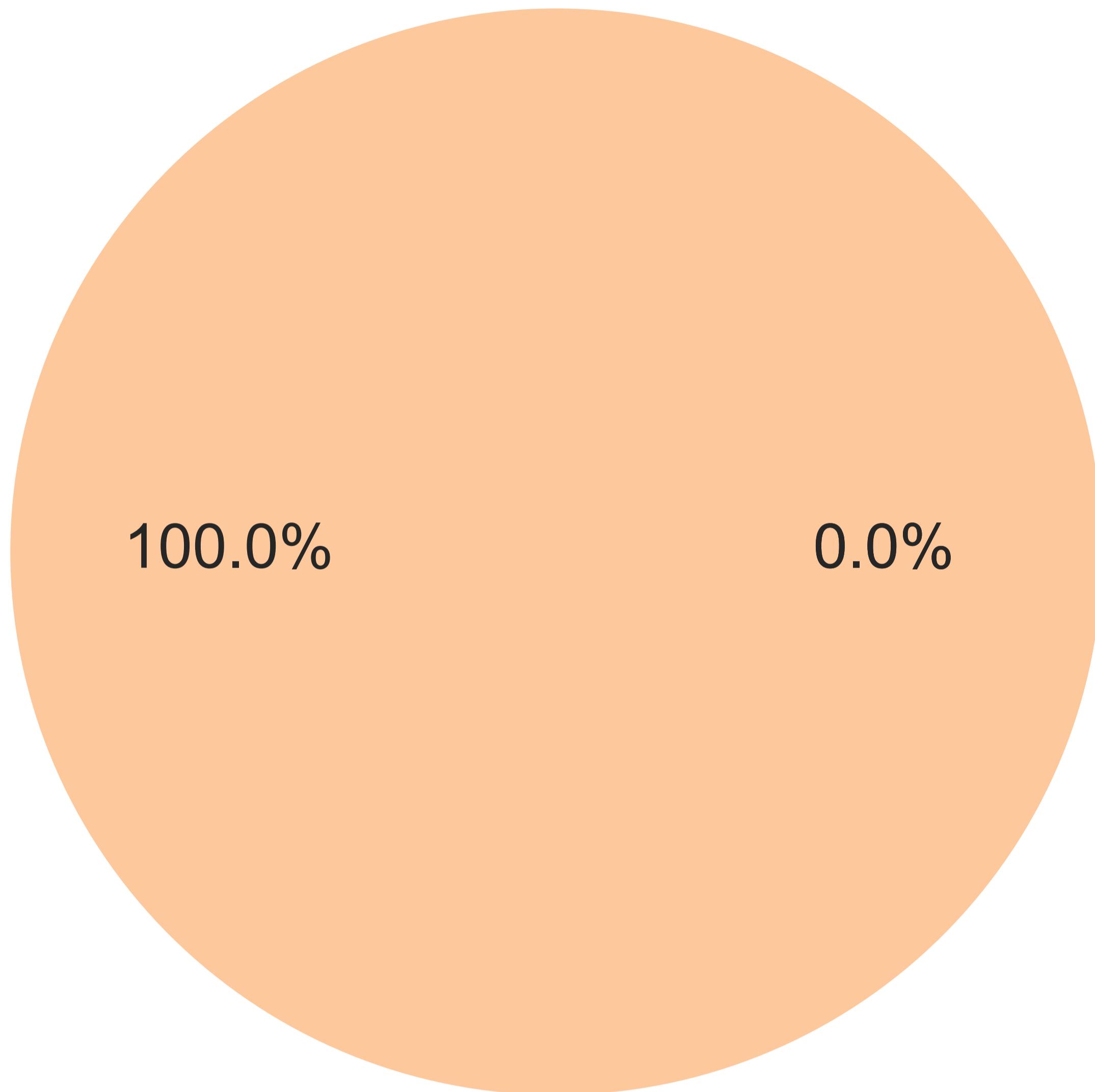


In-frame mutation
(1 reads)

Non-inframe mutation
(0 reads)



- Reference sequence
- Coding sequence/s
- - - Predicted cleavage position
- sgRNA