



Supplementary Material

Short Communication: *In Silico* Profiling of Regulatory MicroRNA Targets in Programmed Cell Death 1 Gene

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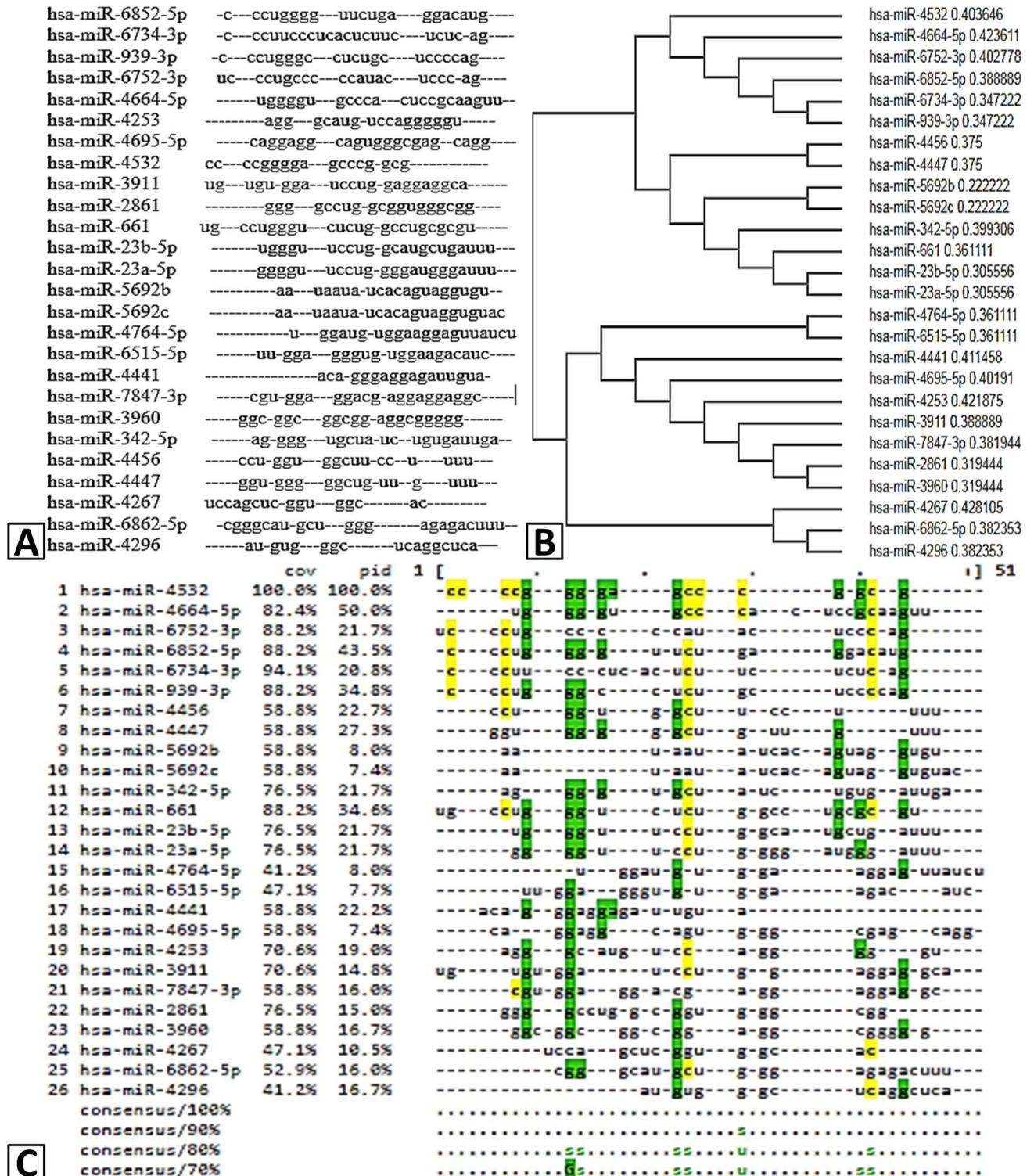
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Supplementary Fig. 1. **A**, represents multiple sequence alignment of 26 selected miRNA genes. **B**, phylogenetic tree analysis of 26 miRNA by using Clustal omega v (1.2.4). Branching represents similarity profile of each miRNA with others. **C**, M view of MSA analysis of 26 miRNA using Clustal omega v (1.2.4). Green represents purine and yellow represents pyrimidine. Cov, covariance; pid, predicted identity.