

## Annotating piRNA processing

It should be possible to distinguish between primary and secondary piRNA processing in Drosophila by using the indicators below:

Feature	Primary	Secondary
<b>Cell type:</b> Follicle cell		
<b>Cell type:</b> Germ		
<b>Subcellular:</b> Nuage/P granule		
<b>Subcellular:</b> Yb body		
<b>Piwi protein:</b> piwi		
<b>Piwi protein:</b> AGO3		
<b>Piwi protein:</b> Aub		
<b>Mechanism of gene silencing:</b> heterochromatin formation		
<b>Mechanism of gene silencing:</b> post-transcriptional		
<b>Gene silencing occurs in:</b> nucleus		
<b>Gene silencing occurs in:</b> cytoplasm		
<b>Retrotransposon target:</b> uni-strand		
<b>Retrotransposon target:</b> dual-strand		
<b>piRNA population:</b> <a href="#">ping-pong pairs</a>		
<b>piRNA population:</b> <a href="#">antisense-bias</a>		

Excellent description of difference between primary and secondary piRNA processing in D.mel is given in PMID:[22902557](#) and PMID: [19395010](#).

Annotate the terms **GO:0140990 primary piRNA processing** or **GO:0140965 secondary piRNA processing** over the more general term **GO:0034587 piRNA processing**.