



30 nt



## Supplementary Table of high throughput datasets analyzed in this study

Sample name	title	used in the following figure panels	GEO Accession
libraries sequenced in this study			
w1118_total_no_2S_rRNA_depletion	small RNAs (18-29nt) from w1118 ovaries without 2S rRNA depletion	Fig. ED4	GSE83698
w1118_total	small RNAs (18-40nt) from w1118 ovaries with 2S rRNA depletion	Fig. 2a, 2b, 3c, ED6c, ED4d, ED4e, ED4, ED4i, ED7b	GSE83698
w1118_PiwiIP	Piwi bound small RNAs (18-40nt) from w1118 ovaries	Fig. 1e, 2c, 3b, 3c, ED6c, ED3j, ED5a, ED5b, ED6a	GSE83698
w1118_AubIP	Aub bound small RNAs (18-40nt) from w1118 ovaries	Fig. 1e, 2c, 2d, 2e, 3b, 3c, ED6c, ED5a, ED5b, ED5c, ED6a	GSE83698
w1118_Ago3IP	Ago3 bound small RNAs (18-40nt) from w1118 ovaries	Fig. 1e, 2c, 2d, 2e, 3b, 3c, ED6c, ED5a, ED5b, ED5c, ED6a	GSE83698
Nibbler_KO_total	small RNAs (18-40nt) from Nibbler knock-out ovaries with 2S rRNA depletion	Fig. 2a, 2b, 3c, ED6c, ED4d, ED4i, ED4e, ED7b	GSE83698
Nibbler_KO_PiwiIP	Piwi bound small RNAs (18-40nt) from Nibbler -/- ovaries	Fig. 1e, 2c, 3b, 3c, ED6c, ED3j, ED5a, ED5b, ED6a	GSE83698
Nibbler_KO_AubIP	Aub bound small RNAs (18-40nt) from Nibbler -/- ovaries	Fig. 1e, 2c, 2d, 2e, 3b, 3c, ED6c, ED5a, ED5b, ED5c, ED6a	GSE83698
Nibbler_KO_Ago3IP	Ago3 bound small RNAs (18-40nt) from Nibbler -/- ovaries	Fig. 1e, 2c, 2d, 2e, 3b, 3c, ED6c, ED5a, ED5b, ED5c, ED6a	GSE83698
Papi_KO_PiwiIP	Piwi bound small RNAs (18-40nt) from Papi -/- ovaries	Fig. 1e, ED3j	GSE83698
Papi_KO_AubIP	Aub bound small RNAs (18-40nt) from Papi -/- ovaries	Fig. 1e	GSE83698
Papi_KO_Ago3IP	Ago3 bound small RNAs (18-40nt) from Papi -/- ovaries	Fig. 1e	GSE83698
Zuc_depletion_AubIP	Aub bound small RNAs (18-40nt) from Zucchini depleted ovaries	Fig. 1b	GSE83698
Zuc_Nbr_depletion_AubIP	Aub bound small RNAs (18-50nt) from Zucchini and Nibbler depleted ovaries	Fig. ED8b, ED9b	GSE83698
Zuc_Nbr_depletion_Ago3IP	Ago3 bound small RNAs (18-50nt) from Zucchini and Nibbler depleted ovaries	Fig. ED8b, ED9b	GSE83698
Biogenisis_repoter_1_two_sites+Zuc_depletion	small RNAs (18-40nt) from flies containing piRNA biogenesis repoter_1 with two target sites with Zucchini depletion and 2S rRNA depletion	Fig. 1d, ED7a, ED8b	GSE83698
Biogenisis_repoter_1_two_sites+Zuc_Papi_depletion	small RNAs (18-40nt) from flies containing piRNA biogenesis repoter_1 with two target sites with Zucchini and Papi depletion and 25 rRNA depletion	Fig. 1d	GSE83698
Biogenisis_repoter_1_two_sites+Zuc_Nbr_depletion	small RNAs (18-40nt) from flies containing piRNA biogenesis repoter_1 with two target sites with Zucchini and Nibbler depletion and 2S rRNA depletion	Fig. 1d, ED7a, ED8b	GSE83698
Biogenisis_repoter_2_two_sites+Zuc_depletion	small RNAs (18-40nt) from flies containing piRNA biogenesis repoter_2 with two target sites with Zucchini depletion and 25 rRNA depletion	Fig. ED7a, ED2f, ED8b	GSE83698
Biogenisis_repoter_2_two_sites+Zuc_Papi_depletion	small RNAs (18-40nt) from flies containing piRNA biogenesis repoter_2 with two target sites with Zucchini and Papi depletion and 2S rRNA depletion	Fig. ED2f, ED3d, ED2g	GSE83698
Biogenisis_repoter_2_two_sites+Zuc_Nbr_depletion	small RNAs (18-40nt) from flies containing piRNA biogenesis repoter_2 with two target sites with Zucchini and Nibbler depletion and 2S rRNA depletion	Fig. ED7a, 4b, ED2f, ED8b	GSE83698
Biogenisis_repoter_2_three_sites+Zuc_Nbr_depletion	small RNAs (18-40nt) from flies containing piRNA biogenesis repoter_2 with three target sites with Zucchini and Nibbler depletion and 2S rRNA depletion	Fig. ED7a, 4b, ED8b	GSE83698
Zuc_Nbr_depletion_rep1	small RNAs (18-40nt) from Zucchini and Nibbler depleted ovaries with 25 rRNA depletion	Fig. ED7a, 4a, 4c, 4d, ED8a, ED8b, ED8c, ED9a, ED10a	GSE83698
Zuc_Nbr_depletion_rep2	small RNAs (18-40nt) from Zucchini and Nibbler depleted ovaries with 2S rRNA depletion	Fig. ED7a, ED8b	GSE83698
Zuc_Nbr_depletion_rep3	small RNAs (18-40nt) from Zucchini and Nibbler depleted ovaries with 2S rRNA depletion	Fig. ED7a, ED8b	GSE83698
Zuc_depletion_rep1	small RNAs (18-40nt) from Zucchini and depleted ovaries with 25 rRNA depletion	Fig. 1b, 2a, 2b, ED7a, 4a, 4c, 4e, ED1a, ED1b, ED2g ED3d, ED8a, ED8b, ED8c, ED9	a GSE83698
Zuc_depletion_rep2	small RNAs (18-40nt) from Zucchini and depleted ovaries with 25 rRNA depletion	Fig. ED7a, ED8b	GSE83698
Zuc_depletion_rep3	small RNAs (18-40nt) from Zucchini and depleted ovaries with 25 rRNA depletion	Fig. ED7a, ED8b	GSE83698
Zuc_depletion_rep4	small RNAs (18-40nt) from Zucchini and depleted ovaries with 25 rRNA depletion	Fig. ED7a, ED8b	GSE83698
Zuc_Nbr_depletion_oxidized	small RNAs (18-40nt) from Zucchini and Nibbler depleted ovaries with 2S rRNA depletion and oxidation	Fig. ED10b	GSE83698
w1118_polyA_plus_RNAseq	polyA selected RNAs from w1118 ovaries	Fig. ED3g	GSE83698
nibbler_null_polyA_plus_RNAseq	polyA selected RNAs from Nibbler -/- ovaries	Fig. ED3g	GSE83698
control_depletion_polyA_plus_RNAseq	polyA selected RNAs from control depletion ovaries	Fig. ED7d	GSE83698
zucchini_depletion_polyA_plus_RNAseq	polyA selected RNAs from Zucchini depleted ovaries	Fig. ED7d	GSE83698
zucchini_and_nibbler_depletion_polyA_plus_RNAseq	polyA selected RNAs from Zucchini and Nibbler depleted ovaries	Fig. ED7d	GSE83698
libraries sequenced previously			
Wsh_total	small RNAs (18-29nt) from control depletion ovaries	Fig. ED6d, ED1a, ED1b	GSE55824

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Wsh_PiwiIP	Piwi bound small RNAs (18-40nt) from control depleted ovaries	Fig. ED1a	GSE64802
Wsh_AubIP	Aub bound small RNAs (18-40nt) from control depleted ovaries	Fig. ED1a	GSE64802
Wsh_Ago3IP	Ago3 bound small RNAs (18-40nt) from control depleted ovaries	Fig. ED1a	GSE64802
ZUCsh_PiwiIP	Piwi bound small RNAs (18-40nt) from Zucchini depleted ovaries	Fig. ED1a, ED5a	GSE64802
ZUCsh_Aub3IP	Aub bound small RNAs (18-40nt) from Zucchini depleted ovaries	Fig. 2c, 2d, 2e, ED8b, ED1a, ED5a, ED5b, ED5c, ED9b	GSE64802
ZUCsh_Ago3IP	Ago3 bound small RNAs (18-40nt) from Zucchini depleted ovaries	Fig. 2c, 2d, 2e, ED8b, ED1a, ED5a, ED5b, ED5c, ED9b	GSE64802