

Table SIII. Predictions for RNA binding proteins of the hsa_circ_0000520, hsa_circ_0006220, hsa_circ_0000977 and hsa_circ_0043278 using the cancer-specific circRNA database.

circRNA ID	Chrom	RBP	Start	End
hsa_circ_0000520	chr14	AGO2_Human_GSE28865_HITS-CLIP	2081144 3	2081148 3
hsa_circ_0000520	chr14	AGO2_Human_GSE28865_PAR-CLIP	2081152 2	2081157 0
hsa_circ_0000520	chr14	FMRP_Human_GSE39682_PAR-CLIP	2081142 5	2081144 6
hsa_circ_0000520	chr14	AGO1_Human_GSE43573_PAR-CLIP	2081153 6	2081157 0
hsa_circ_0000520	chr14	AGO1_Human_GSE43573_PAR-CLIP	2081149 7	2081157 0
hsa_circ_0000520	chr14	AGO2_Human_GSE21918_PAR-CLIP	2081154 8	2081156 8
hsa_circ_0000520	chr14	AGO3_Human_GSE21918_PAR-CLIP	2081154 6	2081157 0
hsa_circ_0000520	chr14	AGO2_Human_GSE28865_HITS-CLIP	2081148 3	2081152 3
hsa_circ_0000520	chr14	AGO2_Human_GSE28865_PAR-CLIP	2081152 5	2081157 0
hsa_circ_0000520	chr14	LIN28A_Human_GSE44615_PAR-CLIP	2081141 8	2081145 3
hsa_circ_0000520	chr14	UPF1_Human_GSE47976_iCLIP	2081144 5	2081144 6
hsa_circ_0000520	chr14	TNRC6_Human_GSE21918_PAR-CLIP	2081147 6	2081157 2

hsa_circ_000052 0	chr14	DGCR8_Human_GSE39086_HITS-CLIP	2081145 0	2081157 1
hsa_circ_000052 0	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081140 1	2081143 9
hsa_circ_000052 0	chr14	HuR_Human_GSE28865_PAR-CLIP	2081148 3	2081152 5
hsa_circ_000052 0	chr14	AGO3_Human_GSE21918_PAR-CLIP	2081149 7	2081152 8
hsa_circ_000052 0	chr14	HuR_Human_GSE28865_PAR-CLIP	2081147 9	2081152 2
hsa_circ_000052 0	chr14	CAPRIN1_Human_GSE38201_PAR-CLIP	2081148 2	2081152 0
hsa_circ_000052 0	chr14	AGO2_Human_GSE21918_PAR-CLIP	2081149 7	2081152 0
hsa_circ_000052 0	chr14	AGO2_Human_GSE28865_HITS-CLIP	2081144 3	2081148 3
hsa_circ_000052 0	chr14	C22ORF28_Human_GSE38201_PAR-CLIP	2081153 1	2081157 0
hsa_circ_000052 0	chr14	FUS_Human_GSE43308_HITS-CLIP	2081149 4	2081157 0
hsa_circ_000052 0	chr14	AGO2_Human_GSE28865_HITS-CLIP	2081148 3	2081152 3
hsa_circ_000052 0	chr14	U2AF65_Human_E-MTAB-1371_iCLIP	2081150 6	2081151 2
hsa_circ_000052 0	chr14	FXR2_Human_GSE39682_PAR-CLIP	2081149 8	2081152 0
hsa_circ_000052 0	chr14	FMRP_Human_GSE39682_PAR-CLIP	2081147 9	2081157 0
hsa_circ_000052	chr14	UPF1_Human_GSE47976_iCLIP	2081149	2081149

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hsa_circ_000052	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081152	2081157
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hsa_circ_000052	chr14	HuR_Human_GSE28865_HITS-CLIP	2081152	2081156
0			9	9
hsa_circ_000052	chr14	FUS_Human_GSE43308_HITS-CLIP	2081144	2081147
0			8	4
hsa_circ_000052	chr14	HuR_Human_GSE28865_PAR-CLIP	2081152	2081156
0			9	9
hsa_circ_000052	chr14	DGCR8_Human_GSE39086_HITS-CLIP	2081140	2081146
0			0	2
hsa_circ_000052	chr14	LIN28A_Human_GSE44615_PAR-CLIP	2081145	2081157
0			8	3
hsa_circ_000052	chr14	UPF1_Human_GSE47976_iCLIP	2081152	2081152
0			3	4
hsa_circ_000052	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081146	2081157
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hsa_circ_000052	chr14	U2AF65_Human_E-MTAB-1371_iCLIP	2081151	2081152
0			8	6
hsa_circ_000052	chr14	HuR_Human_GSE28865_PAR-CLIP	2081152	2081156
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hsa_circ_000052	chr14	FMRP_Human_GSE39682_PAR-CLIP	2081149	2081152
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hsa_circ_000052	chr14	HuR_Human_GSE28865_HITS-CLIP	2081148	2081152
0			5	7
hsa_circ_000052	chr14	eIF4AIII_Human_GSE40778_HITS-CLIP	2081122	2081157
0			6	3
hsa_circ_000052	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081140	2081146
0			4	3

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hsa_circ_000052 0	chr14	AGO2_Human_GSE28865_PAR-CLIP	2081153 3	2081157 0
hsa_circ_000052 0	chr14	FMRP_Human_GSE39682_PAR-CLIP	2081149 2	2081152 0
hsa_circ_000052 0	chr14	LIN28B_Human_GSE44615_PAR-CLIP	2081141 8	2081145 0
hsa_circ_000052 0	chr14	FMRP_Human_GSE39682_PAR-CLIP	2081148 6	2081151 3
hsa_circ_000052 0	chr14	AGO2_Human_GSE28865_PAR-CLIP	2081153 4	2081157 0
hsa_circ_000052 0	chr14	AGO1_Human_GSE43573_PAR-CLIP	2081149 7	2081153 0
hsa_circ_000052 0	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081140 4	2081146 3
hsa_circ_000052 0	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081152 2	2081157 0
hsa_circ_000052 0	chr14	DGCR8_Human_GSE39086_HITS-CLIP	2081152 3	2081157 4
hsa_circ_000052 0	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081143 0	2081157 0
hsa_circ_000052 0	chr14	DGCR8_Human_GSE39086_HITS-CLIP	2081146 8	2081157 1
hsa_circ_000052 0	chr14	LIN28B_Human_GSE44615_PAR-CLIP	2081147 4	2081157 3
hsa_circ_000052 0	chr14	AGO2_Human_GSE41357_HITS-CLIP	2081152 5	2081157 0
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0			6	2
hsa_circ_000052	chr14	PTB_Human_GSE42701_HITS-CLIP	2081152	2081154
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hsa_circ_000052	chr14	AGO2_Human_SupplementaryData_PAR-C	2081148	2081152
0		LIP	5	5
hsa_circ_000052	chr14	DGCR8_Human_GSE39086_HITS-CLIP	2081147	2081150
0			4	9
hsa_circ_000052	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081140	2081143
0			1	9
hsa_circ_000052	chr14	AGO2_Human_GSE28865_PAR-CLIP	2081150	2081153
0			4	1
hsa_circ_000052	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081143	2081157
0			0	0
hsa_circ_000052	chr14	AGO2_Human_GSE44404_HITS-CLIP	2081143	2081152
0			6	2
hsa_circ_000052	chr14	DGCR8_Human_GSE39086_HITS-CLIP	2081146	2081157
0			8	5
hsa_circ_000052	chr14	AGO1_Human_GSE21918_PAR-CLIP	2081154	2081157
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hsa_circ_000052	chr14	AGO2_Human_GSE21918_PAR-CLIP	2081149	2081152
0			7	0
hsa_circ_000052	chr14	eIF4AIII_Human_GSE40778_HITS-CLIP	2081129	2081157
0			7	3
hsa_circ_000052	chr14	U2AF65_Human_E-MTAB-1371_iCLIP	2081151	2081152
0			9	6
hsa_circ_000052	chr14	AGO2_Human_GSE41357_HITS-CLIP	2081152	2081157
0			5	0

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hsa_circ_000622 0	chr17	U2AF65_Human_E-MTAB-1371_iCLIP	3580075 1	3580075 2
hsa_circ_000622 0	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3580060 4	3580067 0
hsa_circ_000622 0	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3580060 4	3580067 0
hsa_circ_000622 0	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3580070 6	3580076 3
hsa_circ_000622 0	chr17	IGF2BP3_Human_GSE21918_PAR-CLIP	3580070 8	3580076 2
hsa_circ_000622 0	chr17	AGO2_Human_GSE28865_HITS-CLIP	3580066 5	3580070 5
hsa_circ_000622 0	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3580067 8	3580079 6
hsa_circ_000097 7	chr2	AGO2_Human_GSE42701_HITS-CLIP	1079555 5	1079557 8
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080884 7	1080886 5
hsa_circ_000097 7	chr2	hnRNPC_Human_E-MTAB-1371_iCLIP	1079703 8	1079704 8
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1080306 4	1080317 2
hsa_circ_000097 7	chr2	hnRNPC_Human_E-MTAB-1371_iCLIP	1080536 2	1080537 3
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080550	1080551

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7			0	1
hsa_circ_000097	chr2	UPF1_Human_GSE47976_iCLIP	1079792	1079792
7			5	6
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079653	1079655
7			4	7
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079783	1079784
7			4	0
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079801	1079801
7			0	7
hsa_circ_000097	chr2	FUS_Human_GSE43308_HITS-CLIP	1080099	1080101
7			6	5
hsa_circ_000097	chr2	UPF1_Human_GSE47976_iCLIP	1080739	1080739
7			0	1
hsa_circ_000097	chr2	hnRNPC_Human_E-MTAB-1371_iCLIP	1080423	1080425
7			7	7
hsa_circ_000097	chr2	FUS_Human_GSE43308_HITS-CLIP	1078483	1078487
7			7	1
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079793	1079794
7			5	2
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080792	1080792
7			1	7
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079778	1079779
7			6	7
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080199	1080200
7			6	3
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079908	1079911
7			3	6

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hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080676 3	1080676 6
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079491 0	1079492 4
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079835 7	1079836 6
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080317 4	1080318 6
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1078876 5	1078879 9
hsa_circ_000097 7	chr2	hnRNPC_Human_E-MTAB-1371_iCLIP	1079670 7	1079671 0
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1080737 3	1080740 3
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080734 3	1080734 9
hsa_circ_000097 7	chr2	AGO2_Human_GSE42701_HITS-CLIP	1079418 8	1079421 6
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079785 6	1079787 0
hsa_circ_000097 7	chr2	IGF2BP3_Human_GSE21918_PAR-CLIP	1079463 9	1079467 1
hsa_circ_000097 7	chr2	FMRP_Human_GSE39682_PAR-CLIP	1079463 7	1079466 2
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1078449 2	1078450 0
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080822	1080824

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7			0	8
hsa_circ_000097	chr2	hnRNP_C_Human_E-MTAB-1371_iCLIP	1079826	1079827
7			5	7
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079929	1079934
7			8	5
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079794	1079794
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hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079122	1079126
7			7	4
hsa_circ_000097	chr2	UPF1_Human_GSE47976_iCLIP	1079799	1079799
7			2	3
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079477	1079478
7			1	1
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079274	1079279
7			9	2
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079486	1079487
7			8	2
hsa_circ_000097	chr2	hnRNP_C_Human_E-MTAB-1371_iCLIP	1079744	1079745
7			2	3
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079462	1079467
7			9	3
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079401	1079405
7			3	8
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080196	1080198
7			6	4
hsa_circ_000097	chr2	AGO2_Human_GSE42701_HITS-CLIP	1079164	1079166
7			7	8

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hsa_circ_000097 7	chr2	hnRNP_C_Human_E-MTAB-1371_iCLIP	1080704 1	1080704 2
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079469 6	1079470 4
hsa_circ_000097 7	chr2	FUS_Human_GSE43308_HITS-CLIP	1080533 2	1080535 9
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079567 8	1079568 5
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079767 3	1079768 7
hsa_circ_000097 7	chr2	AGO2_Human_GSE41437_PAR-CLIP	1080283 5	1080286 1
hsa_circ_000097 7	chr2	hnRNP_C_Human_E-MTAB-1371_iCLIP	1080786 7	1080787 6
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1078447 4	1078449 8
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1080529 5	1080535 2
hsa_circ_000097 7	chr2	ZC3H7B_Human_GSE38201_PAR-CLIP	1079917 7	1079923 7
hsa_circ_000097 7	chr2	UPF1_Human_GSE47976_iCLIP	1079794 9	1079795 0
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1078474 6	1078480 3
hsa_circ_000097 7	chr2	UPF1_Human_GSE47976_iCLIP	1080320 8	1080320 9
hsa_circ_000097	chr2	UPF1_Human_GSE47976_iCLIP	1079842	1079842

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7			5	0
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1080568	1080574
7			6	6
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1080568	1080574
7			6	5
hsa_circ_000097	chr2	hnRNPC_Human_E-MTAB-1371_iCLIP	1078765	1078766
7			5	1
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079798	1079798
7			5	6
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080304	1080304
7			3	4
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080747	1080747
7			4	9
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079469	1079470
7			6	4
hsa_circ_000097	chr2	hnRNPC_Human_E-MTAB-1371_iCLIP	1080834	1080834
7			2	3
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080296	1080296
7			6	7
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080883	1080886
7			8	6
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079929	1079933
7			9	8
hsa_circ_000097	chr2	PTB_Human_GSE42701_HITS-CLIP	1078875	1078879
7			9	0
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079778	1079781
7			6	0

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hsa_circ_000097 7	chr2	hnRNP_C_Human_E-MTAB-1371_iCLIP	1080349 4	1080350 4
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079794 7	1079795 3
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1078738 9	1078741 8
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1080308 3	1080317 5
hsa_circ_000097 7	chr2	AGO2_Human_GSE42701_HITS-CLIP	1078647 3	1078651 9
hsa_circ_000097 7	chr2	FUS_Human_GSE43308_HITS-CLIP	1079764 7	1079767 9
hsa_circ_000097 7	chr2	UPF1_Human_GSE47976_iCLIP	1079466 3	1079466 4
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1078458 4	1078458 8
hsa_circ_000097 7	chr2	PTB_Human_GSE42701_HITS-CLIP	1078844 6	1078849 2
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079280 9	1079286 6
hsa_circ_000097 7	chr2	FMRP_Human_GSE39682_PAR-CLIP	1080311 0	1080313 0
hsa_circ_000097 7	chr2	UPF1_Human_GSE47976_iCLIP	1080739 1	1080739 2
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080728 7	1080728 8
hsa_circ_000097	chr2	hnRNP_C_Human_E-MTAB-1371_iCLIP	1080106	1080107

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7			5	4
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080730	1080730
7			2	6
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079938	1079939
7			6	1
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080505	1080505
7			5	7
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079802	1079804
7			3	2
hsa_circ_000097	chr2	hnRNP_C_Human_E-MTAB-1371_iCLIP	1080694	1080694
7			3	4
hsa_circ_000097	chr2	ALKBH5_Human_GSE38201_PAR-CLIP	1080851	1080853
7			0	8
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080741	1080743
7			2	3
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080746	1080746
7			4	7
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079780	1079780
7			7	8
hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079798	1079800
7			1	7
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1078450	1078452
7			2	0
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080799	1080800
7			2	4
hsa_circ_000097	chr2	UPF1_Human_GSE47976_iCLIP	1079792	1079792
7			3	4

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hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1080239 6	1080243 6
hsa_circ_000097 7	chr2	FUS_Human_GSE43308_HITS-CLIP	1079664 5	1079666 6
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079466 2	1079466 5
hsa_circ_000097 7	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079471 1	1079471 4
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hsa_circ_000097	chr2	UPF1_Human_GSE47976_iCLIP	1079796	1079797

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hsa_circ_000097	chr2	FMRP_Human_GSE39682_PAR-CLIP	1079463	1079466
7			6	7
hsa_circ_000097	chr2	HuR_Human_GSE28865_PAR-CLIP	1079462	1079466
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hsa_circ_000097	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1079103	1079106
7			5	5
hsa_circ_000097	chr2	AGO2_Human_GSE42701_HITS-CLIP	1079786	1079791
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7			6	9
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7			7	2
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7			3	3
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hsa_circ_000097 7	chr2	UPF1_Human_GSE47976_iCLIP	1079794 8	1079794 9
hsa_circ_000097 7	chr2	UPF1_Human_GSE47976_iCLIP	1080313 6	1080313 7
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hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1079608	1079608

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7			6	6
hsa_circ_000097	chr2	U2AF65_Human_E-MTAB-1371_iCLIP	1080755	1080755
7			4	7
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7			8	3
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7			6	0
hsa_circ_000097	chr2	UPF1_Human_GSE47976_iCLIP	1080606	1080606
7			8	9
hsa_circ_000097	chr2	hnRNPC_Human_E-MTAB-1371_iCLIP	1079665	1079666
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hsa_circ_000097 7	chr2	DGCR8_Human_GSE39086_HITS-CLIP	1078795 2	1078797 4
hsa_circ_000097 7	chr2	eIF4AIII_Human_GSE40778_HITS-CLIP	1080010 0	1080014 2
hsa_circ_000097 7	chr2	hnRNPC_Human_E-MTAB-1371_iCLIP	1080683 1	1080684 0
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CircRNA ID	Chrom	RBP	Start	End
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hsa_circ_004327 8	chr17	U2AF65_Human_E-MTAB-1371_iCLIP	3579807 7	3579808 5

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hsa_circ_004327 8	chr17	AGO1_Human_GSE43573_PAR-CLIP	3580066 7	3580068 9
hsa_circ_004327 8	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3579852 9	3579855 5
hsa_circ_004327 8	chr17	UPF1_Human_GSE47976_iCLIP	3579956 0	3579956 1
hsa_circ_004327 8	chr17	U2AF65_Human_E-MTAB-1371_iCLIP	3580075 1	3580075 2
hsa_circ_004327 8	chr17	hnRNPC_Human_E-MTAB-1371_iCLIP	3579825 1	3579826 1
hsa_circ_004327 8	chr17	U2AF65_Human_E-MTAB-1371_iCLIP	3580057 2	3580059 9
hsa_circ_004327 8	chr17	U2AF65_Human_E-MTAB-1371_iCLIP	3579956 0	3579956 1
hsa_circ_004327 8	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3580060 4	3580067 0
hsa_circ_004327 8	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3580060 4	3580067 0
hsa_circ_004327 8	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3580070 6	3580076 3
hsa_circ_004327 8	chr17	IGF2BP3_Human_GSE21918_PAR-CLIP	3580070 8	3580076 2
hsa_circ_004327 8	chr17	FUS_Human_GSE43308_HITS-CLIP	3579875 9	3579877 5
hsa_circ_004327 8	chr17	hnRNPC_Human_E-MTAB-1371_iCLIP	3579955 2	3579956 5
hsa_circ_004327	chr17	TIAL1_Human_SupplementaryData_iCLIP	3579803	3579805

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8			1	9
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8			5	5
hsa_circ_004327	chr17	U2AF65_Human_E-MTAB-1371_iCLIP	3580057	3580059
8			2	9
hsa_circ_004327	chr17	HuR_Human_GSE28865_PAR-CLIP	3579788	3579792
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hsa_circ_004327	chr17	U2AF65_Human_E-MTAB-1371_iCLIP	3579993	3579994
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8			6	1
hsa_circ_004327	chr17	eIF4AIII_Human_GSE40778_HITS-CLIP	3580067	3580079
8			8	6

Chrom/chr, chromosome; circRNA, circular RNA; RBP, RNA binding protein.