Figure S3. The signature associated with Kataegis exhibits a broad range of context sensitivity. 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 each row corresponding to a tri-nucleotide frequency profile for a particular combination of flanking 5’ and 3’ nucleotides. Identities of the flanking nucleotides (positions -2 and +2 relative to the point mutation) are tabulated in the columns on the right. The stack of tri-nucleotide profiles was ordered by hierarchical clustering (dendrogram not shown).