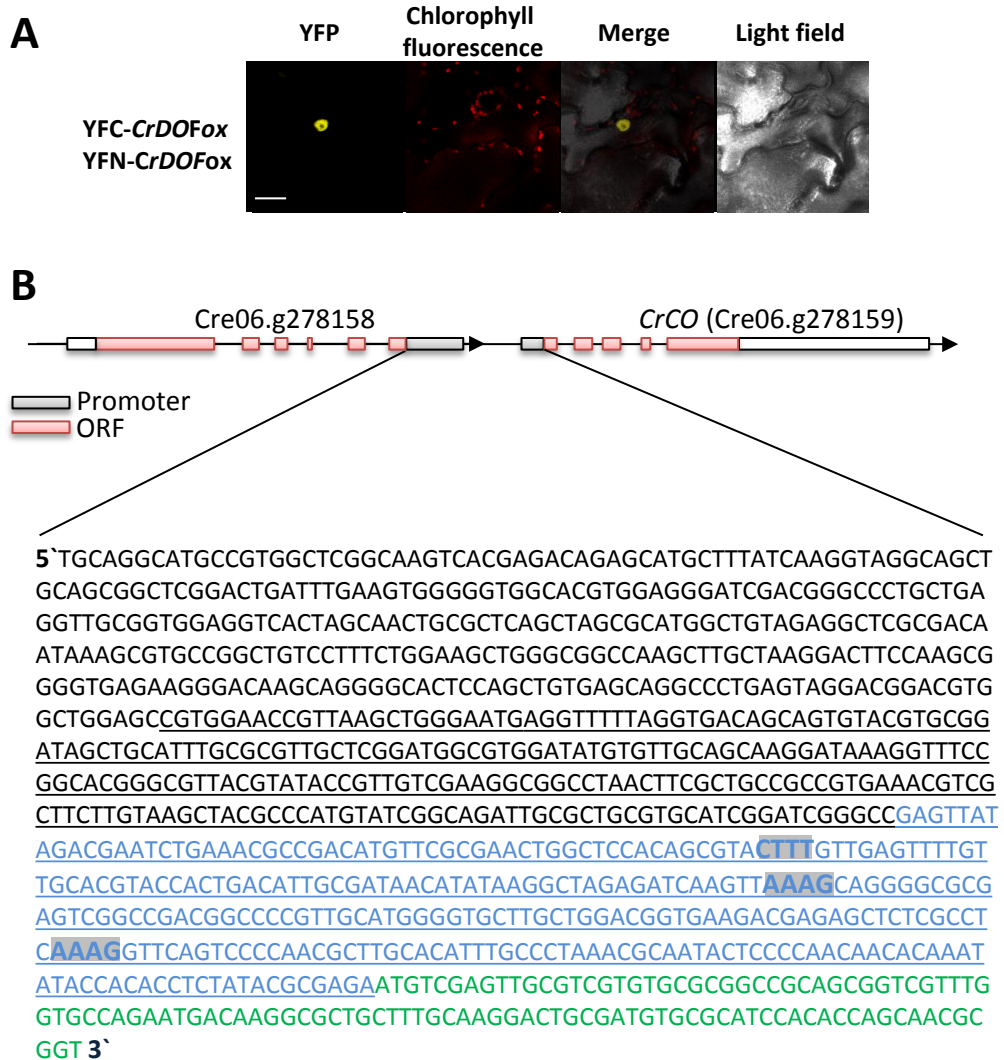


Supplemental Figure S3



Binding site mutations:

1. GGCTCCACAGCGTAC**CaaT**GTTGAGTTTTGTTGC (pCrCO3:GFP*).
2. CTAGAGATCAAGTT**AttG**CAGGGGCGCGAG (pCrCO3:GFP**).
3. GAGCTCTCGCCTC**AttG**GTTTCAGTCCCCA (pCrCO3:GFP***).