

## Evaluating the Efficiency of gRNAs in CRISPR/Cas9 Mediated Genome Editing in Poplars

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### Online Resource 2

#### Sequencing results of *NFP-like1* at first time point

Modification in *NFP-like1* gRNA5 at first time point. Highlighted in yellow: PAM-recognition motif.

Letter in red represent insertions according to wildtype. Gaps in red highlighted represent deletions according to wildtype. Sequences illustrate both alleles.

Plant line	Sequencing results	Commentary
Wildtype	...GGAAATA-ATG <b>GGG</b> ... ...GGAAATA-ATG <b>GGG</b> ...	
N 481-2	...GGAAATA--TG <b>GGG</b> ... ...GGAAATA--TG <b>GGG</b> ...	Homozygous deletion of A
N 481-9	...GGAA----- <b>GGG</b> ... ...GGAAATA--TG <b>GGG</b> ...	Biallelic mutation: deletion ATAAT and deletion A
N 481-15	...GGAAATA--TG <b>GGG</b> ... ...GGAAATA <b>A</b> ATG <b>GGG</b> ...	Biallelic mutation: deletion A, insertion A
N 481-23	...GGAAA--ATG <b>GGG</b> ... ...GGAAATA--TG <b>GGG</b> ... ...GGAAA--TG <b>GGG</b> ...	Chimera: deletion TA, deletion A, deletion TAA
N 481-33	...GGAAATA <b>A</b> ATG <b>GGG</b> ... ...GGAAATA--TG <b>GGG</b> ...	Biallelic mutation: insertion A, deletion A
N 481-97	...GGAAATA( <b>36nt</b> )ATG <b>GGG</b> ... ...GGAAATA( <b>42nt</b> )ATG <b>GGG</b> ...	Biallelic mutation: insertion 36 nt, insertion 42 nt