

gene	circBase RNA ID	RPF count	
		NB	GBM
<i>circ-HGF</i>	<i>hsa_circ_0080914</i>	0	11
<i>circ-E-Cad</i>	<i>hsa_circ_0039992</i>	0	8
<i>circ-Rictor</i>	<i>hsa_circ_0129052</i>	3	5
<i>circ-PINK1</i>	<i>hsa_circ_0111789</i>	1	7
<i>circ-TAZ</i>	<i>hsa_circ_0139927</i>	3	9
<i>circ-Rictor</i>	<i>hsa_circ_0129052</i>	3	7
<i>circ-PTPRZ1</i>	<i>hsa_circ_0133169</i>	6	11
<i>circ-RALY</i>	<i>hsa_circ_0114956</i>	12	4
<i>circ-KIT</i>	<i>hsa_circ_0126617</i>	5	6
<i>circ-CNST</i>	<i>hsa_circ_0112857</i>	13	2
<i>circ-PAX6</i>	<i>hsa_circ_0095689</i>	3	9
<i>circ-GLI3</i>	<i>hsa_circ_0134316</i>	4	7

**Supplementary Figure S1.** Twelve differentially expressed coding circRNAs which were annotated in circBase and numbers of ribosome protected fragments (RFPs) detected in GBM and normal brain (NB).