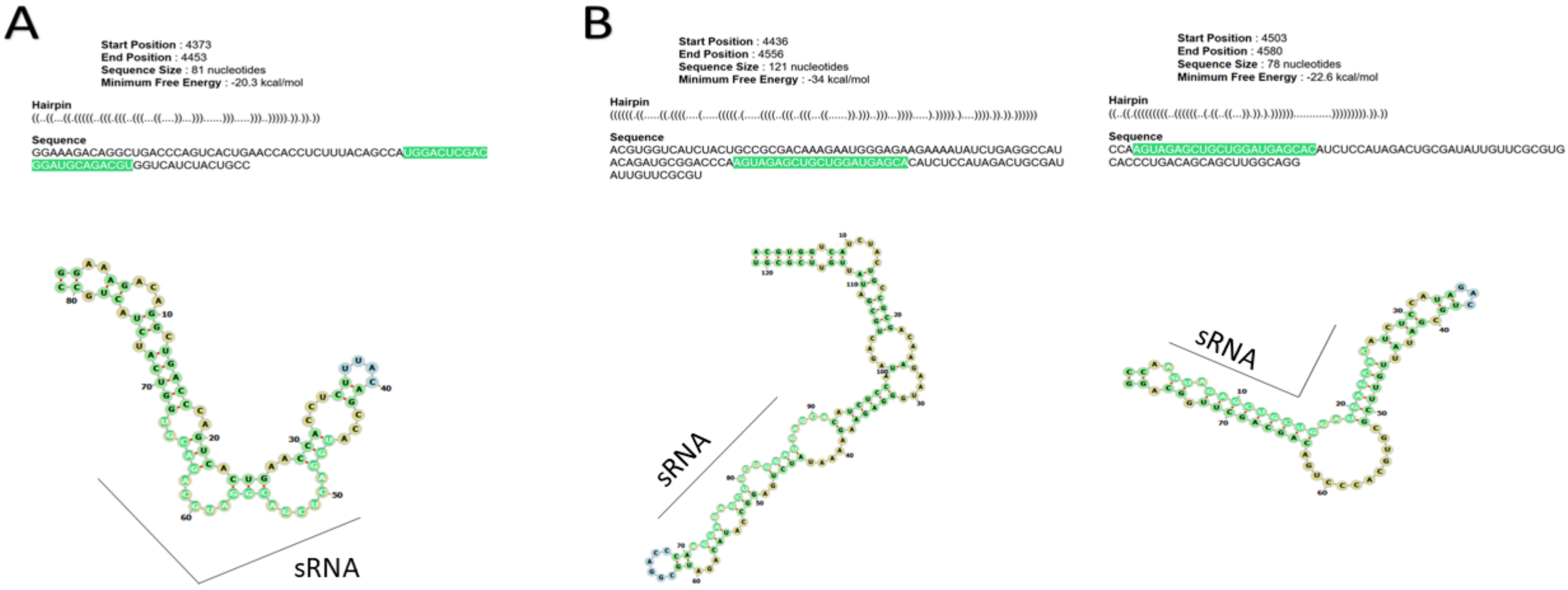Supplemental figure 4. Hairpin predictions computed with miRNAFold. Left: Window between 4373-4453 of the CHIKV 06.21 genome. Candidate small RNA sequence highlighted in green. Right: Two windows (4436-4556 and 4503-4580) for a same candidate viral miRNA sequence.