

Description of Supplementary Files

File name: Supplementary Data 1

Description: 61 cricRNA candidates with highest reads supporting circular junction.

File name: Supplementary Data 2

Description: The Probability of Interaction by Target Accessibility (PITA) algorithm predicts miRNA binding sites on circBIRC6

File name: Supplementary Data 3

Description: The microarray analysis of splicing factors expression in hESC H9, hiPSC CFB50 and its parental cells (HF). (Fold change > 5)