

Table S2: Recurrent^a Low Level Copy Number Gains and Hemizygous Deletions

<u>Gains</u>			
<u>Cytoband</u>	<u>Genome Coordinates</u>	<u>Mean Copy Number^b</u>	<u>Candidate Genes</u>
1q21.1	144634354-144966203	4.475	GPR89A, GPR89B, GPR89C, PDZK1P1
7q34	139089210-139160099	3.396	TBXAS1
7q34	139821372-139834068	3.404	ADCK2
7q36	149654957-149880273	4.584	GIMAP7, GIMAP4, GIMAP2, GIMAP1, GIMAP5
7q36.1	150187117-150189549	3.499	ACCN3, CDK5
8q24.3	141931973-142023392	3.320	PTK2
<u>Deletions</u>			
<u>Cytoband</u>	<u>Genome Coordinates</u>	<u>Mean Copy Number</u>	<u>Candidate Genes</u>
1p36.31	6092828-6096849	1.059	KCNAB2, CHD5
10q26.12	122616028-12265495	0.949	BRWD2
10q26.2	129049838-12914006	1.043	DOCK1
19p13.3	1171321-1172545	0.778	STK11

^a Recurrent = 3 or more cases

^b Mean values determined from tumors with copy number increases at each of the indicated loci.