Supplemental Table II: Cilia-related Gene Expression in Airway Epithelium[1]

Gene symbol	Probe set ID	Gene title	P call (%)	Fold-change[2]	p value[3]	Reference
AKAP14	237281_at	A kinase (PRKA) anchor protein 14	100	-1.17	$1.18x10^{-01}$	[1]
AKAP14	237282_s_at	A kinase (PRKA) anchor protein 14	100	-1.43	$1.35x10^{-02}$	[1]
AKAP14	242698_at	A kinase (PRKA) anchor protein 14	57	-1.59	$4.75x10^{-02}$	[1]
ALMS1	214220_s_at	Alstrom syndrome 1	100	-1.18	$4.16x10^{-02}$	[2]
ALMS1	214221_at	Alstrom syndrome 1	100	-1.08	$7.02x10^{-01}$	[2]
ALMS1	214707__at	Alstrom syndrome 1	100	1.15	$4.86x10^{-01}$	[2]
BBS1	218471_s_at	Bardet-Biedl syndrome 1	100	-1.08	$5.42x10^{-01}$	[3,4]
BBS1	222643_s_at	Bardet-Biedl syndrome 1	75	-1.34	$6.62x10^{-02}$	[3,4]
BBS1	229142_s_at	CDNA FLJ52492 complete cds, highly similar to Bardet-Biedl syndrome 1 protein	4	-1.15	$6.60x10^{-01}$	[3,4]
BBS2	223227_at	Bardet-Biedl syndrome 2	100	-1.x10	$2.x10x10^{-01}$	[5]
BBS4	212744_at	Bardet-Biedl syndrome 4	100	-1.x10	$3.61x10^{-01}$	[6]
BBS4	212745_s_at	Bardet-Biedl syndrome 4	100	-1.19	$1.27x10^{-01}$	[6]
BBS5	230697_at	Bardet-Biedl syndrome 5	94	-1.36	$6.36x10^{-03}$	[7]
BBS7	219688_at	Bardet-Biedl syndrome 7	88	-1.03	$8.92x10^{-01}$	[8]
BBS7	235007_at	Bardet-Biedl syndrome 7	100	-1.15	$2.80x10^{-01}$	[8]
CDKN3	1555758_a_at	cyclin-dependent kinase inhibitor 3	6	1.14	$7.78x10^{-01}$	[9]
CDKN3	209714_s_at	cyclin-dependent kinase inhibitor 3	12	1.21	$4.86x10^{-01}$	[9]
CEP164	1558953_s_at	centrosomal protein 164kDa	100	-1.05	$7.48x10^{-01}$	[10]
CEP164	204250_s_at	centrosomal protein 164kDa	69	-1.14	$5.42x10^{-01}$	[10]
CEP164	204251_s_at	centrosomal protein 164kDa	100	1.00	$9.87x10^{-01}$	[10]
CETN2	209194_at	centrin, EF-hand protein, 2	100	-1.17	$8.74x10^{-02}$	[11]
CRB3	232609_at	crumbs homolog 3 (Drosophila)	75	-1.02	$9.29x10^{-01}$	[12]
CROCC	206274_s_at	ciliary rootlet coiled-coil, rootletin	100	-1.17	$4.86x10^{-01}$	[12]
CROCC	216419_at	ciliary rootlet coiled-coil, rootletin	12	-1.31	$3.65x10^{-01}$	[12]
DNAHx10	1562462_at	dynein, axonemal, heavy chain x10	92	-1.75	$2.40x10^{-03}$	[13]
DNAHx10	1565339_at	dynein, axonemal, heavy chain x10	65	-1.94	$3.24x10^{-03}$	[13]
DNAHx10	229738_at	dynein, axonemal, heavy chain x10	100	-1.26	$5.76x10^{-02}$	[13]
DNAH11	1553159_at	dynein, axonemal, heavy chain 11	98	-1.57	$9.03x10^{-04}$	[14]
DNAH11	1560416_at	dynein, axonemal, heavy chain 11	100	-1.04	$8.35x10^{-01}$	[14]

Supplemental Table II: Cilia-related Gene Expression in Airway Epithelium [1] (cont., page 2)

Gene symbol	Probe set ID	Gene title	P call (%)	Fold-change[2]	p value[3]	Reference
DNAH11	237804_at	Axonemal dynein heavy chain (DNAH11), partial	43	-1.00	$9.98x10^{-01}$	[14]
DNAH3	1560803_at	dynein, axonemal, heavy chain 3	96	-1.08	$6.34x10^{-01}$	[15]
DNAH3	1563290_at	dynein, axonemal, heavy chain 3	29	-1.43	$7.47x10^{-02}$	[15]
DNAH3	215266_at	dynein, axonemal, heavy chain 3	100	-1.36	$5.24x10^{-02}$	[15]
DNAH3	216415_at	dynein, axonemal, heavy chain 3 (DNAH3), mRNA	71	-1.00	$9.86x10^{-01}$	[15]
DNAH3	220725__at	dynein, axonemal, heavy chain 3 (DNAH3), mRNA	100	-1.06	$7.19x10^{-01}$	[15]
DNAH5	232381_s_at	dynein, axonemal, heavy chain 5	100	-1.68	$9.60x10^{-05}$	[16]
DNAH5	243938__at	dynein, axonemal, heavy chain 5	100	-1.83	$9.60x10^{-05}$	[16]
DNAH7	214222_at	dynein, axonemal, heavy chain 7	100	-1.31	$4.79x10^{-03}$	[17]
DNAH7	234476_at	dynein, axonemal, heavy chain 7	100	-1.23	$8.85x10^{-02}$	[17]
DNAH7	234727_at	dynein, axonemal, heavy chain 7	29	1.37	$2.x10x10^{-01}$	[17]
DNAH9	207959_s_at	dynein, axonemal, heavy chain 9	100	-1.19	$1.67x10^{-01}$	[18]
DNAH9	2x10345_s_at	dynein, axonemal, heavy chain 9	100	-1.56	$1.40x10^{-02}$	[18]
DNAH9	240857_at	dynein, axonemal, heavy chain 9	100	-1.39	$3.24x10^{-03}$	[18]
DNAI1	220125_at	dynein, axonemal, intermediate chain 1	100	-1.06	$7.48x10^{-01}$	[19]
DNAI1	233195_at	dynein, axonemal, intermediate chain 1	59	1.09	$4.88x10^{-01}$	[19]
DNAI2	220636_at	dynein, axonemal, intermediate chain 2	100	1.07	$5.42x10^{-01}$	[20]
DNAI2	221668_s_at	dynein, axonemal, intermediate chain 2	100	-1.07	$7.40x10^{-01}$	[20]
DNAL4	204008_at	dynein, axonemal, light chain 4	100	-1.06	$6.95x10^{-01}$	[21]
DNALI1	205186_at	dynein, axonemal, light intermediate chain 1	100	-1.46	$3.12x10^{-02}$	[22]
DNALI1	227081_at	dynein, axonemal, light intermediate chain 1	100	-1.21	$5.58x10^{-02}$	[22]
DYNC1H1	1556831_at	KIAA0325 protein	29	1.11	$7.65x10^{-01}$	[23]
DYNC1H1	211928_at	dynein, cytoplasmic 1, heavy chain 1	100	-1.05	$7.48x10^{-01}$	[23]
DYNC1H1	229042_s_at	dynein, cytoplasmic 1, heavy chain 1	2	1.47	$3.69x10^{-01}$	[23]
DYNC1H1	229115_at	dynein, cytoplasmic 1, heavy chain 1	20	-1.03	$9.20x10^{-01}$	[23]
DYNC1H1	236183_at	dynein, cytoplasmic 1, heavy chain 1	0	1.16	$7.90x10^{-01}$	[23]
DYNC1H1	24x1084__at	dynein, cytoplasmic 1, heavy chain 1	0	-1.09	$8.85x10^{-01}$	[23]
DYNC1H1	244768_at	dynein, cytoplasmic 1, heavy chain 1	0	-1.05	$9.20x10^{-01}$	[23]
DYNC1I2	211684_s_at	dynein, cytoplasmic 1, intermediate chain 2	100	-1.08	$4.51x10^{-01}$	[24]
DYNC2H1	1561939_at	dynein, cytoplasmic 2, heavy chain 1	100	-1.90	$9.03x10^{-04}$	[25]
DYNC2H1	1565149_at	dynein, cytoplasmic 2, heavy chain 1	100	1.05	$7.89x10^{-01}$	[25]
DYNC2H1	219469_at	dynein, cytoplasmic 2, heavy chain 1	100	-1.26	$6.63x10^{-02}$	[25]

Supplemental Table II: Cilia-related Gene Expression in Airway Epithelium [1] (cont., page 3)

Gene symbol	Probe set ID	Gene title	P call (%)	Fold-change[2]	p value[3]	Reference
DYNC2LI1	1554324_s_at	dynein, cytoplasmic 2, light intermediate chain 1	82	-1.03	$8.92x10^{-01}$	[26]
DYNC2LI1	203762_s_at	dynein, cytoplasmic 2, light intermediate chain 1	100	-1.04	$7.97x10^{-01}$	[26]
DYNC2LI1	203763_at	dynein, cytoplasmic 2, light intermediate chain 1	100	-1.14	$2.85x10^{-01}$	[26]
DYNC2LI1	241757__at	dynein, cytoplasmic 2, light intermediate chain 1	100	-1.01	$9.58x10^{-01}$	[26]
DYNLT1	201999_s_at	dynein, light chain, Tcte-type 1	100	-1.08	$3.42x10^{-01}$	[27]
EFHC1	219833_s_at	EF-hand domain (C-terminal) containing 1	100	-1.18	$5.87x10^{-02}$	[28]
EFHC1	225656_at	EF-hand domain (C-terminal) containing 1	100	-1.05	$7.19x10^{-01}$	[28]
EFHC1	23x1026_at	EF-hand domain (C-terminal) containing 1	100	-1.24	$3.12x10^{-02}$	[28]
EML1	204796_at	echinoderm microtubule associated protein like 1	100	-1.41	$2.48x10^{-02}$	[29]
EML1	204797_s_at	echinoderm microtubule associated protein like 1	100	-1.36	$3.24x10^{-03}$	[29]
EZR	208621_s_at	ezrin	100	-1.67	$5.64x10^{-03}$	[30]
EZR	208622_s_at	ezrin	100	-1.41	$1.02x10^{-02}$	[30]
EZR	208623_s_at	ezrin	100	-1.15	$3.88x10^{-02}$	[30]
EZR	217230_at	ezrin	49	1.09	$8.52x10^{-01}$	[30]
EZR	217234_s_at	ezrin	100	-1.69	$4.47x10^{-03}$	[30]
F2RL2	206795_at	coagulation factor II (thrombin) receptor-like 2	0	1.24	$6.79x10^{-01}$	[31]
F2RL2	230147_at	coagulation factor II (thrombin) receptor-like 2	6	2.60	$1.38x10^{-02}$	[31]
FOXJ1	205906_at	forkhead box J1	100	-1.38	$1.53x10^{-01}$	[30]
GAS8	204921_at	growth arrest-specific 8	100	1.01	$9.33x10^{-01}$	[32]
GSK3B	209945_s_at	glycogen synthase kinase 3 beta	100	-1.13	$2.78x10^{-01}$	[33]
IFT122	1563794_s_at	intraflagellar transport 122 homolog (Chlamydomonas)	51	-1.01	$9.75x10^{-01}$	[34]
IFT122	216678_at	intraflagellar transport 122 homolog (Chlamydomonas)	73	-1.06	$7.89x10^{-01}$	[34]
IFT122	220744_s_at	intraflagellar transport 122 homolog (Chlamydomonas)	100	-1.02	$8.77x10^{-01}$	[34]
IFT140	204792_s_at	intraflagellar transport 140 homolog (Chlamydomonas)	100	1.06	$7.89x10^{-01}$	[35]
IFT140	232844_at	KIAA0590 protein	43	1.03	$8.52x10^{-01}$	[35]
IFT140	244585_at	intraflagellar transport 140 homolog (Chlamydomonas)	0	1.42	$3.65x10^{-01}$	[35]
IFT172	226324_s_at	intraflagellar transport 172 homolog (Chlamydomonas)	100	-1.24	$1.50x10^{-02}$	[36]
IFT20	2x10312_s_at	intraflagellar transport 20 homolog (Chlamydomonas)	100	-1.22	$4.75x10^{-02}$	[37]
IFT52	218709_s_at	intraflagellar transport 52 homolog (Chlamydomonas)	100	-1.17	$6.24x10^{-02}$	[38]
IFT52	233532__at	intraflagellar transport 52 homolog (Chlamydomonas)	92	-1.19	$1.82x10^{-01}$	[38]
IFT57	218x100_s_at	intraflagellar transport 57 homolog (Chlamydomonas)	100	-1.17	$1.41x10^{-01}$	[37]
IFT57	222519_s_at	intraflagellar transport 57 homolog (Chlamydomonas)	100	-1.46	$3.12x10^{-02}$	[37]

Supplemental Table II: Cilia-related Gene Expression in Airway Epithelium [1] (cont., page 4)

Gene symbol	Probe set ID	Gene title	P call (%)	Fold-change[2]	p value[3]	Reference
IFT57	222520_s_at	intraflagellar transport 57 homolog (Chlamydomonas)	100	-1.46	$2.40x10^{-03}$	[37]
IFT80	1558956_s_at	intraflagellar transport 80 homolog (Chlamydomonas)	100	-1.03	$8.92x10^{-01}$	[39]
IFT80	1564231_at	intraflagellar transport 80 homolog (Chlamydomonas)	96	1.37	$5.94x10^{-03}$	[39]
IFT80	226098_at	intraflagellar transport 80 homolog (Chlamydomonas)	100	-1.08	$5.42x10^{-01}$	[39]
IFT80	240251_at	CDNA FLJ52072 complete cds, highly similar to Intraflagellar transport 80 homolog	22	1.67	$9.61x10^{-02}$	[39]
IFT88	204703_at	intraflagellar transport 88 homolog (Chlamydomonas)	100	-1.18	$5.82x10^{-02}$	[13]
KIF17	222144_at	kinesin family member 17	18	-1.07	$8.77x10^{-01}$	[40]
KIF3A	213623_at	kinesin family member 3A	100	-1.28	$5.55x10^{-02}$	[23]
KIF3A	228680_at	kinesin family member 3A	100	-1.26	$8.52x10^{-03}$	[23]
KIF3B	203943_at	kinesin family member 3B	100	-1.01	$9.29x10^{-01}$	[23]
KIF3B	225205_at	kinesin family member 3B	100	-1.05	$6.25x10^{-01}$	[23]
KIFAP3	203333_at	kinesin-associated protein 3	100	-1.24	$5.24x10^{-02}$	[41]
KLC1	212877_at	kinesin light chain 1	98	1.06	$6.34x10^{-01}$	[42]
KLC1	212878_s_at	kinesin light chain 1	100	-1.23	$6.29x10^{-02}$	[42]
KLC1	213656_s_at	kinesin light chain 1	100	-1.03	$7.89x10^{-01}$	[42]
KLC1	232036_at	KLC1 mRNA for kinesin light chain 1, complete cds, clone: FLJ08153AAAN	2	1.14	$8.09x10^{-01}$	[42]
KLC2	218906__at	kinesin light chain 2	65	-1.04	$8.52x10^{-01}$	[42]
LBR	201795_at	lamin B receptor	100	-1.x10	$5.57x10^{-01}$	[43]
MKS1	1555820_a_at	Meckel syndrome, type 1	100	-1.20	$1.41x10^{-01}$	[44]
MKS1	218630_at	Meckel syndrome, type 1	100	-1.08	$5.42x10^{-01}$	[44]
MLF1	204783_at	myeloid leukemia factor 1	100	-1.38	$4.05x10^{-03}$	[32]
MLF1	204784_s_at	myeloid leukemia factor 1	100	-1.41	$3.24x10^{-03}$	[32]
MYCBP	203359_s_at	c-myc binding protein	100	1.03	$8.52x10^{-01}$	[32]
MYCBP	203360_s_at	c-myc binding protein	100	-1.14	$2.60x10^{-01}$	[32]
MYCBP	203361_s_at	c-myc binding protein	100	-1.01	$9.54x10^{-01}$	[32]
NASP	201969_at	nuclear autoantigenic sperm protein (histone-binding)	94	-1.09	$4.65x10^{-01}$	[45]
NASP	201970_s_at	nuclear autoantigenic sperm protein (histone-binding)	100	-1.29	$4.75x10^{-02}$	[45]
NASP	237276_at	Nuclear autoantigenic sperm protein (histone-binding) (NASP), transcript variant 1, mRNA	0	1.03	$9.38x10^{-01}$	[45]
NASP	242918_at	Nuclear autoantigenic sperm protein (histone-binding)	65	-1.23	$4.29x10^{-01}$	[45]

Supplemental Table II: Cilia-related Gene Expression in Airway Epithelium [1] (cont., page 5)

Gene symbol	Probe set ID	Gene title	P call (%)	Fold-change[2]	p value[3]	Reference
		(NASP), transcript variant 1, mRNA				
NME5	206197_at	non-metastatic cells 5, protein epressed in (nucleoside-diphosphate kinase)	100	-1.14	$2.x10x10^{-01}$	[32]
NME7	219553_at	non-metastatic cells 7, protein epressed in (nucleoside-diphosphate kinase)	100	-1.x10	$4.77x10^{-01}$	[32]
NME7	227556_at	non-metastatic cells 7, protein epressed in (nucleoside-diphosphate kinase)	100	1.x10	$7.04x10^{-01}$	[32]
NPHP1	206285_at	nephronophthisis 1 (juvenile)	78	1.11	$5.42x10^{-01}$	[46]
NPHP1	238843_at	nephronophthisis 1 (juvenile)	100	1.02	$9.20x10^{-01}$	[46]
NPHP1	238844_s_at	nephronophthisis 1 (juvenile)	100	-1.41	$4.79x10^{-03}$	[46]
NPHP3	1553389_at	nephronophthisis 3 (adolescent)	16	1.11	$7.97x10^{-01}$	[46]
NPHP4	213471_at	nephronophthisis 4	98	1.09	$5.57x10^{-01}$	[46]
NPHP4	216344_at	nephronophthisis 4	6	1.05	$9.20x10^{-01}$	[46]
ODF2	2x10415_s_at	outer dense fiber of sperm tails 2	82	-1.63	$4.05x10^{-03}$	[47]
ODF2	225617_at	outer dense fiber of sperm tails 2	94	-1.17	$5.37x10^{-01}$	[47]
OFD1	203569_s_at	oral-facial-digital syndrome 1	100	-1.18	$3.57x10^{-02}$	[48]
OFD1	241751_at	oral-facial-digital syndrome 1	100	1.x10	$7.18x10^{-01}$	[48]
PACRG	214204_at	PARK2 co-regulated	100	-1.17	$2.31x10^{-01}$	[49]
PACRG	215472_at	PARK2 co-regulated	100	-1.25	$7.47x10^{-02}$	[49]
PCM1	202174_s_at	pericentriolar material 1	100	-1.28	$4.79x10^{-03}$	[50]
PCM1	209996__at	pericentriolar material 1	14	-1.53	$5.76x10^{-02}$	[50]
PCM1	209997__at	pericentriolar material 1	100	-1.20	$1.24x10^{-01}$	[50]
PCM1	214118__at	pericentriolar material 1	100	-1.08	$6.14x10^{-01}$	[50]
PCM1	214937__at	pericentriolar material 1	100	-1.09	$2.21x10^{-01}$	[50]
PCM1	228905_at	pericentriolar material 1	100	-1.37	$9.44x10^{-03}$	[50]
PCNT	203660_s_at	pericentrin	98	-1.05	$7.09x10^{-01}$	[51]
PCNT	233387_s_at	pericentrin	14	-1.71	$5.34x10^{-02}$	[51]
PKD1	202327_s_at	polycystic kidney disease 1 (autosomal dominant)	25	-1.17	$5.68x10^{-01}$	[52]
PKD1	202328_s_at	polycystic kidney disease 1 (autosomal dominant)	98	-1.01	$9.58x10^{-01}$	[52]
PKD1	216949_s_at	polycystic kidney disease 1 (autosomal dominant)	25	-1.47	$3.71x10^{-01}$	[52]
PKD2	203688_at	polycystic kidney disease 2 (autosomal dominant)	100	-1.31	$3.71x10^{-02}$	[52]
PLEKHA8	1560556_a_at	Homo sapiens, clone IMAGE:4617677, mRNA	20	1.22	$7.19x10^{-01}$	[53]

Supplemental Table II: Cilia-related Gene Expression in Airway Epithelium [1] (cont., page 6)

Gene symbol	Probe set ID	Gene title	P call (%)	Fold-change[2]	p value[3]	Reference
PLEKHA8	224208_at	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8	2	1.27	$6.85x10^{-01}$	[53]
PLEKHA8	231543_at	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8	0	-1.x10	$8.52x10^{-01}$	[53]
PLEKHA8	232212_at	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8	2	-1.08	$8.62x10^{-01}$	[53]
PLEKHA8	233943__at	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8	47	1.34	$2.86x10^{-01}$	[53]
PLEKHA8 /// PLEKHA9	220157__at	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8 /// pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 9	61	1.08	$6.64x10^{-01}$	[53]
PPP2R1A	200695_at	protein phosphatase 2 (formerly 2A), regulatory subunit A, alpha isoform	96	-1.20	$2.78x10^{-01}$	[32]
RABL4	205037_at	RAB, member of RAS oncogene family-like 4	100	-1.21	$1.27x10^{-01}$	[32]
RABL4	213784_at	RAB, member of RAS oncogene family-like 4	100	-1.04	$8.32x10^{-01}$	[32]
RABL4	243812_at	RAB, member of RAS oncogene family-like 4	0	-1.03	$9.45x10^{-01}$	[32]
RPGRIP1L	213959_s_at	RPGRIP1-like	100	-1.02	$9.20x10^{-01}$	[54]
RPGRIP1L	24x1028_at	RPGRIP1-like, mRNA (cDNA clone IMAGE:4697938)	6	-1.04	$9.29x10^{-01}$	[54]
RSPH3	1568613_at	radial spoke 3 homolog (Chlamydomonas)	0	-1.03	$8.86x10^{-01}$	[55]
RSPH3	223713_at	radial spoke 3 homolog (Chlamydomonas)	98	-1.16	$3.90x10^{-01}$	[55]
SLC9A3R1	201349_at	solute carrier family 9 (sodium/hydrogen echanger), member 3 regulator 1	100	-1.08	$5.18x10^{-01}$	[56]
SPAG6	2x10032_s_at	sperm associated antigen 6	100	-1.45	$2.48x10^{-03}$	[57]
SPAG6	2x10033_s_at	sperm associated antigen 6	100	-1.34	$3.31x10^{-03}$	[57]
TCTE3	1554400_at	t-comple-associated-testis-epressed 3	8	1.63	$2.x10x10^{-01}$	[58]
TCTE3	1554401_a_at	t-comple-associated-testis-epressed 3	2	-1.40	$4.74x10^{-01}$	[58]
TCTE3	1557945_at	dynein light chain 2 TCTE2	98	1.05	$7.48x10^{-01}$	[58]
TCTE3	232258_at	dynein light chain 2 TCTE2	4	-1.01	$9.77x10^{-01}$	[58]
TEKT1	239216_at	tektin 1	100	-1.45	$2.40x10^{-03}$	[59]
TEKT2	2x10323_at	tektin 2 (testicular)	100	-1.37	$3.77x10^{-02}$	[60]
TEKT3	223867_at	tektin 3	100	-1.27	$5.87x10^{-02}$	[61]

Supplemental Table II: Cilia-related Gene Expression in Airway Epithelium [1] (cont., page 7)

Gene symbol	Probe set ID	Gene title	P call (%)	Fold-change[2]	p value[3]	Reference
TMEM67	1552763_at	transmembrane protein 67	14	-1.58	$3.42x10^{-01}$	[44]
TMEM67	1552765__at	transmembrane protein 67	37	-1.05	$9.29x10^{-01}$	[44]
TMEM67	1563646_a_at	transmembrane protein 67	100	-1.44	$3.12x10^{-02}$	[44]
TMEM67	1569377_at	transmembrane protein 67	100	-1.02	$9.29x10^{-01}$	[44]
TMEM67	1569411_at	transmembrane protein 67	22	-1.57	$2.28x10^{-01}$	[44]
TMEM67	232023_at	transmembrane protein 67	100	-1.38	$4.75x10^{-02}$	[44]
TMEM67	238229_at	transmembrane protein 67	100	-1.17	$2.31x10^{-01}$	[44]
TUBA1C	209251__at	tubulin, alpha 1c	100	-1.01	$9.33x10^{-01}$	[62]
TUBA1C	211750__at	tubulin, alpha 1c	100	-1.03	$7.90x10^{-01}$	[62]
TUBB2A	204141_at	tubulin, beta 2A	100	-1.08	$6.44x10^{-01}$	[62]
TUBB2A /// TUBB2B	209372__at	tubulin, beta 2A /// tubulin, beta 2B	90	-1.64	$4.75x10^{-02}$	[62]
VCP	208648_at	valosin-containing protein	100	-1.01	$9.29x10^{-01}$	[32,63]
VCP	208649_s_at	valosin-containing protein	100	-1.04	$7.60x10^{-01}$	[32,63]
VHL	1559227_s_at	von Hippel-Lindau tumor suppressor	59	1.05	$7.90x10^{-01}$	[64]
VHL	203844_at	von Hippel-Lindau tumor suppressor	29	1.53	$4.19x10^{-01}$	[64]

[1] The HG-U133 Plus 2.0 microarray platform was used to detect differences in the expression of these genes in healthy smokers compared to healthy nonsmokers.

[2] Magnitude of fold change in average expression values for healthy smokers *vs* healthy nonsmokers.

[3] A Benjamini-Hochberg correction was used to limit the false positive rate.

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