

	Sequence	GadFly Coordinates
<b>PCR frag in Fig. 4:</b>		
+7101F	CAGCTCGCGATCGGTCATGT	6979712-6979731
+7101R	CGGCGGAAATGGTGGGAGTG	6979916-6979897
+1218F	GCCAAAGAGGTATGGGTAGC	6985512-6985531
+1218R	GTGGTTATCCCCCATATGGC	6985743-6985724
+138F	CGCTTATCCAGGGTTGCCATACCA	6986594-6986617
+138R	CGGCACCCCTCTGTCGATCC	6986960-6986941
-101F	GTCCTGCGGGCCATCGATCT	6986915-6986934
-101R	CGGGGATCCGCGAAATGCAG	6987117-6987098
-432F	GCCGAAAGGTGGAACGTGGC	6987225-6987244
-432R	GTATTGTGCTGGTATTGCTTC	6987469-6987449
-918F	TTTTGCTAGGATTCGGGTAG	6987721-6987741
-918R	CTGTTCTTGGCGTTCGATTAG	6987944-6987924
-1132F	CTAATCGAACGCCAAGAACAG	6987924-6987944
-1132R	CTAGGGCTTAGGGCTCACTGG	6988169-6988149
-1394F	TTCCAGTGAGCCCTAAGCCCTAGCC	6988147-6988171
-1394R	TAGGCCGCCCCACCCACTTG	6988463-6988444
-1810F	GCTGGAATGGGGTGGTACGG	6988510-6988529
-1810R	CCCCAAAAGTTTTGTCCGTGATGGAC	6988940-6988915
-2224F	GTCCATCACGGACAAAACCTTTGGGG	6988915-6988940
-2224R	AGAACACACATTCACTCAAGCACTGC	6989362-6989337
-2647F	GATGCATGTACACATCTGTGC	6989424-6989444
-2647R	ATATGTTTCGGTTCGCAATG	6989719-6989699
-3488F	GAAGCGAACATCTTCCCTTGC	6990249-6990269
-3488R	CGACCGCCCTACGTCGACGGC	6990558-6990538
-5094F	ACACGGATTGCATGGGGCACA	6991909-6991929
-5094R	TGACCGACAACCGCGAAACGTG	6992109-6992088
-5331F	CACGTTTCGCGGTTGTCGGTCA	6992088-6992109
-5331R	TGAACGAAATCTAACGGGGCTCAAAAAGAC	6992404-6992376
<b>Sxl RT-PCR:</b>		
<i>SxlP1F</i>	CACCGCTGCCCAGCGACAAT	6979801-6979820
<i>SxlP2F</i>	TTGCGTTTCGTTGGCGAGGACC	6982643-6982662
<i>SxlP3R</i>	GAGCGCTGAGCGCCAAAACAATTGA	6985955-6985931
<b>Sxl in situs:</b>		
<i>SxlPeF</i>	GTTTCTAAGCAGATCCCG	6986495-6986512
<i>Sxl PeR</i>	GTTCCACTCGTGACAAGTCC	6986902-6986883
<i>SxlPmF</i>	GGGCGATGCTTGCATGTTGC	6991059-6991078
<i>SxlPmR</i>	GCGAAACGTGCACACTGC	6992097-6992080
<b>ChIP control primers:</b>		
<i>RpA70F</i>	CACCGCCGAGGCTCAAGGAC	4088834-4088853
<i>RpA70R</i>	TGCAAGGAGGTGGGCGAGTTG	4089056-4089036
<i>hbF</i>	GCCTGTCAATGCTGGCGACTTTTCG	4523493-4523516
<i>hbR</i>	GAAAGAGCGAACGCCACGAAGGG	4523867-4523845
<i>kl-2F</i>	GGTTGCCCAATGTGAGTGAAGCGTC	82313-82337
<i>kl-2R</i>	TCGCTTAGCTCGCCGCCGTA	82775-82756