Figure S1 – CRISPR-induced mutation in *FIO1*.
(A) Phenotype of *fio1-3* compared to the Col-0 wildtype when grown in LD conditions.
(B) Determination of flowering by counting the number of rosette leaves (RLN = rosette leaf number) at the bolting stage in LD. Plotted are average leaf number +/- SD, ***p=<0.001, N=10-14.
(C) Nucleotide alignment showing the CRISPR-induced genomic deletion found in *fio1-5*. Gene model on top shows the relative positions of all three *fio1* mutations.