

## SUPPLEMENTARY INFORMATION

### **Elucidation of the Molecular Interaction between miRNAs and the *HOXA9* Gene, Involved in Acute Myeloid Leukemia, by the Assistance of Argonaute Protein through a Computational Approach**

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**Supplementary Table 1.** miRNA target sites in *HOXA9* gene are highlighted using different color

NCBI accession No.	Organism	Gene name	<i>HOXA9</i> DNA sequence
NM_152739.3	<i>Homo sapiens</i>	HomeoboxA9 ( <i>HOXA9</i> ), mRNA	<p>AGTTGTTACATGAAATCTGCAGTTTCATAATTTCCGTGGGTCGGGCCGGG  CGGGCCAGGCGCTGGGCACGGT<b>ATGGCCACC</b>ACTGGGGCCCTGGGGCAAC  <b>TACTACGTGGACTCGTTCC</b>TGCTGGGGCGCCGACGCCGCGATGAGCTGAG  CGTTGGCCGCTATGCGCCGGGGACCCTGGGCCAGCCTCCCCGGCAGGCGG  CGACGCTGGCCGAGCACCCCGACTTCAGCCCGTGCAGCTTCCAGTCCAAG  GCGACGGTGT<b>TTGGCGCCTCGT</b>GGAACCCAGTGCACGCGGGCGGGCGCCAA  CGCTGTACCCGCTGCGGTGTACCACCACCATCACCACCACCCCTACGTGC  ACCCCCAGGCGCCCGTGGCGGGCGGGCGCCGGACGGCAGGTACATGCGC  TCCTGGCTGGAGCCCACGCCCGGTGCGCTCTCCTTCGCGGGCTTGCCCTC  CAGCCGGCCTTATGGCATTAAACCTGAACCGCTGTCCGCCAGAAGGGGTG  ACTGTCCACGCTTGACACTCACACTTTGTCCCTGACTGACTATGCTTGT  GGTTCTCCTCCAGTTGATAGAGAAAAACAACCCAGCGAAGGCGCCTTCTC  TGAAAACAATGCTGAGAATGAGAGCGGGGAGACAAGCCCCCATCGATC  CCAATAACCCAGCAGCCA<b>ACTGGCTTCATGCGCGCTCCACTCGGAAAAAG</b>  CGGTGCCCTATACAAAACACCAGACCC<b>TGGA</b>ACTGGAGAAAGATTTCT  GTTCAACATG<b>TACCTCA</b>CCAGGGACCGCA<b>GGTACGA</b>GGTGGCTCGACTGC  TCAACCTCACCGAGAGG<b>CAGGTCAAG</b>ATCTGGTTCCAGAACCGCAGGATG  AAAATGAAGAAAATCAACAAAGACCGAGCAAAAGACGAGTGATGCCATTT  GGGCTTATTTAGAAAAAAGGGTAAGCTAGAGAGAAAAAGAAAGAACTGTC  CGTCCCCCTTCCGCTTCTCCCTTCTCTCACCCCCACCCTAGCCTCCACC  ATCCCCGCACAAAGCGGCTCTAAACCTCAGGCCACATCTTTTCCAAGGCA  AACCTGTTCAGGCTGGCTCGTAGGCTGCCGCTTTGATGGAGGAGGTAT  TGTAAGCTTCCATTTTCTATAAGAAAAAGGAAAGTTGAGGGGGGGGCA  TTAGTGCTGATAGCTGTGTGTGTTAGCTTGTATATATATTTTTAAAAATC  TACCTGTTCTGACTTAAAACAAAAGGAAAGAA<b>ACTACCTT</b>TTTATAATG  CACAAC<b>TGTGATGGTAGGCTGTATAGTTTTTAGTCTGTGTAGTTAATTT</b>  AATTTGCAGTTTGTGCGGAGATTGCTCTGCCAAGATACTTGAACACTGT  GTTTTATTGTGGTAATTATGTTTTGTGATTCAAAC<b>TCTGTG</b>ACTGGGT  GATGCACCCATTGTGATTGTGGAAGATAGAATTC<b>AATTTGA</b>ACTCAGGTT  GTTTATGAGGGGAAAAAACAGTTGCATAGAGTATAGCTCTGTAGTGGAA  TATGTCTTCTGTATAACTAGGCTGTTAACCTATGATGTAAAGTAGCTGT  AAGAATTTCCAGTGAATAAAAAAAAAATTTAAGTGTCTCGGGGATGC  ATAGATTCATCATTTTCTCCACCTTAAAAATGCGGGCATTTAAGTCTGTC  CATTATCTATATAGTCTGTCTTGTCTATTTGTATATATAATCTATATGAT  TAAAGAAAATATGCATAATCAGACAAGCTTGAATATTTGTTTTGCACCAG  ACGAACAGTGAGGAAATTCGGAGCTATACATATGTGCAGAAAGTT<b>ACTAC</b>  <b>CT</b>AGGGTTTATGCTTAATTTTAATTTGGAGGAAATGAATGCTGATTGTAAC  GGAGTTAATTTTATTGATAATAAATTATACACTATGAAACCGCCATTGGG  CTACTGTAGATTTGATCCTTGATGAATCTGGGGTTTCCATCAGACTGAA  CTTACACTGTATATTTTGAATAGTTACCTCAAGGCC<b>TACTGAC</b>CAAAT  GTTGTGTTGAGATGATATTTAACTTTTTTCCAAATAAAATATATTGATTC  TTTTCTAAAAAAAAAAAAAAAAAAAA</p>

mir-145, yellow; mir-126, sky; let-7a, pink; mir-196b.1, green; mir-196b.2, violet. Coding portion of *HOXA9* is shown in red color. Two binding sites predicted for mir-196b in *HOXA9* represented as mir-196b.1 and mir-196b.2, respectively.