Supplementary figure 3

Dataset A
4189 sequenes

2973

1105

69
42
102

Dataset C
222 sequenes

2324

9

Dataset B
3480 sequenes

Supplementary figure 3. **Sequence overlap of abundant piRNAs**. Venn diagram of overlap between the datasets using piRNA sequences that were cloned three or more times. Sequences similarity was determined by a 95% identity measure. The degree of overlap is similar to the general piRNA population (Fig. 1) suggesting that the abundant piRNAs from different datasets do not originate from the same genomic locations.