

Supplementary Table 2. The predicted AGO2 binding sites in hsa_circ_0007444.

CircRNA	Tag Name	% Identity	Alignment Length	Mismatches	Gap Openings	Tag Start	Tag End	CircRNA Start	CircRNA End
hsa_circ_0007444	HHFCT_129104_cluster -18149_13_51_64	100	64	0	0	1	64	129	192
hsa_circ_0007444	HHFKP_118951_cluster -18149_10_37_43	100	43	0	0	1	43	131	173
hsa_circ_0007444	HHFKP_118952_cluster -18149_12_103_36	100	36	0	0	1	36	212	247
hsa_circ_0007444	HHFCT_129105_cluster -18149_14_24_38	100	38	0	0	1	38	232	269
hsa_circ_0007444	HHFCT_129106_cluster -18149_15_20_49	100	49	0	0	1	49	306	354
hsa_circ_0007444	HHFKP_118950_cluster -18149_8_596_59	100	59	0	0	1	59	30	88
hsa_circ_0007444	HHRBL_11489_BCBL 1-2of3-SCL_9227_95	100	95	0	0	1	95	33	127
hsa_circ_0007444	HHFCT_129102_cluster -18149_11_34_93	100	93	0	0	1	93	3	95