	2.08E-05
CD163	4.24	7.57E-10	7.90E-08	CCL17	-2.96	1.25E-06	2.08E-05
MSR1	3.51	5.46E-10	7.90E-08	CYBB	3.24	1.65E-06	2.67E-05
TLR4	3.42	6.70E-10	7.90E-08	CFD	2.16	1.68E-06	2.67E-05
MERTK	3.10	6.45E-10	7.90E-08	C1QA	2.96	1.88E-06	2.86E-05
CD14	3.68	2.69E-09	2.18E-07	KLRC1	-2.49	1.88E-06	2.86E-05
THBS1	3.63	2.45E-09	2.18E-07	LRP1	2.30	2.10E-06	3.14E-05
CD68	3.29	4.51E-09	3.29E-07	CD40LG	-2.61	2.17E-06	3.17E-05
SLC11A1	2.84	6.97E-09	4.62E-07	IFITM2	2.97	2.25E-06	3.21E-05
LY96	2.68	1.06E-08	6.48E-07	CTSW	-2.64	2.29E-06	3.21E-05
IFIT1	3.56	2.97E-08	1.67E-06	SH2D1A	-2.62	2.53E-06	3.49E-05
CCR1	2.78	4.00E-08	2.08E-06	IFNGR1	3.28	2.67E-06	3.61E-05
IFIT2	2.98	4.80E-08	2.34E-06	CD7	-2.69	2.81E-06	3.73E-05
CDKN1A	2.70	5.80E-08	2.65E-06	ABCB1	-2.56	3.17E-06	4.14E-05
C3AR1	3.12	7.03E-08	3.02E-06	ISG15	2.83	3.31E-06	4.20E-05
FCGR3A	3.51	8.38E-08	3.40E-06	KLRG1	-2.56	3.34E-06	4.20E-05
LILRB3	2.84	9.13E-08	3.51E-06	CMKLR1	3.07	3.95E-06	4.89E-05
BST1	2.83	1.28E-07	4.68E-06	TLR1	2.58	4.11E-06	5.00E-05
NCR1	-3.02	1.39E-07	4.82E-06	LCK	-2.62	4.32E-06	5.16E-05
TIGIT	-2.83	1.65E-07	5.46E-06	CD160	-2.57	5.20E-06	6.12E-05
CD1C	-3.26	1.99E-07	6.33E-06	C1QB	2.81	5.81E-06	6.20E-05
CTSS	3.52	2.69E-07	7.58E-06	CD63	2.70	5.52E-06	6.20E-05
FCER1G	3.32	2.67E-07	7.58E-06	MARCO	2.55	5.38E-06	6.20E-05
IFNG	-2.72	2.80E-07	7.58E-06	ITGA1	-2.35	5.94E-06	6.20E-05
ICOS	-3.15	2.75E-07	7.58E-06	TNFRSF18	-2.40	5.92E-06	6.20E-05
CLEC7A	2.61	3.09E-07	8.05E-06	TBX21	-2.46	5.63E-06	6.20E-05
CD27	-2.98	3.45E-07	8.68E-06	TNFRSF4	-2.49	5.62E-06	6.20E-05
TLR2	2.51	3.92E-07	9.54E-06	CD1E	-2.99	5.71E-06	6.20E-05
LY86	2.65	4.13E-07	9.73E-06	S100A8	3.14	6.15E-06	6.26E-05
FOS	3.49	4.38E-07	9.98E-06	FPR2	2.58	6.17E-06	6.26E-05
PDGFC	4.19	6.79E-07	1.30E-05	KLRB1	-2.64	6.44E-06	6.44E-05
MX1	2.97	6.17E-07	1.30E-05	TNFRSF9	-2.63	7.50E-06	7.40E-05
LILRA5	2.91	6.51E-07	1.30E-05	LTB	-2.58	7.79E-06	7.58E-05
FCGR2A	2.88	6.71E-07	1.30E-05	HLA-B	2.97	1.03E-05	9.86E-05
IL1R2	2.62	6.66E-07	1.30E-05	PMCH	-2.69	1.09E-05	0.000103
IL2RB	-2.52	6.18E-07	1.30E-05	RORA	-2.46	1.15E-05	0.000107
LILRB2	2.97	7.56E-07	1.35E-05	DUSP6	2.26	1.17E-05	0.000108
CEBPB	2.80	7.22E-07	1.35E-05				

* Positive values indicate upregulation at C2D4

Suppl. Table 1A (cont'd)

	log2Fold Change	pvalue	padj		log2Fold Change	pvalue	padj
CCL22	-2.57	1.19E-05	0.000108				
HLA-A	2.99	1.27E-05	0.000113				
TNFSF13B	2.27	1.26E-05	0.000113	KIR_Activati ng_Subgrou			
BLK	-2.76	1.37E-05	0.000121	p_1	-2.59	9.56E-05	0.000582
NOD2	2.01	1.51E-05	0.000131	CD5	-2.29	9.68E-05	0.000584
TNFRSF1A	2.35	1.59E-05	0.000136	C2	2.18	0.000109	0.000655
DDX58	2.26	1.66E-05	0.000141	CD1B	-2.74	0.000119	0.000705
CSF3R	2.38	1.76E-05	0.000148	SPP1	2.81	0.000121	0.000713
CTSL	2.75	1.90E-05	0.000158	CD79B	-2.17	0.000125	0.000729
CD180	2.36	1.93E-05	0.000158	OAS3	2.34	0.000133	0.000773
CXCR3	-2.35	2.13E-05	0.000171	CD36	2.85	0.000137	0.000778
PPBP	-2.69	2.11E-05	0.000171	VEGFA	1.81	0.000137	0.000778
LY9	-2.19	2.23E-05	0.000177	EOMES	-2.30	0.000136	0.000778
GZMM	-2.36	2.39E-05	0.000187	CD247	-2.24	0.000143	0.000803
PECAM1	2.17	2.48E-05	0.000193	BCL2	-1.99	0.000161	0.000899
TLR6	2.04	2.56E-05	0.000196	CD84	2.22	0.000172	0.000953
LTBR	2.17	2.60E-05	0.000198	ITGB2	2.49	0.000177	0.000974
CD8B	-2.46	2.63E-05	0.000198	CMA1	-2.34	0.0002	0.001089
CD96	-2.46	3.23E-05	0.000241	SIGIRR	-1.91	0.000203	0.001098
CXCL5	-2.93	3.54E-05	0.000261	TCF7	-2.44	0.000225	0.001207
IL10	2.42	3.58E-05	0.000262	LYN	1.93	0.000227	0.00121
ENG	2.50	3.85E-05	0.000278	ITGAM	2.10	0.000253	0.001331
CTSH	2.28	4.46E-05	0.000319	CCR2	1.83	0.000252	0.001331
HCK	2.32	4.62E-05	0.000327	STAT2	1.96	0.000262	0.001368
CLEC4C	-2.50	4.88E-05	0.000342	BAX	2.07	0.000285	0.001455
BTLA	-2.07	5.46E-05	0.00038	CASP1	1.77	0.000284	0.001455
IRF7	2.10	5.62E-05	0.000387	IL11RA	-2.05	0.000282	0.001455
CTLA4	-2.33	5.76E-05	0.000393	NCAM1	-2.02	0.000298	0.001508
ZAP70	-2.31	5.88E-05	0.000398	KIR_Inhibiti ng_Subgrou			
CARD11	-2.46	5.94E-05	0.000398	p_1	-2.03	0.0003	0.00151
SLAMF1	-2.27	6.29E-05	0.000418	ITGA5	1.68	0.000311	0.001556
TNFRSF1B	2.14	7.07E-05	0.000461	IFI35	2.08	0.000344	0.001706
PSEN1	1.97	7.03E-05	0.000461	HLA-C	2.07	0.000353	0.00174
IL2RA	-2.09	7.18E-05	0.000464	IL1RN	2.43	0.000357	0.001748
LAMP2	1.77	7.39E-05	0.000465	CCR6	-2.36	0.00036	0.001751
SLAMF7	-1.87	7.33E-05	0.000465	CD53	2.42	0.000365	0.001764
CD8A	-2.24	7.29E-05	0.000465	CD2	-2.04	0.000371	0.00178
KLRD1	-2.03	7.72E-05	0.000482	KLRF1	-1.88	0.000416	0.001986
SYK	2.39	8.12E-05	0.000503	LGALS3	2.98	0.000425	0.002014
IL13RA1	2.43	9.43E-05	0.000578	FOXP3	-2.42	0.000429	0.002019
				IFI16	2.10	0.000435	0.002022
				CFP	1.79	0.000433	0.002022
				CD6	-2.37	0.000462	0.00212
				CR2	-2.44	0.000461	0.00212

* Positive values indicate upregulation at C2D4

Suppl. Table 1A (cont'd)

	log2Fold Change	pvalue	padj		log2Fold Change	pvalue	padj
UBC	2.49	0.000485	0.002214	KIR_Activ			
BIRC5	-1.99	0.000494	0.002238	ating_Subg			
CD79A	-1.94	0.000571	0.002573	roup_2	-1.98	0.001723	0.00629
CXCL16	1.72	0.000612	0.002741	PLA2G6	-1.62	0.001768	0.006421
PRKCD	1.63	0.000616	0.002743	LILRA4	-1.91	0.00191	0.006902
HLA-G	1.98	0.000631	0.002768	CD24	-1.85	0.002091	0.007519
PLAUR	1.66	0.000637	0.002768	JAM3	-1.86	0.002155	0.007712
MEF2C	1.59	0.000634	0.002768	CFB	1.57	0.002264	0.008062
IL32	-2.21	0.000637	0.002768	APP	1.43	0.002374	0.008372
TLR8	1.90	0.000662	0.002858	ITGA4	-1.61	0.002368	0.008372
MAPK11	-2.23	0.000675	0.002897	ANXA1	2.29	0.002438	0.008557
CCR4	-1.83	0.000722	0.003084	HLA-E	2.02	0.002466	0.008614
MRC1	2.05	0.000734	0.003116	IL1A	-1.91	0.002532	0.008801
KLRK1	-2.12	0.000741	0.003125	TREM2	1.74	0.002582	0.008907
CD74	2.40	0.00084	0.003523	SPN	-1.89	0.002587	0.008907
HAVCR2	1.77	0.000895	0.003733	TREM1	2.46	0.002679	0.009183
IFIH1	1.87	0.000944	0.003914	PSMB8	1.88	0.002839	0.009685
GNLY	-1.82	0.000987	0.00407	GZMK	-2.02	0.002866	0.009685
MAP3K1	1.51	0.001009	0.004117	MAGEC1	-2.05	0.002864	0.009685
CXCL13	-2.11	0.001007	0.004117	SPINK5	-1.86	0.002899	0.009752
BCL10	1.65	0.001026	0.004161	IL11	-1.98	0.002972	0.009951
STAT1	2.11	0.001123	0.004518	ADORA2			
CXCL9	-1.76	0.001133	0.004518	A	-1.76	0.003006	0.009974
CD3E	-2.11	0.001128	0.004518	CREBBP	-2.12	0.003003	0.009974
C3	1.72	0.001182	0.004688	IFITM1	2.10	0.003028	0.010001
CD81	2.24	0.00119	0.004695	IL18RAP	-1.78	0.003083	0.010092
ITGAE	-1.42	0.001207	0.004739	DOCK9	-1.84	0.003075	0.010092
CCL13	-2.29	0.00125	0.004881	LTK	-2.06	0.003194	0.010408
BAGE	-2.20	0.001264	0.004906	PTPRC	1.90	0.003268	0.010511
MYD88	1.98	0.001271	0.004907	STAT6	1.50	0.003248	0.010511
TLR9	-2.22	0.001299	0.00499	SH2D1B	-2.33	0.003262	0.010511
CCR7	-1.87	0.001405	0.00537	AIRE	-2.09	0.003331	0.010666
TXNIP	2.44	0.001459	0.005548	IL19	-2.02	0.003352	0.010685
CD164	2.13	0.001495	0.005651	MCAM	-1.91	0.003444	0.01093
IL23R	-2.13	0.001502	0.005651	TLR5	1.52	0.003479	0.010993
PAX5	-2.10	0.001567	0.005865	IL21	-1.93	0.003552	0.011177
SLAMF6	-1.71	0.001598	0.005923	PRG2	-1.72	0.00361	0.01131
CSF2	-2.32	0.001598	0.005923	IL3	-2.38	0.003627	0.011315
CX3CL1	-2.00	0.001654	0.0061	F12	-2.27	0.003656	0.011356
LAMP1	1.89	0.001682	0.006171	ITK	-1.85	0.003841	0.011881
				IL10RA	1.75	0.004069	0.012535
				IL13	-1.86	0.004211	0.012916
				HLA-DRA	2.11	0.004787	0.014528

* Positive values indicate upregulation at C2D4

Suppl. Table 1A (cont'd)

log2Fold				log2Fold			
	Change	pvalue	padj		Change	pvalue	padj
CCL19	-1.71	0.004797	0.014528				
S100A7	-1.78	0.004787	0.014528	CTAGE1	-1.86	0.009494	0.024751
IL7R	-1.82	0.004816	0.014528	HLA-			
TLR10	-1.91	0.004905	0.014735	DMA	1.45	0.009651	0.025049
LILRB1	1.72	0.00498	0.014899	IL6ST	1.44	0.009677	0.025049
ITGA2B	-2.10	0.005106	0.015214	CDK1	-1.45	0.009755	0.025163
BCL2L1	1.55	0.005249	0.015578	LCPI	1.81	0.0099	0.025447
THBD	1.40	0.005341	0.015786	PPARG	1.52	0.010292	0.026363
STAT3	1.51	0.005463	0.016082	FOXJ1	-1.71	0.010906	0.027837
HLA-				CD46	1.56	0.010983	0.027937
DRB3	1.88	0.005523	0.016088	SPANXB1	-2.14	0.011034	0.027967
IL23A	-1.40	0.005559	0.016088	CCL5	-1.68	0.011143	0.028146
RUNX3	-1.46	0.005561	0.016088	SERPINB2	1.47	0.01127	0.028369
IRF4	-1.48	0.005576	0.016088	FCER2	-1.48	0.011561	0.028903
KLRC2	-1.67	0.005552	0.016088	CD34	-1.85	0.011538	0.028903
IL7	-1.69	0.005598	0.016088	PSMB9	1.37	0.012036	0.029987
HLA-				CCRL2	1.34	0.012082	0.029999
DPA1	1.90	0.005671	0.016117	CD4	1.38	0.012457	0.030707
IL22RA2	-1.76	0.005674	0.016117	VCAM1	1.22	0.012535	0.030707
TNFSF11	-1.93	0.00563	0.016117	MS4A1	-1.58	0.012443	0.030707
MAP2K1	1.21	0.005843	0.016532	FUT5	-1.66	0.012505	0.030707
CXCR2	1.69	0.005937	0.016732	GZMH	-1.47	0.012636	0.030772
IGF2R	1.50	0.00598	0.016791	XCL2	-1.55	0.012646	0.030772
HLA-				TPSAB1	-1.62	0.012689	0.030775
DQB1	1.84	0.006106	0.017079	NEFL	-1.81	0.013544	0.032739
ICAM2	1.28	0.006156	0.017153	CHIT1	-1.56	0.013662	0.032914
TNFRSF				IL5RA	-1.57	0.014339	0.034433
13C	-2.24	0.006378	0.017704	PSMD7	1.47	0.014482	0.034662
HLA-				IL12A	-1.58	0.014758	0.035208
DMB	1.59	0.007019	0.019398	LRRN3	-1.55	0.015151	0.036028
NT5E	-1.69	0.007042	0.019398	TNFRSF1			
STAT4	-1.27	0.007281	0.019981	7	-1.58	0.016213	0.038427
CCL14	-1.68	0.00737	0.020151	PYCARD	1.77	0.016721	0.039375
IRF8	1.34	0.007578	0.02064	CASP10	1.26	0.016706	0.039375
CD3D	-1.81	0.007757	0.02105	EWSR1	1.52	0.017246	0.040481
TNFRSF				ATG7	1.33	0.018232	0.042473
11A	1.42	0.007983	0.021583	MAPK3	1.17	0.018269	0.042473
BATF	-1.22	0.008133	0.021904	TXK	-1.49	0.01818	0.042473
IL4	-1.73	0.008161	0.021904	HLA-			
HLA-				DRB4	1.51	0.018557	0.043005
DPB1	1.89	0.008416	0.022463	KIR_Inhibi			
KIT	-1.69	0.008462	0.022463	ting_Subgr			
C8G	-1.80	0.008433	0.022463	oup_2	-1.83	0.018753	0.043321
IRF5	1.25	0.008645	0.022866	IL1B	-1.29	0.018908	0.043541
BST2	1.22	0.008721	0.022982	HLA-DOB	-1.31	0.018981	0.043572
IL13RA2	-1.89	0.009245	0.024275	C4BPA	-1.83	0.019528	0.044687
TTK	-1.60	0.00933	0.024411				

* Positive values indicate upregulation at C2D4

Suppl. Table 1A (cont'd)

	log2Fold				log2Fold		
	Change	pvalue	padj		Change	pvalue	padj
NFKBIA	1.57	0.019594	0.044698	TGFB1	1.23	0.032973	0.066863
CXCL12	1.96	0.020354	0.045916	JAK2	1.00	0.033339	0.067416
PSMB7	1.24	0.020323	0.045916	IL17RA	0.95	0.033837	0.068234
IFNAR2	1.17	0.020429	0.045916	CD19	-1.21	0.034405	0.069
MICB	1.09	0.020484	0.045916	CEACAM			
CXCR5	-1.59	0.020409	0.045916	1	-1.33	0.03435	0.069
ATM	-1.71	0.020505	0.045916	IL12RB2	-1.35	0.034816	0.069633
ALCAM	1.36	0.02125	0.047438	CLU	-1.74	0.034993	0.069795
S100A12	1.26	0.021396	0.047618	PTGDR2	-1.31	0.035535	0.070683
POU2AF1	-1.58	0.021498	0.0477	TLR7	1.58	0.035948	0.071309
IL34	-1.65	0.022277	0.049279	IL3RA	-1.16	0.036162	0.07154
THY1	-1.18	0.022399	0.049399	CTAG1B	-1.28	0.036412	0.071793
C1R	1.19	0.022512	0.0495	LTF	-1.41	0.036487	0.071793
BLNK	1.24	0.02263	0.049609	TANK	1.15	0.036613	0.071848
CXCR6	-1.74	0.02403	0.052521	CYFIP2	-1.11	0.037509	0.073409
MAGEC2	-1.52	0.024282	0.052912	FCGR2B	1.09	0.038069	0.074306
GPI	1.09	0.024393	0.052997	TNFRSF8	1.29	0.038728	0.075391
CRP	-1.57	0.024554	0.053188	TOLLIP	0.92	0.039024	0.075765
LTA	-1.49	0.024973	0.053935	MME	-1.31	0.042062	0.081446
SMPD3	-1.36	0.025071	0.053987	AICDA	-1.49	0.042362	0.08181
FLT3LG	-1.31	0.025835	0.055433	PSMB10	1.06	0.043028	0.082442
MS4A2	-1.51	0.025894	0.055433	CD58	1.02	0.042951	0.082442
RORC	-1.29	0.026288	0.056003	CCR3	-1.22	0.043024	0.082442
CCL21	-1.43	0.026314	0.056003	CD22	-1.15	0.043615	0.083349
CD97	1.42	0.026982	0.057258	TNFRSF1			
DUSP4	-1.22	0.027378	0.05789	4	1.01	0.044055	0.08397
ICAM4	-1.41	0.027438	0.05789	CD38	0.98	0.044648	0.08422
CSF1R	1.12	0.028407	0.059761	PRF1	-1.24	0.044585	0.08422
TNFRSF1				IL1RAPL			
OC	1.34	0.028739	0.060286	2	-1.30	0.044579	0.08422
CASP3	1.04	0.029137	0.060945	MAGEA3	-1.49	0.044442	0.08422
IL6R	1.08	0.030122	0.062826	MAP2K2	1.20	0.044946	0.084564
CTCFL	-1.52	0.030612	0.063665	LAG3	-1.04	0.045173	0.084771
CD47	1.48	0.030864	0.064008	GZMA	-1.40	0.045695	0.085531
PRAME	-1.30	0.030989	0.064085	LAIR2	1.09	0.046228	0.086308
KIR3DL2	-1.44	0.031308	0.064199	TRAF2	-1.00	0.047251	0.087994
CCL1	-1.54	0.031147	0.064199	TAP1	1.03	0.047824	0.088833
DMBT1	-1.55	0.031274	0.064199	IL27	1.48	0.048099	0.089118
IL22	-1.60	0.032284	0.066015	S100B	-1.50	0.048512	0.089656
NFATC2	-1.28	0.032669	0.066616	KIR3DL1	-1.37	0.049535	0.091314
C4B	1.69	0.032944	0.066863	FCER1A	-1.16	0.049744	0.091468

* Positive values indicate upregulation at C2D4

1B. List of DE genes (n=411) identified in IP wash samples (C3D11 v C3D1), q<0.1

	log2Fold Change*	pvalue	padj		log2Fold Change*	pvalue	padj
LILRA5	3.50	8.96E-08	6.54E-05	THBS1	2.88	7.15E-05	0.001273
LILRB2	3.24	2.38E-07	8.7E-05	HAMP	-2.36	7.33E-05	0.001275
SLC11A1	2.50	4.64E-06	0.000548	CTSS	2.44	7.54E-05	0.00128
TNFRSF1B	2.40	5.08E-06	0.000548	BST1	2.51	8.33E-05	0.001382
CFP	2.24	6.07E-06	0.000548	ISG15	2.29	8.55E-05	0.001387
TNFSF13B	2.14	9.75E-06	0.000548	CD1C	-2.40	9.04E-05	0.001435
SAA1	-2.82	9.35E-06	0.000548	CEACAM8	-2.70	9.31E-05	0.001447
CCL22	-2.85	2.25E-06	0.000548	IFITM2	2.22	0.000107	0.001622
BAGE	-2.97	9.46E-06	0.000548	CTSH	1.96	0.000112	0.001669
S100A7	-3.09	5.59E-06	0.000548	CD36	2.19	0.000115	0.001681
CCL17	-3.15	9.66E-06	0.000548	ITGAM	2.30	0.000118	0.001694
CXCL12	-3.89	8.18E-06	0.000548	IL11	-3.43	0.000126	0.001764
CCL25	-4.17	7.09E-06	0.000548	MAP3K1	1.91	0.000128	0.001765
CLEC7A	2.80	1.79E-05	0.00063	XCL2	-2.09	0.000131	0.001765
FCER1G	2.66	1.72E-05	0.00063	CCR2	2.31	0.000136	0.001802
CSF3R	2.65	1.3E-05	0.00063	MYD88	2.07	0.000141	0.001836
FCGR2A	2.46	1.59E-05	0.00063	IRF7	1.91	0.000152	0.001951
CCR1	2.30	1.81E-05	0.00063	CD68	2.16	0.000155	0.001951
DOCK9	-2.46	1.64E-05	0.00063	CXCL14	-2.70	0.000158	0.001961
GAGE1	-2.69	1.39E-05	0.00063	CD180	2.43	0.000169	0.002061
IL13RA2	-3.81	1.58E-05	0.00063	LILRB1	1.78	0.000175	0.002089
TREM1	2.80	2.16E-05	0.000662	LGALS3	1.78	0.000188	0.002143
CD86	2.63	2.18E-05	0.000662	ZAP70	-2.02	0.000184	0.002143
TCF7	-2.73	2.04E-05	0.000662	KLRG1	-2.06	0.000186	0.002143
CD160	-3.18	2.89E-05	0.000845	SYK	1.98	0.000194	0.002143
LCK	-2.25	3.04E-05	0.000853	IL1RL1	-2.34	0.000193	0.002143
S100A8	2.71	3.27E-05	0.000883	SERPING1	2.28	0.000197	0.002144
FCGR3A	2.54	3.97E-05	0.000945	TLR1	2.20	0.000213	0.002251
CEBPB	2.51	3.67E-05	0.000945	CD8A	-2.01	0.00021	0.002251
CEACAM1	-2.65	4.14E-05	0.000945	CD24	-2.21	0.000229	0.002387
CCL14	-2.75	3.83E-05	0.000945	IFIT1	2.36	0.000246	0.002517
CD1E	-3.58	4.09E-05	0.000945	SIGLEC1	2.22	0.000248	0.002517
MASP1	-2.64	4.55E-05	0.001006	C3AR1	2.71	0.000265	0.002651
TIGIT	-2.31	4.98E-05	0.001069	LYN	2.17	0.000272	0.002657
FPR2	2.51	5.6E-05	0.001077	IL2RA	-2.14	0.000277	0.002657
PECAM1	2.40	5.33E-05	0.001077	ITGA2	-2.26	0.000275	0.002657
NOD2	2.38	5.34E-05	0.001077	CXCR5	-2.33	0.000303	0.002875
CTAGE1	-2.89	5.61E-05	0.001077	MARCO	2.29	0.000315	0.002951
CYBB	2.47	6.13E-05	0.001147	IFI27	2.28	0.000346	0.003185
TNFSF10	2.48	6.36E-05	0.001161	NEFL	-2.28	0.000352	0.003185
				IL4	-3.00	0.000353	0.003185

* Positive values indicate upregulation at C3D11

Suppl. Table 1B (cont'd)

log2Fold				log2Fold			
	Change*	pvalue	padj		Change*	pvalue	padj
CD14	2.64	0.000364	0.003244	HLA-A	1.94	0.00098	0.005862
CTSW	-1.90	0.000375	0.003298	MASP2	-2.89	0.001006	0.005968
IL1RN	1.83	0.000388	0.003376	CHIT1	-2.25	0.001015	0.005976
IFNG	-2.11	0.000407	0.003496	GPI	1.65	0.001035	0.005996
IFNA7	-2.19	0.000419	0.003557	LTA	-2.27	0.001033	0.005996
PLAUR	1.83	0.00043	0.003609	THBD	2.15	0.001054	0.006056
HCK	1.80	0.000443	0.003678	LILRB3	2.01	0.00111	0.006236
TNFRSF13				CCR5	1.93	0.001111	0.006236
C	-2.40	0.000453	0.003714	THY1	-1.85	0.001108	0.006236
CTLA4	-2.20	0.000465	0.003728	PYCARD	1.64	0.001196	0.006664
IFNA2	-2.41	0.00046	0.003728	TLR4	2.04	0.001208	0.006681
IL5RA	-2.98	0.00047	0.003728	CD63	1.67	0.001222	0.006706
ITGB2	2.05	0.000496	0.003872	GZMK	-1.96	0.001291	0.007031
ITGA1	-2.04	0.000499	0.003872	CD27	-2.07	0.001332	0.007205
LRRN3	-2.41	0.000551	0.004231	MSR1	2.12	0.001353	0.00723
CDKN1A	1.72	0.000586	0.004459	C6	-2.00	0.001357	0.00723
VEGFA	2.16	0.00062	0.004574	CXCL16	1.90	0.00139	0.00731
TPTE	-2.36	0.000616	0.004574	PAX5	-2.14	0.001392	0.00731
TMEFF2	-2.46	0.000614	0.004574	IFIT2	2.04	0.001458	0.007551
CD53	2.06	0.000631	0.004587	KLRC2	-2.00	0.001451	0.007551
HLA-B	1.91	0.000647	0.004587	BTK	1.52	0.001524	0.007675
MCAM	-2.48	0.000643	0.004587	DMBT1	-2.28	0.0015	0.007675
IL22	-2.65	0.000639	0.004587	IFNL1	-2.37	0.001513	0.007675
PSEN1	1.84	0.000675	0.004737	IL17F	-2.46	0.001514	0.007675
PRKCD	1.89	0.000686	0.004768	STAT2	1.66	0.001602	0.008008
CD3EAP	-2.12	0.000703	0.004838	ICAM4	-1.97	0.001639	0.008142
KLRD1	-1.71	0.000715	0.004878	LY86	2.07	0.001725	0.008496
OAS3	1.80	0.000731	0.004943	AMBP	-2.32	0.001734	0.008496
CTSL	1.73	0.000746	0.00496	SELPLG	2.02	0.001794	0.008658
CFD	1.68	0.000747	0.00496	NCR1	-2.09	0.001794	0.008658
DUSP6	1.83	0.000761	0.005003	MBL2	-2.39	0.001803	0.008658
CCL1	-2.45	0.000776	0.005055	CXCR6	-1.69	0.001858	0.008809
IFNGR1	2.00	0.000785	0.005069	TNFSF11	-2.13	0.00185	0.008809
CT45A1	-2.00	0.000792	0.005074	IFNL2	-2.02	0.001985	0.00935
CD79A	-2.02	0.000838	0.005282	KLRK1	-1.66	0.002015	0.00937
EPCAM	-2.08	0.000839	0.005282	CD40LG	-2.26	0.002004	0.00937
TLR9	-2.28	0.00086	0.005364	HLA-G	1.70	0.002047	0.009453
CCR7	-2.31	0.000871	0.005385	TLR2	1.66	0.002059	0.009453
FOS	2.17	0.000928	0.005695	MNX1	-2.69	0.002103	0.009577
LRP1	2.04	0.000964	0.00586	PMCH	-2.75	0.002112	0.009577
NCF4	1.84	0.000971	0.00586				

* Positive values indicate upregulation at C3D11

Suppl. Table 1B (*cont'd*)

	log2Fold Change*	pvalue	padj		log2Fold Change*	pvalue	padj
CD5	-1.76	0.002166	0.009759	IL5	-2.25	0.004309	0.015574
IL22RA2	-2.20	0.002185	0.009786	HLA-DOB	-1.60	0.004349	0.015639
ARG2	-1.96	0.00222	0.009884	CCR6	-1.93	0.004403	0.015757
PIK3CG	1.74	0.002261	0.009941	AKT3	-1.42	0.004546	0.016189
IL13	-2.20	0.00225	0.009941	CXCL5	-2.18	0.004695	0.016638
TLR6	1.65	0.002298	0.010045	UBC	1.66	0.004825	0.016935
IFIH1	1.56	0.002403	0.010441	LCN2	-1.78	0.004823	0.016935
CD83	1.95	0.00246	0.010625	CD97	1.64	0.004902	0.01696
PLA2G1B	-2.22	0.002481	0.010653	LCP1	1.60	0.004896	0.01696
IFNB1	-2.23	0.002504	0.01069	KIT	-2.09	0.004902	0.01696
MAGEA12	-2.12	0.002538	0.010773	ITGA6	-2.14	0.004928	0.01697
S100B	-1.83	0.002598	0.010902	KIR_Activati ng_Subgrou p_2	-2.07	0.00502	0.017206
IL1RAPL2	-2.06	0.002591	0.010902	FCGR2B	1.53	0.005068	0.017289
TNFRSF18	-1.81	0.002634	0.010987	IRF5	1.44	0.005137	0.017421
SPANXB1	-2.07	0.002687	0.011147	CFI	-1.71	0.005179	0.017421
TNFRSF1A	1.61	0.002758	0.011374	SELE	-2.25	0.005175	0.017421
MAGEA3	-2.18	0.002839	0.011644	CD96	-1.70	0.005324	0.017827
RIPK2	1.75	0.002881	0.011671	FOXJ1	-1.98	0.00535	0.017832
IL26	-2.28	0.002894	0.011671	HLA-DRB3	1.44	0.005475	0.018097
CTCF	-2.29	0.00289	0.011671	CD7	-1.81	0.005479	0.018097
CD79B	-1.74	0.003015	0.012092	ABCB1	-1.52	0.005684	0.018692
STAT3	1.50	0.003077	0.012276	PTPRC	1.70	0.005812	0.019025
CR2	-2.15	0.003111	0.012341	MAPK11	-2.62	0.005843	0.01904
HLA-DRA	1.75	0.003174	0.012486	HLA-DMB	1.58	0.00595	0.019221
TNFSF4	-1.69	0.003181	0.012486	NT5E	-1.52	0.005951	0.019221
ITGA5	1.41	0.003338	0.01303	LAMP2	1.63	0.006033	0.0194
HAVCR2	1.58	0.003376	0.01304	CD84	1.62	0.006146	0.019677
SH2D1A	-1.90	0.003366	0.01304	FEZ1	-1.62	0.006254	0.019936
ICOS	-1.89	0.003422	0.013146	MPPED1	-2.26	0.006305	0.020012
BCL10	1.65	0.003476	0.013223	HLA-E	1.53	0.006381	0.020165
ICAM2	1.46	0.003478	0.013223	CD22	-1.50	0.006663	0.020967
TLR5	1.45	0.003533	0.013365	HLA-DPA1	1.61	0.006715	0.021037
NLRP3	1.74	0.003598	0.013409	ITGAX	1.47	0.007063	0.021894
CD74	1.73	0.0036	0.013409	IL11RA	-1.90	0.007078	0.021894
FCER1A	-1.79	0.003582	0.013409	KIR3DL2	-1.95	0.007067	0.021894
MX1	1.64	0.003664	0.013576	MAP2K1	1.25	0.007171	0.021995
MEF2C	1.75	0.00371	0.01368	IFNA8	-2.37	0.007147	0.021995
CCRL2	1.73	0.003933	0.014426	ENG	1.77	0.007297	0.022288
CD6	-1.73	0.004089	0.014926	CD247	-1.48	0.00739	0.022477
BTLA	-1.52	0.004302	0.015574	PSMB10	1.40	0.007524	0.022791

* Positive values indicate upregulation at C3D11

Suppl. Table 1B (*cont'd*)

	log2Fold Change*	pvalue	padj		log2Fold Change*	pvalue	padj
CREBBP	-1.78	0.007589	0.022894	IL18RAP	-2.02	0.013969	0.03616
LTK	-1.96	0.007689	0.023099	TNFRSF14	1.33	0.014412	0.037132
IGLL1	-2.00	0.008244	0.024665	CD19	-1.51	0.014446	0.037132
ARG1	-2.36	0.008294	0.024711	DDX58	1.54	0.014818	0.037954
IL6	-1.83	0.008427	0.025008	PDGFRB	-1.38	0.014879	0.037979
IFI16	1.38	0.008645	0.025446	HLA-DPB1	1.44	0.014978	0.038096
F2RL1	-1.69	0.008636	0.025446	IFITM1	1.56	0.015208	0.038547
TLR10	-1.74	0.008702	0.025513	IL13RA1	1.26	0.015699	0.039654
HLA-C	1.55	0.0088	0.025696	ATG7	1.26	0.015879	0.039697
CCL28	-2.06	0.008909	0.025911	CCL23	-1.67	0.015844	0.039697
STAT6	1.29	0.008952	0.025932	ITGB3	-2.02	0.015847	0.039697
ITK	-1.55	0.009092	0.026235	TNFRSF4	-1.57	0.015995	0.039852
POU2AF1	-2.24	0.009186	0.026402	JAK2	1.15	0.016071	0.039903
SPACA3	-2.05	0.009356	0.026782	IL24	-1.69	0.016314	0.040234
ANXA1	1.60	0.009527	0.027109	MAGEC2	-1.76	0.016306	0.040234
IL7	-1.68	0.009544	0.027109	S100A12	3.68	0.016392	0.04029
KLRC1	-1.37	0.009594	0.027146	SPN	-1.93	0.016548	0.040538
IFI35	1.98	0.00966	0.027228	STAT1	1.37	0.016616	0.040567
IL6R	1.55	0.01011	0.028234	IL10RA	1.65	0.016735	0.040703
HLA-DMA	1.37	0.010087	0.028234	HLA-DQB1	-1.42	0.016783	0.040703
CCL13	-1.78	0.010133	0.028234	SYT17	-2.25	0.016842	0.040712
CMKLR1	1.34	0.010218	0.028362	RELA	1.44	0.016936	0.040804
TPSAB1	-1.65	0.010605	0.029213	TXNIP	1.50	0.017206	0.041275
PPBP	-1.85	0.010594	0.029213	CSF2RB	1.13	0.017282	0.041275
BAX	1.40	0.010987	0.030151	CD2	-1.34	0.017302	0.041275
LAMP1	1.21	0.011205	0.03056	IL23R	-1.77	0.01757	0.041778
AIRE	-1.91	0.011219	0.03056	ULBP2	-1.50	0.018192	0.042565
CD8B	-1.42	0.011411	0.030966	IL12B	-1.67	0.018183	0.042565
CD47	1.39	0.011564	0.031266	CX3CL1	-1.68	0.018142	0.042565
BCL2	-1.48	0.011771	0.031708	CD1A	-1.72	0.018037	0.042565
CD28	-1.52	0.012065	0.03238	LILRA4	-1.82	0.017997	0.042565
TARP	-1.28	0.012575	0.033614	TLR8	1.29	0.018697	0.043607
TTK	-1.46	0.012617	0.033614	EOMES	-1.50	0.018924	0.043995
IL21	-2.07	0.012733	0.033799	CXCL9	-1.56	0.01903	0.044101
PSMB9	1.21	0.012933	0.034206	MAPK3	1.12	0.019151	0.044102
IRF8	1.44	0.013064	0.034428	CCL11	-2.00	0.019103	0.044102
EWSR1	1.43	0.013382	0.035139	ELANE	-1.60	0.019316	0.044342
BCL2L1	1.18	0.013769	0.035921	CSF1R	1.19	0.019609	0.044874
IL17B	-2.06	0.013778	0.035921	TBX21	-1.37	0.020046	0.045731
IL12RB2	-1.65	0.013844	0.035965	IRF4	-1.17	0.020309	0.046185

* Positive values indicate upregulation at C3D11

Suppl. Table 1B (cont'd)

	log2Fold Change*	pvalue	padj		log2Fold Change*	pvalue	padj
CLEC4A	1.25	0.020377	0.046197	IL23A	-1.43	0.031849	0.06335
A2M	-1.43	0.02088	0.04719	CRP	-1.74	0.032174	0.063823
IL17RA	1.15	0.021043	0.047411	TGFB1	1.22	0.032758	0.064806
APOE	-1.68	0.021697	0.048735	NFKBIA	1.32	0.032982	0.065073
CEACAM6	-2.01	0.021981	0.049221	MAPK14	0.97	0.033341	0.065426
IL17RB	-2.11	0.022054	0.049233	GZMM	-1.33	0.033277	0.065426
CD1D	1.29	0.022189	0.049385	SYCP1	-1.63	0.034169	0.066726
IFNAR2	1.08	0.022806	0.05045	CCL24	-1.74	0.034186	0.066726
ATM	-2.02	0.022754	0.05045	NOD1	1.89	0.034301	0.066772
CXCR4	1.69	0.023132	0.051017	DPP4	-1.32	0.034732	0.067254
IL2RB	-1.22	0.023744	0.052208	CCL16	-1.84	0.034709	0.067254
PSEN2	-1.33	0.023957	0.052518	CCL21	-1.67	0.034887	0.067374
CD164	1.22	0.024107	0.052688	CXCR3	-1.17	0.035559	0.068491
COL3A1	-1.72	0.025351	0.055242	CXCL1	-1.30	0.03568	0.068544
C8B	-1.62	0.025908	0.056159	TNFRSF1			
C8A	-1.69	0.025926	0.056159	7	-1.42	0.035785	0.068565
TNFRSF10				MUC1	-1.56	0.036043	0.068878
C	1.28	0.026422	0.056875	TNFRSF9	-1.53	0.036865	0.070264
SPINK5	-1.77	0.026393	0.056875	RELB	1.07	0.037417	0.071131
ROPN1	-1.95	0.02649	0.056875	BATF	-1.15	0.038267	0.072558
IKBK	1.27	0.026752	0.057271	PTGDR2	-1.52	0.038548	0.072902
KIR3DL3	-1.66	0.027197	0.058053	SSX1	-1.44	0.03872	0.073038
IL1A	-1.60	0.027743	0.059046	BIRC5	-1.45	0.039092	0.073549
CCL20	-1.33	0.027906	0.059047	NFKB2	1.03	0.039391	0.073921
CXCL13	-1.58	0.027867	0.059047	HSD11B1	-1.29	0.039625	0.074169
BCL6	1.31	0.028064	0.059175	CD40	1.12	0.040962	0.076477
CCL26	-1.39	0.028128	0.059175	PSMD7	1.23	0.041463	0.077214
TANK	1.37	0.028374	0.05935	C8G	-1.51	0.041845	0.077728
TLR7	1.35	0.028337	0.05935	PSMB8	0.99	0.042513	0.078767
SSX4	-1.68	0.028776	0.060018	CD81	1.07	0.044033	0.081171
ITGAE	-1.14	0.028873	0.060049	IL3	-1.55	0.043956	0.081171
CD46	1.22	0.029078	0.060303	F12	-1.62	0.044213	0.081299
CD163	1.31	0.029239	0.060467	TYK2	1.28	0.0445	0.081551
MICB	1.20	0.029594	0.061026	CD48	1.20	0.044686	0.081551
CD55	-1.38	0.029924	0.061533	ICAM1	1.08	0.044797	0.081551
AMICA1	1.16	0.030053	0.061625	RAG1	-1.50	0.044578	0.081551
ZNF205	-1.60	0.030209	0.061772	PSMB7	0.92	0.045387	0.082419
CCL15	-1.80	0.030418	0.062026	IGF2R	1.04	0.046228	0.083738
IGF1R	-1.13	0.030504	0.062029	CD58	1.02	0.046388	0.08382
CD3D	-1.27	0.031193	0.062941	ITGA2B	-1.69	0.046639	0.084065
IL7R	-1.28	0.031384	0.062941	CD3E	-1.24	0.047148	0.084773
CD34	-1.58	0.031365	0.062941	IFNA17	-1.43	0.047339	0.084908
IL34	-1.60	0.031363	0.062941	RORA	-1.16	0.048285	0.086392
NOS2A	-1.75	0.031225	0.062941	BLK	-1.50	0.048631	0.086799
MAPK1	1.17	0.031527	0.063054	CASP1	1.05	0.049204	0.087606
IL32	-1.25	0.031778	0.06335	CD44	1.18	0.049378	0.087702

* Positive values indicate upregulation at C3D11

1C. List of DE genes identified in IP wash samples (C2D1 v C1D1, p<0.05)

	log2FoldChange*	pvalue	padj		log2FoldChange*	pvalue	padj
CCL18	2.362746	0.000116	0.085035	TNFRSF1A	-0.76575	0.021718	0.493789
TNFSF10	-1.40053	0.000454	0.165849	CCL13	1.20673	0.021791	0.493789
CD163	-1.40386	0.001228	0.23964	CYBB	-0.92241	0.022322	0.493789
PDGFC	-1.81903	0.001313	0.23964	CR2	1.032712	0.025269	0.542538
C3	-1.44217	0.002933	0.345052	TLR4	-0.83353	0.026112	0.542819
DPP4	1.023591	0.003161	0.345052	CCL23	0.993382	0.02722	0.542819
FCGR3A	-1.29092	0.003758	0.345052	MSR1	-0.83711	0.029689	0.542819
SERPING1	-1.10618	0.003781	0.345052	IFI27	-0.96597	0.030928	0.542819
CCL19	1.206259	0.005556	0.424631	IFITM2	-0.90907	0.031278	0.542819
NCR1	1.158027	0.005931	0.424631	HLA-A	-0.90752	0.031844	0.542819
IFIT2	-1.25362	0.006625	0.424631	USP9Y	1.055435	0.03245	0.542819
MX1	-1.07126	0.00698	0.424631	C2	-0.82941	0.033368	0.542819
MARCO	-1.1296	0.00801	0.44979	CD46	-0.78241	0.034325	0.542819
C4B	-1.48578	0.009179	0.478632	IGF2R	-0.66215	0.034695	0.542819
CCL17	0.895152	0.010356	0.49302	APP	-0.66456	0.03553	0.542819
KIR_Inhibiting_				IFI16	-0.81594	0.036088	0.542819
Subgroup_1	1.086075	0.011444	0.49302	NT5E	0.74658	0.037479	0.542819
IL5	1.198787	0.011661	0.49302	IFIH1	-0.78148	0.037481	0.542819
IFIT1	-1.12952	0.012361	0.49302	TLR1	-0.84047	0.038095	0.542819
FN1	-1.16608	0.012832	0.49302	CD40	-0.60901	0.038333	0.542819
CEBPB	-0.89286	0.014309	0.493789	ISG15	-0.7584	0.038643	0.542819
S100A8	-0.99948	0.015498	0.493789	LILRB2	-0.7478	0.038667	0.542819
IL12B	1.214228	0.015651	0.493789	CD3G	0.806443	0.040042	0.551522
CD68	-0.91713	0.016446	0.493789	STAT1	-0.80816	0.042842	0.579058
IFITM1	-1.12828	0.017346	0.493789	FOXP3	1.002639	0.043777	0.579058
FCER1G	-0.87889	0.018281	0.493789	NCAM1	0.704666	0.044945	0.579058
CD1C	0.914589	0.018709	0.493789	CD79B	0.649191	0.045775	0.579058
IL2RA	0.757874	0.018843	0.493789	IFNG	0.743265	0.047385	0.579058
DDX58	-1.0192	0.02031	0.493789	PLA2G6	0.663122	0.047634	0.579058
CTSS	-1.01816	0.020645	0.493789	EPCAM	-1.01309	0.048533	0.579058
CD63	-0.86938	0.02129	0.493789	MPPED1	0.957639	0.048912	0.579058
				IL13RA2	0.969963	0.04918	0.579058

* Positive values indicate upregulation at C2D1

1D.List of DE genes identified in tumor tissue (interval vs baseline, $p < 0.05$; genes in bold $q < 0.1$).

	log2Fold Change*	pvalue	padj		log2Fold Change*	pvalue	padj
CHIT1	2.61	5.56E-07	0.000249	MIF	-1.31	0.010035	0.101949
CFD	2.55	1.83E-06	0.000409	CCL18	2.76	0.01056	0.102884
CD36	3.22	4.66E-06	0.000695	CCRL2	1.11	0.010588	0.102884
FOXJ1	-2.13	1.3E-05	0.001456	ECSIT	-1.01	0.010994	0.104494
S100A12	2.85	1.64E-05	0.001469	CD209	1.39	0.011253	0.104494
IL1RN	2.20	2.67E-05	0.001989	MUC1	-1.20	0.011712	0.104494
CCL13	1.80	4.62E-05	0.002949	MAP3K1	-0.99	0.011849	0.104494
ITGAX	1.54	7.64E-05	0.003952	ITK	1.27	0.011922	0.104494
CFB	-1.97	7.96E-05	0.003952	PVR	-0.99	0.01367	0.117505
S100A8	2.46	8.94E-05	0.003996	PECAM1	1.32	0.014075	0.118706
CXCR2	2.44	0.000126	0.005101	C4B	-1.47	0.015036	0.124465
SAA1	3.36	0.000144	0.005377	SLC11A1	1.10	0.015786	0.1283
MME	1.81	0.000216	0.00742	RPS6	-1.24	0.016698	0.131505
IL7R	1.68	0.000415	0.013248	CD34	1.07	0.016769	0.131505
GZMB	1.53	0.000473	0.013831	HCK	0.92	0.02035	0.153508
CXCL2	2.42	0.000495	0.013831	IL11RA	-0.97	0.02036	0.153508
CDH1	-2.04	0.000583	0.015335	CCL19	1.47	0.020644	0.153508
PPARG	1.69	0.001155	0.028579	LCN2	-1.26	0.020949	0.153508
VEGFC	1.37	0.001215	0.028579	MFGE8	-1.01	0.021768	0.156944
THBD	1.32	0.001347	0.030102	A2M	0.95	0.022424	0.159104
EPCAM	-1.73	0.001621	0.034513	TREM1	1.13	0.023374	0.163255
CMKLR1	1.16	0.002618	0.053188	ELK1	-1.06	0.024878	0.168558
CR1	1.35	0.003022	0.058728	TNFRSF1B	0.76	0.025179	0.168558
SPA17	-1.28	0.003612	0.065258	CTSS	1.17	0.026038	0.168558
RUNX1	-1.40	0.00375	0.065258	TNFRSF10C	1.16	0.026083	0.168558
F2RL1	-1.28	0.003875	0.065258	CYFIP2	-0.91	0.026449	0.168558
CSF2RB	1.35	0.003942	0.065258	MRC1	0.96	0.026557	0.168558
IL1RAP	-1.09	0.004438	0.070845	MAPK8	-0.88	0.026773	0.168558
LAIR2	1.12	0.00583	0.087627	TNFAIP3	0.92	0.027794	0.172555
MARCO	1.60	0.005881	0.087627	ITGAM	0.87	0.028766	0.174285
CX3CR1	-1.38	0.00688	0.095925	ITGA2	-0.93	0.028853	0.174285
TLR8	1.14	0.006964	0.095925	CCL14	1.44	0.029821	0.177735
PRF1	1.33	0.007082	0.095925	APP	-1.13	0.030612	0.179861
PPBP	2.68	0.0075	0.097858	BAX	-0.96	0.031041	0.179861
TLR4	1.02	0.007745	0.097858	ENG	0.81	0.031385	0.179861
TAB1	-1.02	0.007881	0.097858	ILF3	-1.04	0.032794	0.185557
CD24	-1.75	0.008255	0.098091	ITGB3	0.92	0.033813	0.188931
CFI	-1.23	0.008785	0.098091	TGFB2	-0.86	0.035427	0.194316
CDH5	1.18	0.009176	0.098091	ZNF205	-0.93	0.035646	0.194316
NFATC4	-0.85	0.009189	0.098091	IRF3	-0.82	0.036463	0.196371
PDGFC	-1.11	0.009199	0.098091	RELA	-1.01	0.03933	0.209293
CD200	-0.98	0.009424	0.098091	TNFRSF1A	-0.80	0.040765	0.214378
LILRB2	0.95	0.009436	0.098091	CXCL3	1.26	0.041506	0.215736
				STAT2	-0.92	0.0421	0.216306
				BMI1	-0.93	0.042643	0.216609
				AXL	-0.73	0.043305	0.217498
				TP53	-0.74	0.044489	0.218613
				ITGA4	-0.79	0.044505	0.218613
				BIRC5	-1.19	0.047821	0.232348

* Positive values indicate upregulation at interval debulking. Genes in bold have $p \text{ adj} < 0.1$.