Supplementary Table 1: 14q13.3 amplicon-associated gains/losses

Cytoband[a]	Gain/loss	Selected resident cancer genes
1q21.1	gain	
1q21.2	gain	
1q21.3	gain	
2p23.2	gain	
2p23.1	gain	
5p15.33	gain	*TERT*
5p15.32	gain	
5p15.31	gain	
5p15.2	gain	
5p15.1	gain	
8p23.3	loss	
8p23.2	loss	
8p23.1	loss	
8p22	loss	
14q11.2	gain	
14q12	gain	
14q13.1	gain	
14q13.2	gain	
14q13.3	gain	*TITF1*
14q21.1	gain	
14q21.2	gain	
14q21.3	gain	
14q22.1	gain	
14q22.2	gain	
16p13.13	gain	
17q23.2	gain	*RPS6KB1* , *PPM1D*

[a]To detect associations between DNA copy number alterations at distinct loci, a Pearson's correlation was computed between the mean copy number of a given cytoband and that of all other cytobands. Statistically significant correlations were determined by randomly permuting cytoband labels and recalculating correlations 100 times; a false discovery rate (FDR) of 1% was used to establish a significance threshold. Positive associations with 14q13.3 (*TITF1*) are shown. Note, positive associations elsewhere on 14q reflect DNA segments co-amplified with 14q13.3.