

**E-CRISP**

Design of CRISPR constructs

De-novo

Evaluation

Help

Links

[Download a tabular report for all query sequences together](#)
**Query name: hG19\_dnA Query length: 201 Query location: ::0::0**

Total number of possible designs = 30

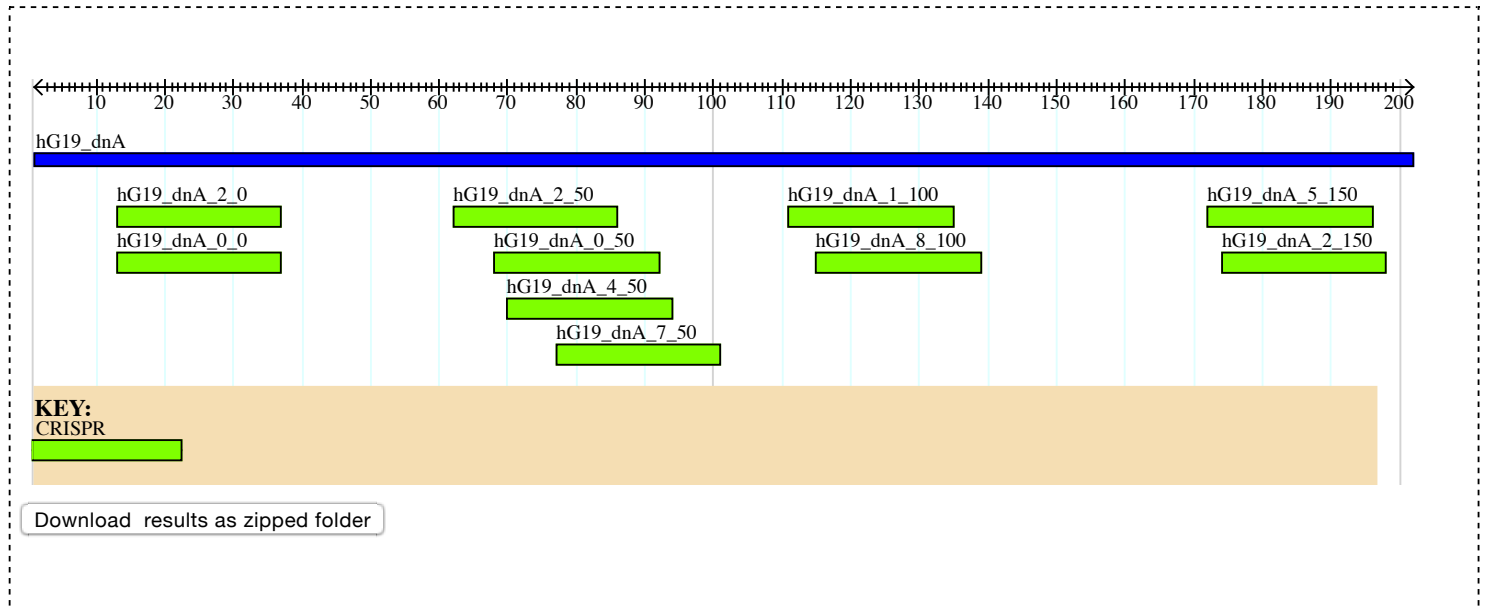
Number of successful designs = 10

Number of designs that hit a specific target = 30

Number of designs excluded because the maximum of designs per exon was exceeded = 20

S: Specificity score A: Annotation score E: Efficiency score  
for more information please see the [Help](#) pages

Name	Nucleotide sequence	SAE-Score	Target	Matchstring	Number of Hits
hG19_dnA_4_50	GGCCGGGGTTGCCA GGAAGT NGG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_2_0	CTGATCATCAAAAGT GAACG NGG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_1_100	AAGTCTGAAACTGTA GGGAA NGG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_0_0	CCACGTTCACTTTTG ATGAT NAG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_7_50	TATGCATGGCCGGGG TTGCC NGG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_5_150	ACCTTGGGTTTCAGTT GTTTC NGG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_2_150	AGAAACAACCTGAACC CAAGG NAG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_2_50	TTGCCAGGAAGTGG GAGCTG NAG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_0_50	CTCCCACTTCCTGGC AACCC NGG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1
hG19_dnA_8_100	ACCACCTTTCCTAC AGTTT NAG	S A E	ENSG00000182533::C AV3	<a href="#">Matchstring Info</a>	1



[Like](#) [Share](#) 105 people like this. [Sign Up](#) to see what your friends like.

### Boutros lab, E-CRISP-Version 4.2

For suggestions please contact us at [crispr@dkfz.de](mailto:crispr@dkfz.de)