Mutation position distribution

- Reference amplicon position (bp)

Sequences: % Total (no.)
- 3.9% (921)
- 3.2% (737)
- 2.4% (552)
- 1.6% (368)
- 0.8% (184)
- 0.0% (0)

Quantification window

- Insertions
- Deletions
- Substitutions
- sgRNA
- Quantification window