

CORRECTION

# Correction: CCTop: An Intuitive, Flexible and Reliable CRISPR/Cas9 Target Prediction Tool

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The first sentence in the Material and Methods section under the subheading "Off-target search" is incorrect. The correct sentence is: For each off-target site of any sgRNA a score is computed that indicates the likelihood of an unstable sgRNA/DNA heteroduplex.

## Reference

1. Stemmer M, Thumberger T, del Sol Keyer M, Wittbrodt J, Mateo JL (2015) CCTop: An Intuitive, Flexible and Reliable CRISPR/Cas9 Target Prediction Tool. PLoS ONE 10(4): e0124633. <https://doi.org/10.1371/journal.pone.0124633> PMID: 25909470



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