Supplementary Table 5: Regions of recurrent amplification in 56 triple negative breast cancers.

Chromosome	Cytobands	Start (Mb)	End (Mb)	length MB	Frequency	Cases	Genes	mirnas	Copy number variations (CGH)
1	p32.2-p31.3	58.26	61.78	3.519815	5%	TN.12523, TN.BC3044, TN.NT005	DAB1, OMA1, TACSTD2, MYSM1, JUN, FGGY, HOOK1, CYP2J2, C1orf87, NFIA		V_4226_LC0413_Wong et al. (2007), V_4227_LC0416_Wong et al. (2007)
1	p31.3	64	64.86	0.856887	4%	TN.BC3044, TN.NT005	EFCAB7, DLEU2L, PGM1, ROR1, UBE2U		
1	p31.3-p31.2	66.64	69.36	2.720669	5%	TN.B6, TN.BC3044, TN.NT005	PDE4B, SGIP1, TCTEX1D1, INSL5, WDR78, MIER1, SLC35D1, C1orf141, IL23R,	hsa-mir-1262	V_0009_LC0463_Iafrate et al. (2004)
1	p22.3	87.52	87.98	0.459519	4%	TN.12523, TN.BC2697	HS2ST1, LMO4		
1	p12	119.26	119.88	0.614307	4%	TN.1878, TN.BC3045	TBX15, WARS2		
1	p12-p11.2	120.45	121.35	0.89588	13%	TN.B18, TN.B41, TN.B6, TN.B81, TN.BC3045, TN.NT004,	NOTCH2, FAM72B, FCGR1B		V_4246_LC0743_Wong et al. (2007)
1	q21.1-q31.1	143.51	189.98	46.474323	14%	TN.10030, TN.11093, TN.3319, TN.4274, TN.7880,	PPIAL4G, FAM72D, SRGAP2P2, PPIAL4B, NBPF9, PDE4DIP, SEC22B, NOTCH2NL,	hsa-mir-554, hsa-mir-190b, hsa-	V_0014_LC0941_Iafrate et al. (2004), V_0015_LC0954_Iafrate et al. (2004), V_0685_LC0752_Sharp et al. (2005),
1	q31.2-q31.3	191.26	197.59	6.326642	5%	TN.10030, TN.B6, TN.NT0015	RGS18, RGS1, RGS13, RGS2, UCHL5, TROVE2, GLRX2, CDC73, B3GALT2, KCNT2,	hsa-mir-1278	V_0686_LC1055_Sharp et al. (2005), V_0687_LC1055_Sharp et al. (2005), V_2053_LC1052_Locke et al. (2006),
1	q31.3-q32.1	197.93	200.85	2.915265	5%	TN.10030, TN.B6, TN.NT0015	NEK7, ATP6V1G3, PTPRC, NR5A2, C1orf98, ZNF281, KIF14, DDX59, CAMSAP1L1,	hsa-mir-181b-1, hsa-mir-181a-1	V_4267_LC1097_Wong et al. (2007)
1	q32.1	203.64	204.03	0.393461	4%	TN.10030, TN.NT0015	ATP2B4, LAX1, ZC3H11A, SNRPE, C1orf157		
1	q32.1-q32.2	206.61	211.1	4.489412	7%	TN.10030, TN.7880, TN.B6, TN.BC3045, TN.NT0015	SRGAP2, IKBKE, C1orf147, RASSF5, LGTN, DYRK3, MAPKAPK2, IL10, IL19, IL20, IL24,	hsa-mir-29c, hsa-mir-29b-2, hsa-	
1	q32.3-q41	213.38	223.71	10.327949	11%	TN.10030, TN.B41, TN.B6, TN.BC3045, TN.NT0015,	RPS6KC1, PROX1, SMYD2, PTPN14, CENPF, KCNK2, KCTD3, USH2A, ESRRG,	hsa-mir-215, hsa-mir-194-1	V_4270_LC1176_Wong et al. (2007)
1	q42.12-q42.13	226.39	227.63	1.233736	4%	TN.7880, TN.NT0015	MIXL1, LIN9, PARP1, C1orf95, ITPKB, PSEN2, CABC1, CDC42BPA		
1	q42.13-q44	228.79	248.88	20.086973	9%	TN.10030, TN.10137, TN.11155, TN.12303, TN.1878,	RHOU, TMEM78, RAB4A, SPHAR, C1orf96, ACTA1, NUP133, ABCB10, TAF5L, URB2,	hsa-mir-1182, hsa-mir-1537	V_0016_LC1285_Iafrate et al. (2004), V_0017_LC1401_Iafrate et al. (2004), V_2056_LC1401_Locke et al. (2006),
3	p26.1	4.38	4.86	0.47423	4%	TN.10662, TN.NT0024	SUMF1, ITPR1		
3	q23	139.29	140.59	1.294612	4%	TN.B6, TN.B78	NMNAT3, CLSTN2, TRIM42		
3	q24-q25.1	148.9	149.31	0.41387	5%	TN.12303, TN.1267, TN.B78	CP, TM4SF18, TM4SF1, TM4SF4, WWTR1		
3	q25.1-q25.2	149.65	153.02	3.365157	7%	TN.B17, TN.B6, TN.B64, TN.B78	RNF13, PFN2, TSC22D2, SERP1, EIF2A, C3orf44, SIAH2, C3orf76, CLRN1, MED12L,		V_4355_LC3723_Wong et al. (2007), V_4356_LC3737_Wong et al. (2007)
3	q25.31-q25.32	155.87	157.66	1.790988	4%	TN.1878, TN.B64	KCNAB1, SSR3, TIPARP, METT5D2, LEKR1, CCNL1, VEPH1, PTX3, C3orf55		V_4358_LC3775_Wong et al. (2007)
3	q25.32-q25.33	158.38	160.26	1.882166	4%	TN.1878, TN.B64	GFM1, LXN, RARRES1, MFSD1, IQCJ, SCHIP1, IL12A, IFT80, SMC4, TRIM59, KPNA4	hsa-mir-15b, hsa-mir-16-2	
3	q26.1-q26.32	166.13	176.8	10.669782	5%	TN.1878, TN.B2, TN.B6, TN.B65	ZBBX, SERPINI2, WDR49, PDCD10, SERPINI1, GOLIM4, EVI1, MYNN, LRRC34,	hsa-mir-551b, hsa-mir-569	V_0039_LC3834_Iafrate et al. (2004), V_4362_LC3840_Wong et al. (2007), V_4363_LC3872_Wong et al. (2007)
4	p16.3	3.39	3.66	0.268738	4%	TN.B64, TN.B86	RGS12, HGFAC, DOK7, LRPAP1		V_4371_LC4159_Wong et al. (2007), V_4372_LC4159_Wong et al. (2007)
5	p15.33-p15.1	1.56	16.72	15.160152	9%	TN.1878, TN.3701, TN.B65, TN.B78, TN.BC2968,	MRPL36, NDUFS6, IRX4, IRX2, C5orf38, IRX1, ADAMTS16, MED10, NSUN2, SRD5A1,	hsa-mir-887	V_2082_LC6572_Locke et al. (2006), V_4430_LC6506_Wong et al. (2007)
5	p14.3	20.56	21.2	0.639397	4%	TN.3701, TN.NT004			V_0708_LC6659_Sharp et al. (2005), V_4434_LC6659_Wong et al. (2007)
5	p14.3-p14.1	22.32	26.56	4.237383	5%	TN.10646, TN.B65, TN.NT004	CDH12, PRDM9, CDH10		
5	p12	44.25	44.49	0.244421	4%	TN.10646, TN.NT004	FGF10		
6	q22.31-q22.33	124.14	129.07	4.928908	5%	TN.12303, TN.3319, TN.BC2968, TN.NT0024	NKAIN2, RNF217, TPD52L1, HDDC2, HEY2, NCOA7, HINT3, TRMT11, C6orf173,	hsa-mir-588	V_0090_LC8741_Iafrate et al. (2004), V_0091_LC8757_Iafrate et al. (2004)
6	q23.3	135.24	137.36	2.121275	4%	TN.12303, TN.BC2968	ALDH8A1, HBS1L, MYB, AHI1, PDE7B, FAM54A, BCLAF1, MAP7, MAP3K5, PEX7,	hsa-mir-548a-2	
7	p11.2	54.99	55.7	0.713173	4%	TN.10665, TN.BC184	EGFR, LANCL2		
7	q11.1	61.06	61.27	0.207558	4%	TN.4643, TN.B2			V_4538_LC9545_Wong et al. (2007), V_4539_LC9545_Wong et al. (2007)
7	q11.21	64.42	65.19	0.769661	4%	TN.4643, TN.7880	ZNF117, ERV3, ZNF92		V_0722_LC9554_Sharp et al. (2005), V_2101_LC9554_Locke et al. (2006)
7	q21.13-q21.2	90.86	91.46	0.604123	4%	TN.B45, TN.BC2968	FZD1, MTERF		
7	q21.2-q21.3	91.6	93.68	2.082652	4%	TN.4643, TN.B45	AKAP9, CYP51A1, KRIT1, ANKIB1, GATAD1, PEX1, C7orf64, FAM133B, CDK6, SAMD9,	hsa-mir-1285-1, hsa-mir-653, hsa-	
7	q33	136.97	137.46	0.490678	4%	TN.7880, TN.NT005	PTN, DGKI, RPL41		
7	q34	138.98	139.72	0.738026	4%	TN.7880, TN.B18	UBN2, LUC7L2, C7orf55, KLRG2, CLEC2L, HIPK2, TBXAS1		
7	q36.1	150.84	151.65	0.813473	4%	TN.B18, TN.NT0001	AGAP3, GBX1, ASB10, ABCF2, SMARCD3, NUB1, WDR86, CRYGN, RHEB, PRKAG2,	hsa-mir-671	
8	p12-q24.3	34.93	146.24	111.305555	32%	TN.10646, TN.10662, TN.10665, TN.11093, TN.12303,	UNC5D, KCNU1, ZNF703, ERLIN2, PROSC, GPR124, BRF2, RAB11FIP1, GOT1L1,	hsa-mir-486, hsa-mir-124-2, hsa-	V_0119_LC10711_Iafrate et al. (2004), V_0121_LC10727_Iafrate et al. (2004), V_0123_LC10906_Iafrate et al. (2004),
9	p24.3-p22.3	1.98	15.14	13.161039	9%	TN.7880, TN.B6, TN.B65, TN.BC2968, TN.BC3044,	SMARCA2, VLDLR, KCNV2, KIAA0020, CARM1L, RFX3, GLIS3, SLC1A1, C9orf68,	hsa-mir-101-2	V_4615_LC11143_Wong et al. (2007)
9	p22.3-p22.2	15.98	16.87	0.890872	4%	TN.BC2968, TN.NT005	C9orf93, C9orf92, BNC2		
9	p22.2-p21.3	17.65	20.48	2.82604	4%	TN.BC2968, TN.NT005	SH3GL2, ADAMTSL1, FAM154A, RRAGA, HAUS6, ADFP, DENND4C, RPS6, ACER2,		
10	p15.3-p12.1	0.06	26.98	26.92066	16%	TN.10137, TN.10752, TN.12523, TN.7880, TN.B17,	ZMYND11, DIP2C, C10orf108, LARP5, GTPBP4, IDI2, IDI1, WDR37, ADARB2, PFKP,	hsa-mir-1265, hsa-mir-511-1, hsa-	V_4664_LC11931_Wong et al. (2007), V_4665_LC12095_Wong et al. (2007), V_4666_LC12101_Wong et al. (2007),
10	p11.21	36.01	36.33	0.319526	4%	TN.NT005, TN.NT0067			
10	q11.22	46.69	48.28	1.589695	4%	TN.12303, TN.NT0067	FAM35B, SYT15, GPRIN2, ANXA8L1, PPYR1, FAM25B, BMS1P2, CTSLL7, FAM25HP,		V_0764_LC12274_Sharp et al. (2005), V_2153_LC12274_Locke et al. (2006), V_4685_LC12274_Wong et al. (2007),
10	q11.23	51.77	52.58	0.804264	4%	TN.B78, TN.NT0067	FAM21A, ASAH2, SGMS1, ASAH2B, A1CF		V_2156_LC12305_Locke et al. (2006)
10	q21.1-q21.3	60.43	66.41	5.988516	5%	TN.B17, TN.B41, TN.BC3044, TN.NT0067	BICC1, PHYHIPL, FAM13C, SLC16A9, CCDC6, C10orf40, ANK3, ARL4P, CDC2,	hsa-mir-1296	V_4697_LC12385_Wong et al. (2007), V_4698_LC12387_Wong et al. (2007), V_4699_LC12389_Wong et al. (2007)
10	q22.1	71.84	73.6	1.766933	5%	TN.10752, TN.11093, TN.NT0067	H2AFY2, AIFM2, TYSND1, SAR1A, PPA1, NPFFR1, LRRC20, EIF4EBP2, NODAL,		
10	q22.3	78.65	81.08	2.42975	4%	TN.B69, TN.NT0067	KCNMA1, DLG5, POLR3A, RPS24, ZMIZ1		V_0138_LC12468_Iafrate et al. (2004), V_2157_LC12477_Locke et al. (2006), V_2158_LC12478_Locke et al. (2006),
10	q26.12-q26.13	123.05	125.68	2.630053	4%	TN.10752, TN.9776	FGFR2, ATE1, NSMCE4A, TACC2, BTBD16, PLEKHA1, HTRA1, DMBT1, C10orf120,		V_4714_LC12735_Wong et al. (2007)
11	p13-p12	34.17	39.22	5.055173	5%	TN.B65, TN.B73, TN.B79	NAT10, ABTB2, CAT, ELF5, EHF, APIP, PDHX, CD44, SLC1A2, PAMR1, FJX1, TRIM44,		V_4736_LC13107_Wong et al. (2007), V_4737_LC13107_Wong et al. (2007)
11	q13.3-q13.4	69.6	70.53	0.925982	5%	TN.10499, TN.4643, TN.BC3047	FGF3, ANO1, FADD, PPFIA1, CTTN, SHANK2	hsa-mir-548k	
11	q23.3	119.48	119.96	0.480124	4%	TN.3701, TN.9961	PVRL1		
12	p13.33-p13.31	0.15	6.14	5.995011	13%	TN.12303, TN.12523, TN.4643, TN.9961, TN.B69,	FAM138D, IQSEC3, SLC6A12, SLC6A13, KDM5A, CCDC77, B4GALNT3, NINJ2, WNK1,		V_0161_LC13778_Iafrate et al. (2004), V_4765_LC13757_Wong et al. (2007), V_4766_LC13773_Wong et al. (2007)
12	p13.31-p13.2	9.25	12.33	3.087283	5%	TN.9961, TN.B6, TN.BC3044, TN.BC3047, TN.NT004	A2M, PZP, PTMAP4, DDX12, KLRB1, CLEC2D, CLEC1, CD69, KLRF1, CLEC2B,		V_4769_LC13838_Wong et al. (2007)
12	p13.2-p12.3	12.47	19	6.529615	4%	TN.BC3047, TN.NT004	MANSC1, LOH12CR2, LOH12CR1, DUSP16, CREBL2, GPR19, CDKN1B, APOLD1,	hsa-mir-613, hsa-mir-614	V_4770_LC13869_Wong et al. (2007)
12	p12.3-p11.1	19.74	33.44	13.704219	7%	TN.B78, TN.BC3047, TN.NT0001, TN.NT004, TN.NT005	PDE3A, SLCO1C1, SLCO1B3, SLCO1B1, SLCO1A2, IAPP, PYROXD1, RECQL,	hsa-mir-920	V_0162_LC13887_Iafrate et al. (2004), V_0163_LC13898_Iafrate et al. (2004), V_4772_LC13891_Wong et al. (2007),
13	q31.3	93.64	94.49	0.850316	4%	TN.7880, TN.BC2697	GPC6		
13	q32.3	101.23	101.68	0.448493	4%	TN.7880, TN.BC2697	A2LD1, TMTC4		V_4816_LC15155_Wong et al. (2007)
13	q33.2-q33.3	106.37	109.06	2.686443	4%	TN.7880, TN.B65	EFNB2, ARGLU1, FAM155A, LIG4, ABHD13, TNFSF13B	hsa-mir-1267	V_4817_LC15210_Wong et al. (2007), V_4818_LC15212_Wong et al. (2007), V_0361_LC15215_Shaw-Smith et al. (2004)
13	q33.3-q34	109.83	115.07	5.241634	5%	TN.4274, TN.7880, TN.BC2697	MYO16, IRS2, COL4A1, COL4A2, RAB20, CARKD, CARS2, ING1, C13orf29, ANKRD10,		V_4819_LC15248_Wong et al. (2007), V_4820_LC15248_Wong et al. (2007)
14	q11.2	19.97	20.41	0.446278	4%	TN.B2, TN.B45	OR11H2, OR4Q3, OR4H12P, OR4M1, OR4N2, OR4K2, OR4K5, OR4K1		
15	q11.2	20.83	21.11	0.284766	4%	TN.B2, TN.B78	BCL8		V_4879_LC16090_Wong et al. (2007)
15	q26.2	95.9	97.64	1.741426	4%	TN.12523, TN.B18	NR2F2, FAM149B2, SPATA8	hsa-mir-1469	V_0192_LC16541_Iafrate et al. (2004)
15	q26.2-q26.3	98.31	99.78	1.476949	4%	TN.B18, TN.B64	ARRDC4, FAM169B, IGF1R, SYNM, TTC23		V_2201_LC16579_Locke et al. (2006)
15	q26.3	101.37	102.27	0.905945	4%	TN.B18, TN.B64	ALDH1A3, LRRK1, CHSY1, SNRPA1, PCSK6, TM2D3, TARSL2		
19	p13.12	14.47	15.44	0.965629	4%	TN.11093, TN.7880	CD97, DDX39, PKN1, PTGER1, GIPC1, DNAJB1, GPSN2, NDUFB7, CLEC17A, EMR3,	hsa-mir-639	V_5090_LC18847_Wong et al. (2007)
19	q11-q13.11	27.95	32.91	4.954227	9%	TN.7880, TN.B41, TN.B6, TN.BC2968, TN.NT0015,	UQCRFS1, VSTM2B, POP4, PLEKHF1, C19orf12, CCNE1, C19orf2, ZNF536, TSHZ3,		V_5096_LC18924_Wong et al. (2007)
19	q13.11	33.76	34.52	0.763047	4%	TN.B41, TN.B81	CEBPG, PEPD, CHST8, KCTD15		
19	q13.12	37.85	38.21	0.362527	4%	TN.7880, TN.B41	ZNF527, ZNF569, ZNF570, ZNF793, ZNF540, ZNF571, ZFP30, ZNF781, ZNF607		
19	q13.2	40.43	41	0.567833	4%	TN.B41, TN.NT0015	FCGBP, PSMC4, ZNF546, ZNF780B, ZNF780A, MAP3K10, TTC9B, CNTD2, AKT2,	hsa-mir-641	
21	q11.2-q21.1	15.74	20.45	4.707721	5%	TN.B18, TN.B81, TN.NT0067	HSPA13, SAMSN1, NRIP1, USP25, C21orf34, C21orf37, CXADR, BTG3, C21orf91,	hsa-mir-99a, hsa-let-7c, hsa-mir-	V_5159_LC20079_Wong et al. (2007)
21	q21.2-q21.3	24.3	30.26	5.957184	4%	TN.3319, TN.NT0067	MRPL39, JAM2, ATP5J, GABPA, APP, CYYR1, ADAMTS1, ADAMTS5, N6AMT1	hsa-mir-155	
21	q21.3-q22.11	31.06	31.59	0.526838	4%	TN.3319, TN.NT0067	GRIK1, CLDN17, CLDN8		
21	q22.13-q22.3	39	47.37	8.371448	5%	TN.3319, TN.BC2697, TN.NT005, TN.NT0067	KCNJ6, DSCR4, DSCR8, KCNJ15, ERG, ETS2, C21orf104, PSMG1, BRWD1, C21orf87,		V_5162_LC20269_Wong et al. (2007), V_5163_LC20269_Wong et al. (2007)
22	q11.1	16.06	16.51	0.444161	4%	TN.B2, TN.B81	POTEH, OR11H1		
22	q11.21	18.72	18.96	0.239596	4%	TN.B53, TN.BC3044	DGCR6, PRODH		
23	q23	112.83	114.73	1.904359	4%	TN.B41, TN.B64	HTR2C, IL13RA2, LRCH2, LUZP4		
23	q26.3	134.8	134.98	0.177372	4%	TN.B41, TN.B64	CT45A3, CT45A5, SAGE1		V_0826_LC21120_Sharp et al. (2005), V_2281_LC21120_Locke et al. (2006)
1	p12-q21.3	120.16	152.13	31.96597	14%	TN.11093, TN.4274, TN.B18, TN.B41, TN.B6, TN.B81, TN.BC3045, TN.NT0015, TN.NT004, TN.NT005	NBPF7, ADAM30, NOTCH2, LOC440607, NBPF20, PDE4DIP, SEC22B, NOTCH2NL, HFE2, TXNIP, POLR3GL, ANKRD34, LIX1L, RBM8A, PEX11B, ITGA10, ANKRD35, PIAS3, NUDT17, POLR3C, ZNF364, CD160, PDZK1, FAM108A2, PRKAB2, FMO5, CHD1L, BCL9, ACP6, GJA5, GJA8, GPR89A, NBPF11, NBPF14, NBPF10, NBPF15, LOC388692, LOC440686, PPIAL4, FCGR1A, HIST2H2BF, H2A/R, HIST2H2AA, H3/o, HIST2H3C, H4/o, HIST2H4, HIST2H2BE, HIST2H2AC, HIST2H2AB, BOLA1, SV2A, SF3B4, MTMR11, OTUD7B, VPS45A, PLEKHO1, ANP32E, CA14, APH1A, C1orf54, C1orf51, MRPS21, PRPF3, KIAA0460, TARSL1, ECM1, ADAMTSL4, MCL1, ENSA, GOLPH3L, HORMAD1, CTSS, CTSK, ARNT, SETDB1, LASS2, ANXA9, FAM63A, PRUNE, BNIPL, C1orf56, CDC42SE1, MLLT11, MGC29891, SEMA6C, TNFAIP8L2, LYSMD1, SCNM1, TMOD4, VPS72, PIP5K1A, PSMD4, ZNF687, PIK4CB, RFX5, SELENBP1, PSMB4, POGZ, CGN, TUFT1, SNX27, TNRC4, MRPL9, OAZ3, TDRKH, LINGO4, RORC, THEM5, THEM4, S100A10, S100A11, TCHHL1, HRNR, FLG, FLG2, CRNN, LCE5A, C1orf42, LCE3E, LCE3D, LCE3C, LCE3B, LCE3A, LCE2D, LCE2C, LCE2B, LCE2A, LCE4A, C1orf45, LCE1F, LCE1E, LCE1D, LCE1C, LCE1B, LCE1A, SMCP, IVL, SPRR4, SPRR1A, SPRR3, SPR	hsa.mir.554	V_0683_LC0751_Sharp et al. (2005), V_0684_LC0752_Sharp et al. (2005), V_0685_LC0752_Sharp et al. (2005), V_2050_LC0752_Locke et al. (2006), V_2051_LC0752_Locke et al. (2006), V_2052_LC0752_Locke et al. (2006), V_4245_LC0743_Wong et al. (2007), V_4246_LC0743_Wong et al. (2007), V_4247_LC0752_Wong et al. (2007), V_4248_LC0752_Wong et al. (2007), V_4249_LC0752_Wong et al. (2007), V_4250_LC0752_Wong et al. (2007), V_4251_LC0752_Wong et al. (2007), V_4252_LC0752_Wong et al. (2007), V_4253_LC0752_Wong et al. (2007), V_4254_LC0752_Wong et al. (2007), V_4255_LC0752_Wong et al. (2007), V_4256_LC0752_Wong et al. (2007)
1	q21.3-q24.3	153.23	169.95	16.728232	16%	TN.11093, TN.3319, TN.4274, TN.7880, TN.B41, TN.B6, TN.BC3045, TN.NT0015, TN.NT004, TN.NT005	FLAD1, LENEP, ZBTB7B, DCST2, DCST1, ADAM15, EFNA4, EFNA3, EFNA1, RAG1AP1, DPM3, KRTCAP2, TRIM46, MUC1, THBS3, MTX1, GBA, C1orf2, SCAMP3, CLK2, HCN3, PKLR, FDPS, C1orf104, RUSC1, ASH1L, MSTO1, YY1AP1, DAP3, GON4L, SYT11, RIT1, KIAA0907, RXFP4, ARHGEF2, SSR2, UBQLN4, MAPBPIP, RAB25, LMNA, SEMA4A, KIAA0446, PMF1, BGLAP, PAQR6, SMG5, TMEM79, C1orf85, VHLL, CCT3, C1orf182, RHBG, C1orf61, MEF2D, IQGAP3, APOA1BP, GPATC4, HAPLN2, BCAN, NES, CRABP2, ISG20L2, C1orf66, MRPL24, HDGF, PRCC, SH2D2A, INSRR, NTRK1, PEAR1, C1orf92, ARHGEF11, FLJ16478, ETV3, FCRL5, FCRL4, FCRL3, FCRL2, FCRL1, CD5L, KIRREL, CD1D, CD1A, CD1C, CD1B, CD1E, OR10T2, OR10K2, OR10K1, OR10R2, OR6Y1, OR10X1, OR10Z1, SPTA1, OR6K2, OR6K3, OR6K6, OR6N1, OR6N2, MNDA, PYHIN1, IFI16, AIM2, IGSF4B, DARC, FCER1A, OR10J3, OR10D1, OR10J5, APCS, CRP, DUSP23, FCRL6, SLAMF8, VSIG8, CCDC19, TAGLN2, IGSF9, SLAMF9, PIGM, KCNJ10, KCNJ9, IGSF8, ATP1A2, ATP1A4, CASQ1, PEA15, WDR42A, PEX19, COPA, NCSTN, NHLH1, VANGL2, SLAMF6, CD84, SLAMF1, CD48, SLAMF7, LY9, CD244, ITLN1, ITLN2, F11R, USF1, ARHGAP30, PVRL4, KARCA1, PFDN2, NIT1, DEDD, UFC1, USP21, PPOX,	hsa.mir.92b, hsa.mir.555, hsa.mir.9.1, hsa.mir.765, hsa.mir.556, hsa.mir.921, hsa.mir.1255b.2, hsa.mir.557, hsa.mir.1295	V_4257_LC0831_Wong et al. (2007), V_4258_LC0863_Wong et al. (2007), V_4259_LC0863_Wong et al. (2007), V_4260_LC0878_Wong et al. (2007), V_4261_LC0907_Wong et al. (2007)
8	p12-q24.3	35.07	146.21	111.140043	27%	TN.10646, TN.10662, TN.10665, TN.11093, TN.12303, TN.12523, TN.1267, TN.3319, TN.4643, TN.B18, TN.B6, TN.B62, TN.B78, TN.B81, TN.BC184, TN.BC2697, TN.BC3045, TN.BC3047, TN.NT0001, TN.NT0015, TN.NT0024, TN.NT0039, TN.NT004, TN.NT005, TN.NT0067	UNC5D, FKSG2, ZNF703, SPFH2, PROSC, GPR124, BRF2, RAB11FIP1, GOT1L1, ADRB3, EIF4EBP1, ASH2L, STAR, LSM1, BAG4, DDHD2, PPAPDC1B, WHSC1L1, LETM2, FGFR1, FLJ43582, TACC1, HTRA4, TM2D2, ADAM9, ADAM32, ADAM18, ADAM2, INDO, C8orf4, ZMAT4, SFRP1, GOLGA7, SLD5, AGPAT6, FLJ25169, ANK1, MYST3, AP3M2, PLAT, IKBKB, POLB, DKK4, VDAC3, SLC20A2, C8orf40, CHRNB3, CHRNA6, THAP1, RNF170, HOOK3, FNTA, FLJ23356, POTE8, KIAA0146, CEBPD, PRKDC, MCM4, UBE2V2, EFCAB1, SNAI2, C8orf22, SNTG1, PXDNL, PCMTD1, ST18, UNQ9433, RB1CC1, NPBWR1, OPRK1, ATP6V1H, RGS20, TCEA1, LYPLA1, MRPL15, SOX17, RP1, XKR4, TMEM68, NCOA6IP, LYN, RPS20, MOS, PLAG1, CHCHD7, RDHE2, PENK, IMPAD1, C8orf72, LOC137886, CYP7A1, SDCBP, NSMAF, TOX, CA8, RAB2, CHD7, MGC34646, ASPH, FAM77D, FLJ39198, GGH, TTPA, YTHDF3, BHLHB5, CYP7B1, ARMC1, MTFR1, PDE7A, DNAJC5B, TRIM55, CRH, RRS1, ADHFE1, C8orf46, MYBL1, VCPIP1, C8orf44, SGK3, C8orf45, LOC286187, COPS5, CSPP1, ARFGEF1, CPA6, DEPDC2, C8orf34, SULF1, SLCO5A1, PRDM14, NCOA2, TRAM1, LACTB2, XKR9, EYA1, MSC, TRPA1, KCNB2, TERF1, RPESP, RPL7, RDH10, STAU2, UBE2W, TCEB1, TMEM70, LY96, JPH1, GDAP1, PI15	hsa.mir.486, hsa.mir.124.2, hsa.mir.599, hsa.mir.875, hsa.mir.1273, hsa.mir.548a.3, hsa.mir.548d.1, hsa.mir.1205, hsa.mir.1206, hsa.mir.1207, hsa.mir.1208, hsa.mir.30b, hsa.mir.30d, hsa.mir.151, hsa.mir.1302.7, hsa.mir.937, hsa.mir.661, hsa.mir.939, hsa.mir.1234	V_0119_LC10711_Iafrate et al. (2004), V_0121_LC10727_Iafrate et al. (2004), V_0123_LC10906_Iafrate et al. (2004), V_0124_LC10922_Iafrate et al. (2004), V_2126_LC10727_Locke et al. (2006), V_4589_LC10531_Wong et al. (2007), V_4590_LC10569_Wong et al. (2007), V_4591_LC10578_Wong et al. (2007), V_4592_LC10583_Wong et al. (2007), V_4593_LC10587_Wong et al. (2007), V_4594_LC10611_Wong et al. (2007), V_4595_LC10619_Wong et al. (2007), V_4596_LC10633_Wong et al. (2007), V_4597_LC10634_Wong et al. (2007), V_4598_LC10640_Wong et al. (2007), V_4599_LC10697_Wong et al. (2007), V_4600_LC10722_Wong et al. (2007), V_4601_LC10727_Wong et al. (2007), V_4602_LC10727_Wong et al. (2007), V_4603_LC10727_Wong et al. (2007), V_4604_LC10727_Wong et al. (2007), V_4605_LC10727_Wong et al. (2007), V_4606_LC10731_Wong et al. (2007), V_4607_LC10749_Wong et al. (2007), V_4608_LC10811_Wong et al. (2007), V_4609_LC10877_Wong et al. (2007), V_4610_LC10912_Wong et al. (2007), V_4611_LC10941_Wong et al. (2007), V_4612_LC10994_Wong et al. (2007), V_4613_LC11006_Wong et al. (2007), V_4614_LC11012_Wong et al. (2007), V_4791_LC10979_Wong et