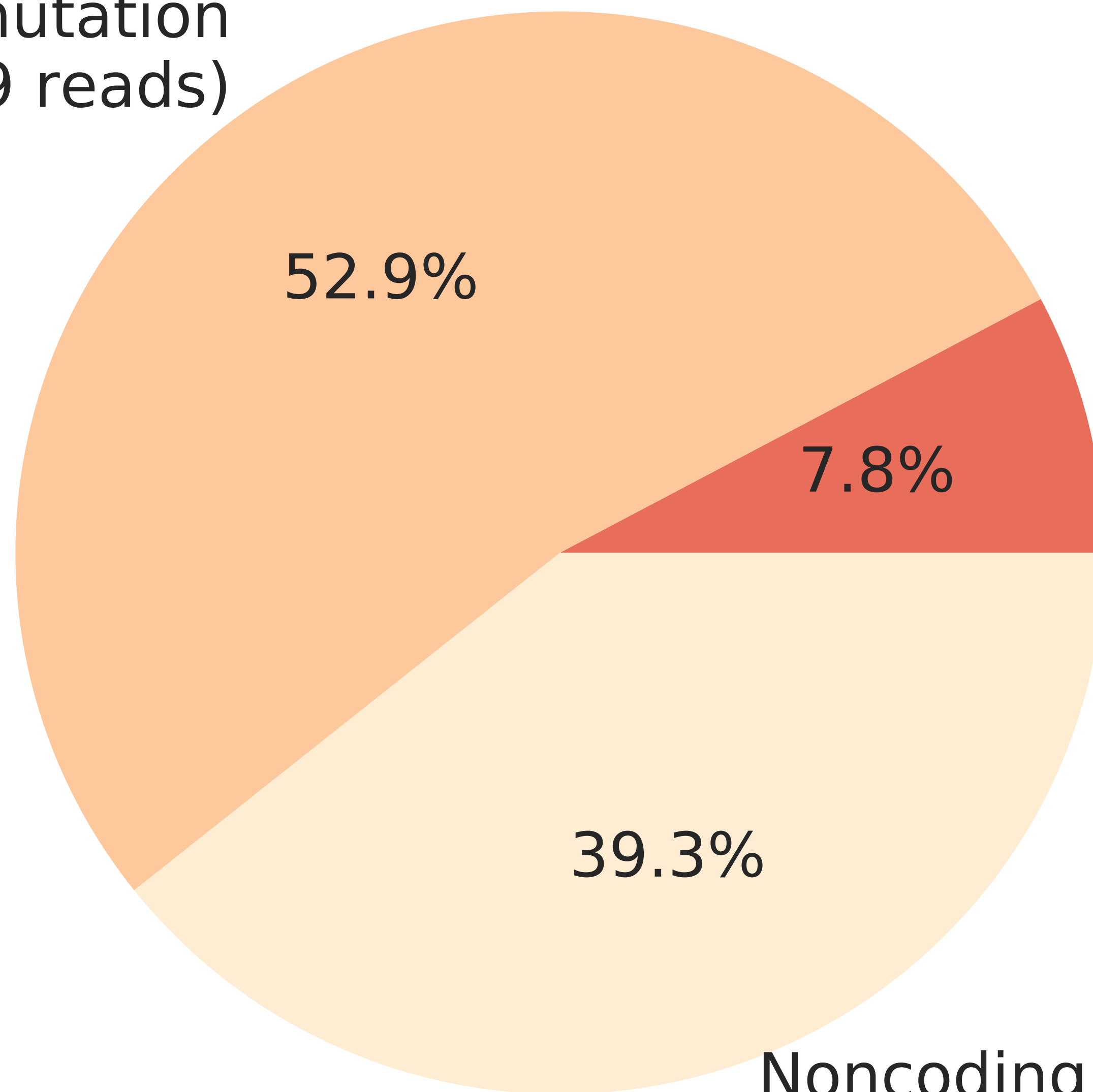


In-frame mutation  
(22099 reads)



Frameshift mutation  
(3240 reads)

Noncoding mutation  
(16397 reads)



- Amplicon sequence
- Coding sequence/s
- ▼ Predicted Cas9 cleavage site/s