



Supplementary Figure 1. Venn-diagram presentation of common prediction binding sites on CHIKV negative sense RNA genome of each genotype/lineage. miRNAs that were sorted by the conservation of target sites, might show a more general miRNA-vRNA interaction. IOL, Indian Ocean Lineage, AM258992; WA, West African, HM045816; Asian, EU703762; ECSA, East/Central/South African, HM045811.