

## Annotations for all recurrently gained regions

cytoband	1q25.1	1q25.1 (continued)	1q31.3	3q27.3	6q15	7p12.2
q value		6,38E-06		7,93E-06	0,057707	0,095294
residual q value		6,38E-06		0,012405	0,057707	0,095294
wide peak boundaries	chr1:166155454-171863618		chr1:195253751-195672280	chr3:188956472-189175357	chr6:91249512-91370379	chr7:49995112-50127527
genes in wide peak	FASLG	<b>U81</b>	PTPRC	[BCL6]	MAP3K7	IKZF1
	SERPINC1	<b>U47</b>	<b>hsa-mir-181b-1</b>			
	F5	<b>U80</b>	<b>hsa-mir-213</b>			
	FMO1	<b>U79</b>	<b>hsa-mir-213</b>			
	FMO2	<b>U78</b>				
	FMO3	<b>U44</b>				
	FMO4	<b>U77</b>				
	MYOC	<b>U76</b>				
	PIGC	<b>U75</b>				
	PRRX1	<b>U74</b>				
	SELE					
	SELL					
	SELP					
	TNFSF4					
	VAMP4					
	TNFSF18					
	GPR52					
	PRDX6					
	KIAA0040					
	RABGAP1L					
	SLC19A2					
	KIFAP3					
	BAT2D1					
	DNM3					
	CACYBP					
	KLHL20					
	C1orf9					
	KIAA0859					
	DARS2					
	C1orf112					
	SCYL3					
	TNN					
	MRPS14					
	C1orf129					
	ZBTB37					
	CENPL					
	C1orf156					
	SCYL1BP1					
	C1orf105					
	RC3H1					
	SLC9A11					
	ANKRD45					
	<b>hsa-mir-214</b>					
	<b>hsa-mir-199a-2</b>					
	<b>hsa-mir-199a-2</b>					

7q34	8q24.13	11q23.3	11q23.3 (Continued)	11q24.3	12q15	12q15 (Continued)
	0,10948	0,16127	0,10948		0,011853	0,0050948
	0,10948	0,16127	0,10948		0,011853	0,0050948
chr7:141162879-142244396	chr8:121784261-129821355	chr11:115828212-118580638		chr11:127825348-127931093	chr12:61724591-82263925	
EPHB6	ANXA13	APOA1	FLJ11783	ETS1	AVPR1A	TMEM19
KEL	HAS2	APOA4	TMPRSS13		CPM	SLC35E3
PRSS1	MYC	APOC3	BUD13		CSRP2	IL26
PRSS2	NDUFB9	ARCN1	TMEM25		GNS	CAND1
MGAM	SNTB1	FXYD2	APOA5		IFNG	MDM1
TRPV6	SQLE	BLR1	AMICA1		KCNC2	NUP107
TRPV5	MTSS1	CD3D	LOC143941		LYZ	SRGAP1
OR9A2	KIAA0196	CD3E	LOC196264		MDM2	TBC1D15
C7orf34	TRIB1	CD3G	DKFZp547C195		MYF5	FLJ21963
TRY1	RNF139	DDX6	BCL9L		MYF6	BBS10
	ZHX1	DPAGT1	FOXR1		PPP1R12A	THAP2
	ZHX2	SLC37A4	LOC283152		NAP1L1	C12orf26
	ATAD2	H2AFX	CCDC84		CNOT2	C12orf31
	TRMT12	HMBS			PAWR	CAPS2
	C8orf32	IL10RA			PTPRB	NAV3
	DERL1	MLL			PTPRR	HELB
	TATDN1	PAFAH1B2			RAP1B	OSBPL8
	C8orf76	RPS25			SYT1	RAB3IP
	FAM83A	SCN2B			TSPAN8	TPH2
	WDR67	SCN4B			YEATS4	GLIPR1L2
	FBXO32	TAGLN			HMGA2	BEST3
	ZNF572	UPK2			DYRK2	E2F7
	C8orf78	ZNF259			PPFIA2	FLJ32549
	TMEM65	PCSK7			LGR5	TMTC2
	FAM84B	UBE4A			LIN7A	CCDC131
	FAM91A1	TMEM24			TMEM5	C12orf28
	NSMCE2	EVA1			CCT2	MSRB3
	C8orf54	HYOU1			FRS2	GLIPR1L1
	TMEM75	ATP5L			GLIPR1	FLJ90579
	C8ORFK23	TREH			CPSF6	RASSF3
		CEP164			KRR1	LOC283392
		PHLDB1			WIF1	DPY19L2
		KIAA0999			IRAK3	LRRC10
		BACE1			XPOT	
		MIZF			PHLDA1	
		SIDT2			RAB21	
		TRAPPC4			ZDHHHC17	
		FXYD6			GRIP1	
		VPS11			LEMD3	
		TMPRSS4			KCNMB4	
		C11orf60			CCDC59	
		DSCAML1			TBK1	
		ABCG4			TRHDE	
		NLRX1			IL22	
		PDZD3			TMBIM4	

12q24.33	13q31.3	17p11.2	18q21.2
	0,19182	1,16E-05	0,066881
	0,19182	1,16E-05	0,066881
chr12:128465362-132449811	chr13:90362259-90843579	chr17:18755702-20858373	chr18:49301605-55260237
STX2	[GPC5]	ALDH3A1	DCC
GOLGA3	<b>hsa-mir-17</b>	ALDH3A2	FECH
MMP17	<b>hsa-mir-17</b>	MFAP4	GRP
POLE	<b>hsa-mir-18a</b>	MAPK7	LMAN1
PXMP2	<b>hsa-mir-18a</b>	PRPSAP2	NARS
RAN	<b>hsa-mir-19a</b>	ZNF179	ATP8B1
SFRS8	<b>hsa-mir-20a</b>	ULK2	RAB27B
ZNF10	<b>hsa-mir-19b-1</b>	GRAP	TCF4
ZNF26	<b>hsa-mir-92-1</b>	AKAP10	MBD2
ZNF84		EPN2	TXNL1
ZNF140		EPPB9	ONECUT2
ULK1		FLJ10847	MALT1
PIWIL1		SPECC1	POLI
ZNF268		SLC5A10	NEDD4L
FZD10		FLJ31196	WDR7
P2RX2		LOC284194	RAX
RIMBP2		MGC87631	ST8SIA3
GALNT9		FAM83G	ZNF532
CHFR		<b>U3-2</b>	CCDC68
EP400		<b>U3-2</b>	SEC11C
NOC4L		<b>U3-4</b>	HAK
PUS1		<b>U3</b>	STARD6
ZNF605		<b>U3-3</b>	CCBE1
TMEM132D		<b>ACA59</b>	C18orf54
PGAM5			C18orf26
GPR133			CPLX4
DDX51			FLJ45743
LOC338797			<b>ACA37</b>
EP400NL			<b>hsa-mir-122a</b>
<b>ACA49</b>			