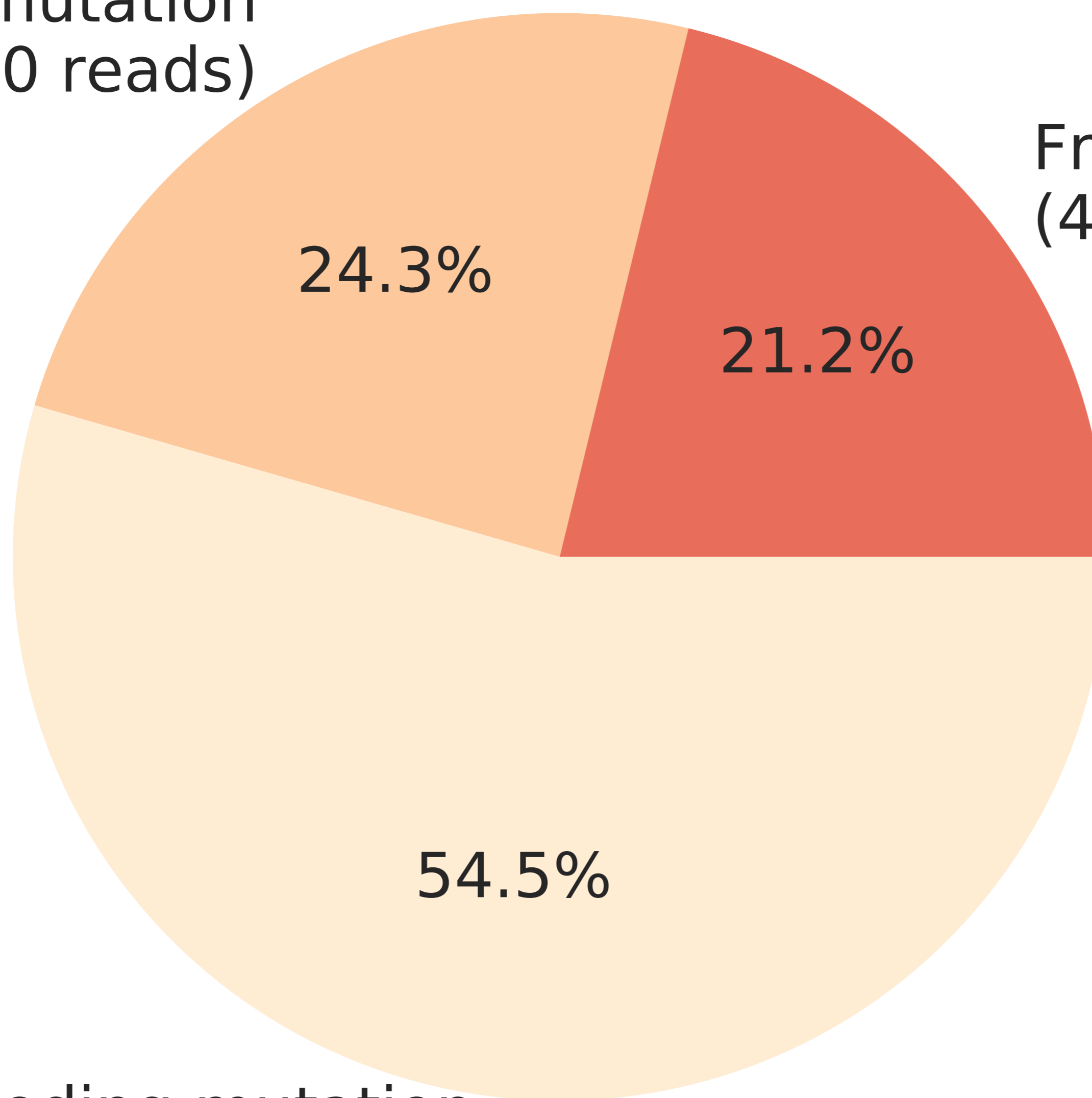In-frame mutation
(47190 reads)

Frameshift mutation
(41217 reads)

24.3%

21.2%

54.5%

Noncoding mutation
(105879 reads)

— Amplicon sequence

— Coding sequence/s

▼ Predicted Cas9 cleavage site/s