

Additional file 2: Scatterplot showing the positions of high frequency (AAF $\geq$ 0.9) frameshift InDels (in X axis) located at beginning of cDNA and of the nearest downstream ATG start codon (in Y axis). Both the start codon and InDel positions are represented as relative to the cDNA lengths. Data from 72 FS InDels which were located within 0.1 length of cDNA have been used to create the graph.

