

**Supplementary table 1.** Comparison of the SSFinder with other available tools.

<b>Name of the tool</b>	<b>Link</b>	<b>Input sequence/genome limit</b>	<b>Excess</b>	<b>Adjustability</b>
CRISPR Design tool	<a href="http://www.broadinstitute.org/mpg/crispr_design/">http://www.broadinstitute.org/mpg/crispr_design/</a>	1000 bp	Internet dependent	Non-editable
CRISPR Design	<a href="http://crispr.mit.edu/">http://crispr.mit.edu/</a>	Limited to 15 genome (only 1 from plants )	Internet dependent	Non-editable
CRISPR Target	<a href="http://bioanalysis.otago.ac.nz/CRISPRTarget/crispr_analysis.html">http://bioanalysis.otago.ac.nz/CRISPRTarget/crispr_analysis.html</a>	Limited to the 50 Mb user database at web interface	Internet dependent	Flexible
ZiFiT Targeter	<a href="http://zifit.partners.org/ZiFiT/ChoiceMenu.aspx">http://zifit.partners.org/ZiFiT/ChoiceMenu.aspx</a>	Limited to small number of sequences	Internet dependent	Non-editable
SSFinder	<a href="http://code.google.com/p/ssfinder">http://code.google.com/p/ssfinder</a>	Useful for large genomes	Internet independent	Editable