

**TABLE S1: PCR primers used in this study**

**2S rRNA depletion:**

5'-AGTCTTACAACCCCTCAACCATATGTAGTCCAAGCAGCACT-3'

**qPCR (miScript assay):**

bantam-fwd	5'-GAGATCATTGAAAGCTGATT-3'	
miR-12-fwd	5'-GAGTATTACATCAGGTACTGGT-3'	
mir-13b-1-2-fwd	5'-TATCACAGCCATTGACGAGT-3'	
mir-13a-fwd	5'-TATCACAGCCATTGATGAGT-3'	
mir-283-fwd	5'-TAAATATCAGCTGGTAATTCT-3'	
mir-304-fwd	5'-AATCTCAATTGAAATGTGAG-3'	
mir-210-fwd	5'-TTGTGCGTGTGACAGCGGCTA-3'	
mir-iab-4-5p-fwd	5'-ACGTATACTGAATGTATCCTGA-3'	
mir-iab-4-3p-fwd	5'-GTATACCTTCAGTATACGTAAC-3'	
mir-318-fwd	5'-TCACTGGGCTTGTTATCTCA-3'	
mir-985-fwd	5'-CAAATGTTCCAATGGTCGGGCA-3'	
mir-979-fwd	5'-TTCTTCCCGAACTCAGGCTAA-3'	
mir-981-fwd	5'-TTCGTTGTCGACGAAACCTGCA-3'	
mir-977-fwd	5'-TGAGATATTCACGTTGTCTAA-3'	
mir-285-fwd	5'-TAGCACCATTCGAAATCAGTGC-3'	
mir-969-fwd	5'-GAGTCCACTAAGCAAGTTT-3'	
mir-972-fwd	5'-TGTACAATACGAATATTTAGGC-3'	
mir-978-fwd	5'-TGTCCAGTGCCGTAAATTGCAG-3'	
mir-314-fwd	5'-TATTCGAGCCAATAAGTCGG-3'	
mir-100-fwd	5'-AACCGTAAATCCGAACTTGTG-3'	
mir-137-Fwd	5'-TATTGCTTGAGAACATACGTAG-3'	
mir-989-fwd	5'-ATGTGATGTGACGTAGTGGAAC-3'	
mir-1013-fwd	5'-AATAAAAGTATGCCGAACTCG-3'	
mir-Spike-fwd	5'-CTCAGGATGGCGGAGCGGTG-3'	external reference
U6snRNA-fwd	5'-ATTGGAACGATAACAGAGAAAGATTAG-3'	reference
Dspt-4-fwd	5'-TTGACGCGATACCCAAGGAT-3'	reference
Dspt4-rev	5'-CTAGTGTGATCATAGACATTGTCCTTGT-3'	

**qPCR on sex-specific transcripts:**

RpL32-fwd	5'-AAGATGACCATCCGCCAGC-3'	endogenous control
RpL32-rev	5'-GTCGATACCCTGGGCTTGC-3'	
Esg-fwd	5'-CGCCCAGTGGATCTGAAATC-3'	
Esg rev	5'-GGTCTTGTCAACATCCTTG-3'	(Chau <i>et al.</i> 2009)
DsxM fwd	5'-TCCTTGGGAGCTGATGCCAC-3'	
DsxM rev	5'-GGCTACAGTGCATTTATT-3'	
Yp1-fwd	5'-TGAGCGTCTGGAGAACATGAA-3'	
Yp1-rev	5'-GCGACAGGTGGTAGACTTGCT-3'	
tra1-fwd	5'-GGAACCCAGCATCGAGATT-3'	
tra1-rev	5'-ATGCCCATGGTATTCTTTTC-3'	
Sxl-fwd	5'-ACAACGACAGCAGCAGGCCA-3'	
Sxl-rev	5'-TTGTAACCACGACGCGACGAT-3'	(Hashiyama <i>et al.</i> 2011)