Figure S2. The signature associated with Kataegis exhibits a broad range of context sensitivity. (A) Non-normalized mutation frequencies for Kataegis using a tri-nucleotide signature. Mutation frequency is given on the verticle axis. (B) Mutation frequencies for Kataegis using a penta-nucleotide signature (normalized to the penta-nucleotide frequency distribution of the exome). The penta-nucleotide signature is represented as a stack of tri-nucleotide frequency profiles, with the flanking 5’ and 3’ nucleotides (positions -2 and +2 relative to the point mutation) tabulated adjacentlly. The stack of tri-nucleotide profiles was ordered by hierarchical clustering (see the corresponding distance dendrogram shown on the left side).