

**Table S3**

<b>gene</b>	<b>region (chr.: position bp)</b>	<b>forward</b>	<b>reverse</b>
<i>HLHm<math>\alpha</math>-m<math>\beta</math></i>	3R: 21,832,944-21,833,048	GCCGGCGACGATGACGCTCGC	TCGAGTAGCTGCCGTCTCGGAGCT
<i>HLHm5-m6</i>	3R: 21,856,326-21,856,404	ACCGCTCGAGACGCGTCGTGT	CGCTGGCTTTCCGTGTATCCGTGCC
<i>HLHm7</i>	3R: 21,862,621-21,862,714	TGCTCCGCAGGTGGTGGTTCTTCG	CGCCGCGAGAGCGTCTCGTGC
<i>cut</i>	X: 7,532,437-7,532,570	GCGCCACCACCCAAAGGTTCAACC	ACAACAGCGCGTGTCTAGCTGCCA
<i>Eip75B</i>	3L: 17,985,690-17,985,807	TGTGCCCAGGACCGCGATGTTGT	TGCCAGCAACAGGCCCTTGCCC
<i>BrC</i>	X: 1,515,021-1,515,108	CCGCGAGGGGCGGCAACAAGTT	TGTGCGTGTGTAGGTTGCCCGATCG