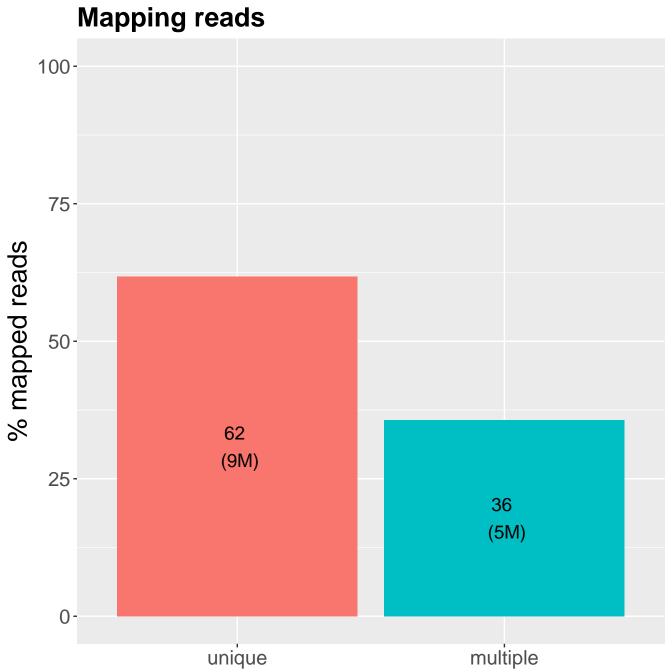


## Length filtering 100-75-% discarded reads 50-25-0-(242K) (64K) reads length < 15nt reads length > 32nt



## **Read distribution** 100-75-% reads 99 50-(9M) 25-0-(60K) (36) (2K)(131)piRNA miRNA snoRNA rRNA snRNA