

Table S4 Candidate miRNAs binding to circ_0007386 predicted by CircInteractome.

CircRNA Mirbase ID	CircRNA (Top) - miRNA (Bottom) pairing	Site Type
hsa_circ_0007386 (5' ... 3')	<pre> CUC AAGUCAGACTUACUCAGAU CCUUCGGUAGUUCAGC-GACGUCC </pre>	7mer-1a
hsa-miR-1184 (3' ... 5')	<pre> CUC AAGUCAGACTUACUCAGAU GAGUUUCGUUUUG--GACGUCC </pre>	7mer-1a
hsa_circ_0007386 (5' ... 3')	<pre> GUGCAACCCCGCAGGCUGUCUGC CUUUCACGUADCUUAGACAGACC </pre>	7mer-m8
hsa-miR-1205 (3' ... 5')	<pre> UUUUUUUUUUUAGAGA---AGCCAGAU CCUCACUUCUGUCCUUGGUUU </pre>	7mer-1a
hsa_circ_0007386 (5' ... 3')	<pre> G AUGGUUGCUGUACUUUGCCAAC UCACACUCAAGAGGUUACGGUUU </pre>	7mer-m8
hsa-miR-182 (3' ... 5')	<pre> UCCUGAAGAUUCUGUUCUGAUCG UCGGUGUUAGUGGAAGACTAGA </pre>	7mer-m8
hsa_circ_0007386 (5' ... 3')	<pre> UCGAGGGUUUUGUCUCCUCCUGGG UCCCGGAGUCGGAGGACCA </pre>	7mer-m8
hsa-miR-665 (3' ... 5')	<pre> UUUUUUUUUUUUUUUUUUUUUUUUUUUUUU GUGUACCGGUUUUGUCUCUUCU </pre>	7mer-m8
hsa_circ_0007386 (5' ... 3')	<pre> AGGAGAGCCAGAUUCUCUCCAGG GGUGGGUUCACUGGAUGAGGUUCU </pre>	7mer-m8
hsa-miR-432 (3' ... 5')	<pre> CUGUGACCCUCUAGGAGUGCAAAC AAGUGAGGUUUUUCCAGGUUUU </pre>	7mer-m8
hsa_circ_0007386 (5' ... 3')	<pre> CUGUGACCCUCUAGGAGUGCAAAC UCUGUUCUCCGGUGGGCAGGUUUG </pre>	7mer-m8
hsa-miR-507 (3' ... 5')	<pre> UUUCUCCAGGGCCGUCUGAAG UACUGUGAGGGACACUU </pre>	7mer-1a
hsa_circ_0007386 (5' ... 3')	<pre> CUCGAGAUUGGUUCUGUACUUUG UUGUUUUAGCGGUUCAUGAAAA </pre>	7mer-m8
hsa-miR-513a-5p (3' ... 5')	<pre> UUGGGAGUGCUGUCCUUAACCA UAGUCUCCAAGUUGAAUGGGG </pre>	8mer-1a
hsa_circ_0007386 (5' ... 3')	<pre> CAGCCGGGAACCUU---AACAUACU CAAGACAGGACCUUCUUUAUGA </pre>	7mer-m8
hsa-miR-567 (3' ... 5')		