Figure S1: Overview of the RNAsnp results predicted for 29,290 SNVs. a) Distribution of the p-values calculated for 29,290 using $d_{\text{max}}$ and $r_{\text{min}}$ measures of RNAsnp. The dotted line indicates the threshold value of 0.1. b) Distribution of the distance value between the SNV position and the predicted local region c) Length distribution of the predicted local region.