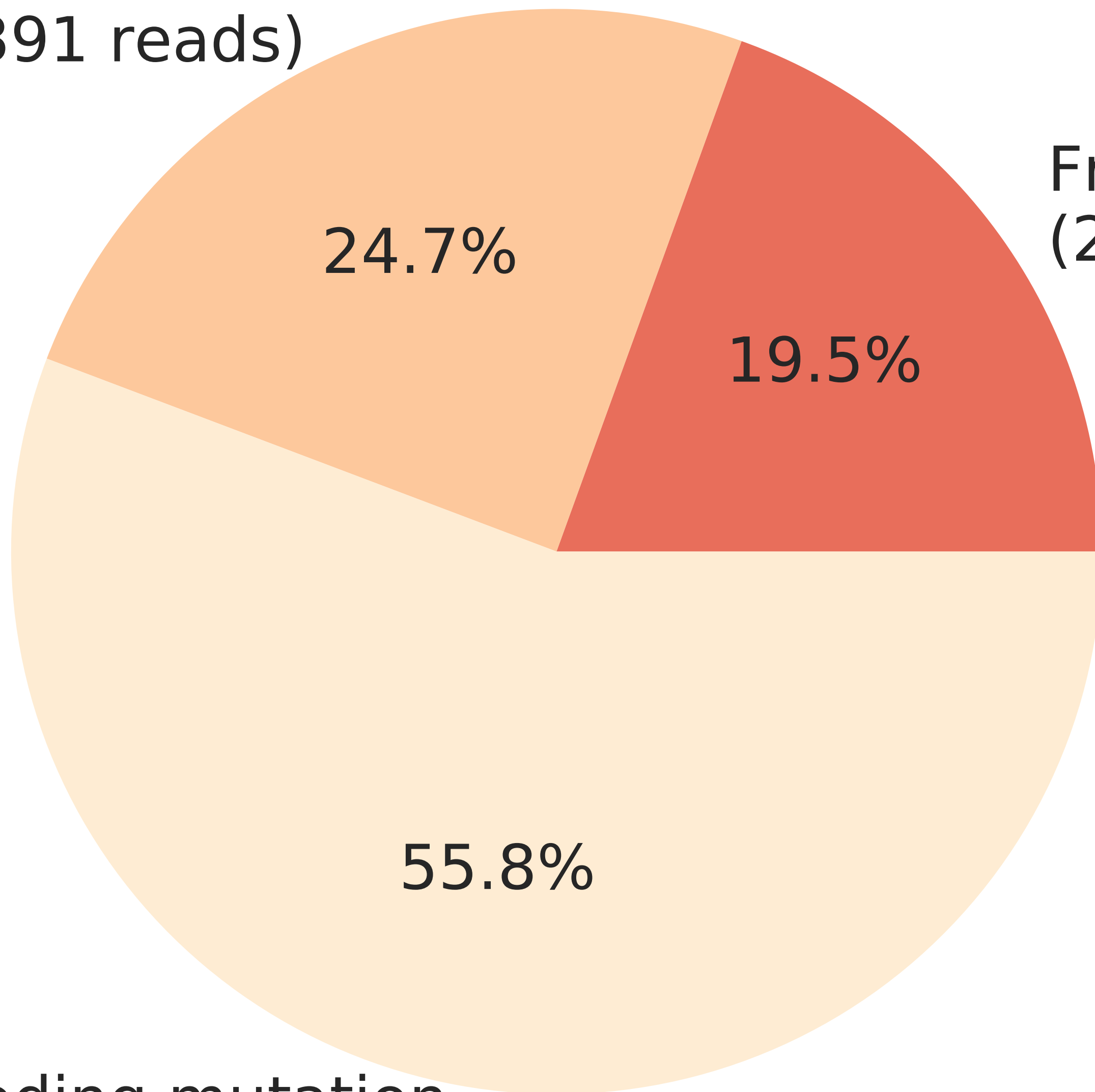In-frame mutation
(36391 reads)

Frameshift mutation
(28686 reads)

24.7%

19.5%

55.8%

Noncoding mutation
(82108 reads)

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