

Supplementary Figure 1 Legend: Multispecies sequence alignment of the MCS-9.7 segment corresponding to nucleotides from –14470 to –14654 with respect to *IRF6* translation initiation site, that contains four putative AP-2 α binding sites (highlighted in yellow) predicted by the TESS software <http://www.cbil.upenn.edu/tess>. Variants -14474A>G, rs642961 (G>A) and -14523G>A, are indicated by vertical arrows.