

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 GGG GGAAATA-ATG GGG . . .	
N 481-2	. . . GGAAATA- - TG GGG GGAAATA- - TG GGG . . .	Homozygous deletion of A
N 481-9	. . . GGAA- - - - - GGG GGAAATA- - TG GGG . . .	Biallelic mutation: deletion ATAAT and deletion A
N 481-15	. . . GGAAATA- - TG GGG GGAAATA A ATG GGG . . .	Biallelic mutation: deletion A, insertion A
N 481-23	. . . GGAAA- - - ATG GGG GGAAATA- - TG GGG GGAAA- - - TG GGG . . .	Chimera: deletion TA, deletion A, deletion TAA
N 481-33	. . . GGAAATA A ATG GGG GGAAATA- - TG GGG . . .	Biallelic mutation: insertion A, deletion A
N 481-97	. . . GGAATA(36nt)ATG GGG GGAATA(42nt)ATG GGG . . .	Biallelic mutation: insertion 36 nt, insertion 42 nt