

Table S1. Primer sequences

Primer	Sequence (5' to 3')
Sequencing	
ago 3' sense-seq	TGTGCGTCTGTGTGTATGCAG
ago 3 antisense-seq	CGCCGTAGGCAGAGATTGG
Gl1-sense-seq	ATGTATCTTATGGTACTGTA ACTG
Gl2-antisense-seq	CTTTATGTTTTTGGCGTCTTCCA
Mutagenesis	
Su(H) site 1-sense	TTGCGCCTGTATGTGAGACAGTGGGCCCAAAC
Su(H) site 1-antisense	GTTTGGGGCCCACTGTCTCACATACAGGCGCAA
Su(H) site 2-sense	GCGTATCGAAATGCGGAGTCAAAGCGGAGACCCTAA
Su(H) site 2-antisense	TTAGGGTCTCCGCTTTGACTCCGCATTTTCGATACGC
Su(H) site 3-sense	GAAGCAGTAGAAGGAGGGAGGCAAGAGAATGAATACGAATA
Su(H) site 3-antisense	TATTCGTATTCATTCTCTTGCCCTCCCTCCTTCTACTGCTTC
Su(H) site 4-sense	GGAAGAGAATGAATACGAATAAGGCAA AATGTTTCGGGAGAAATGCC
Su(H) site 4-antisense	GGCATTCTCCCGAACATTTTGCCTTATTCGTATTCATTCTCTTCC
Su(H) site 5-sense	GAGCAACGTCCTTAAAACGTAAGTCAAAAATTCGATTGAACGTGTG
Su(H) site 5-antisense	CACACGTTCAATCGAATTTTTGACTTACGTTTTAAGGACGTTGCTC
Su(H) site 6-sense	CATCCCTAACTTAAGCCGTGAATTGCCTTGGTATATTTGAACG
Su(H) site 6-antisense	CGTTCAAATATACCAAGGCAATTCACGGCTTAAGTTAGGGATG