

A	BIRC1A1	TTAAACTAGTAAGGAGATCAACATTTTCAAATTAGATGTTTCAACTGTGCTCTAAGCTTTTAA
A	BIRC1A2	TTAAAAGCTTAGAGCACAGTTGAAACATCTAATTTGAAAATGTTGATCTCCTTACTAGTTTAA
A	BIRC1B1	TTAAACTAGTAAGGAGATCAACATTTTCAAATTATATGTGTCAACTGTGCTCTAAGCTTTTAA
A	BIRC1B2	TTAAAAGCTTAGAGCACAGTTGACACATATAATTTGAAAATGTTGATCTCCTTACTAGTTTAA
B	BIRC2A1	TTAAACTAGTGCAGTGTCCCTTTTGCTAGAGCTGACAGCTTTGTTTCGCGTGGAAGCTTTTAA
B	BIRC2A2	TTAAAAGCTTCCACGCGAACAAAGCTGTCAGCTCTAGCAAAAGGGACACTGCACTAGTTTAA
B	BIRC2B1	TTAAACTAGTGCAGTGTCCCTTTTGCTAGAGCTGACATCTGTGTTTCGCGTGGAAGCTTTTAA
B	BIRC2B2	TTAAAAGCTTCCACGCGAACACAGATGTCAGCTCTAGCAAAAGGGACACTGCACTAGTTTAA
C	320F	CCTCCCAAAGTGTGGGATTA
C	320R	TTCTAGGTGGGGGAGATACCA
C	494F	TCTTGACCACAGAATTCGAGA
C	494R	TTGTGGAAGTCCAGGCCA

*Key:*

A: Primers for creation of 3'-end survivin target (for mir-494). BIRC1A1/2 creates an exact target, and BIRC1B1/2 a mismatched target.

B: Primers for 3' survivin target (for mir-320a) BIRC2A1/2 create an exact target and BISC2B1/2 a mismatch target.

C: Primer sequences for chromatin immunoprecipitation-PCR analysis.