

Supplemental Table 2. Recurrent regions of copy number gain and loss together with the candidate genes located within the regions.

A. Regions of gain.

Region	Cytoband	Start	End	Candidate genes
G1	1p36.33	1,688,462	1,960,160	<i>CDK11A/B, CALML6, NADK</i>
G2	1q21.1	146,023,079	149,331,955	<i>NBPF10, NOTCH2NL</i>
G3	1q21.3	150,542,525	150,594,423	<i>MCL1</i>
G4	1q22	156,189,235	156,298,108	<i>BGLAP, SMG5</i>
G5	1q23.3	160,642,011	160,825,109	<i>SLAMF7, CD48, LY9, SLAMF1</i>
G6	1q24.2	167,212,297	167,970,941	<i>POU2F1, CREG1, DCAF6</i>
G7	1q25.2	178,226,546	179,305,784	<i>RALGPS2, ABL2, RASAL2</i>
G8	1q32.1	203,246,823	203,377,011	<i>BTG2, FMOD</i>
G9	1q44	247,158,599	248,946,422	<i>NLRP3</i>

B. Regions of deletion.

Region	Cytoband	Start	End	Candidate genes
D1	1p36.33	1	1,808,460	<i>TNFRSF4, DVL1, CDK11A/B, CCNL2, TNFRSF18, MIB2</i>
D2	1p35.3	29,315,089	31,907,425	<i>SDC3, PUM1</i>
D3	1p33	50,183,629	51,109,627	<i>FAF1, CDKN2C</i>
D4	1p31.1	70,904,779	74,501,017	<i>NEGR1, PTGER3</i>
D5	1p31.1	76,627,494	87,337,216	<i>BCL10, FUBP1, PRKACB, PTGFR, DNAJB4</i>
D6	1p22.1	92,383,744	93,156,082	<i>RPL5, EVI5, GFI1, MTF2</i>
D7	1p12	117,501,979	117,869,477	<i>TENT5C</i>