

5' CCA GUA A CUG GC A GGC AGGA AGAG G A G C C C U G G G G C U G A G G U G A G G
3' C C G G G G A C C G G C G G C U U U C C A C C C G G A C U C U G G C C C U U G G U A



hsa-miR-671-3p

[illegible]