

Table S3. Comparison of proTRAC and piRNApredictor outputs.

Samples	Input	Common Output	Distribution of common piRNAs											
			Number of unique reads	Number of unique piRNAs	Repeat	Unannotated	Intron		Exon		snRNA	rRNA	miRNA	tRNA
							sense	antisense	sense	antisense				
PGCs	687544	35538	28520	5845	633	259	213	46	20	1	1	0		
Stage X	350113	4097	3695	301	65	31	4	0	0	1	0	0		
GSCs	350647	1795	1162	562	38	23	8	0	1	0	1	0		
CEFs	159803	27	1	19	3	0	4	0	0	0	0	0		