

**N**\* Novosibirsk  
State  
University  
\*THE REAL SCIENCE



# Regulation of Transcription Activity of *MAKR4* in *Arabidopsis thaliana* L.

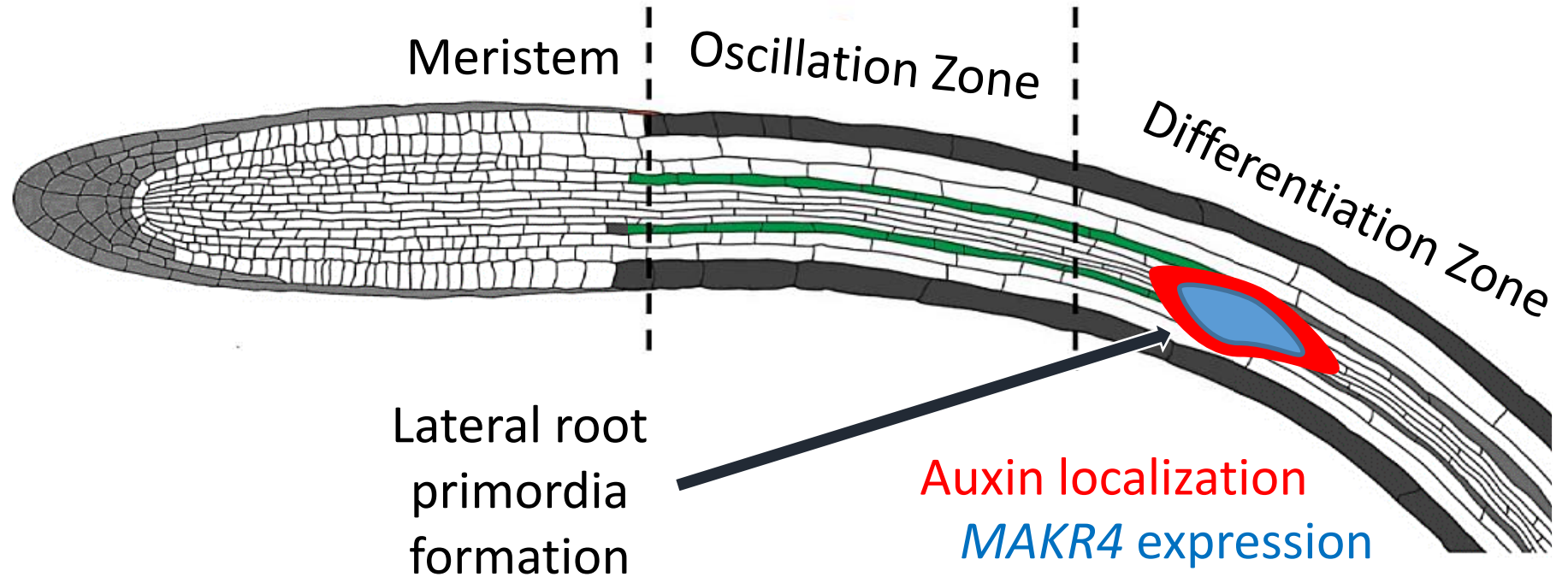
Anastasia Korosteleva  
Novosibirsk State University  
Novosibirsk, Russia  
kartzeva.kar@gmail.com

Daria Novikova  
Institute of Cytology and Genetics SB RAS  
Novosibirsk, Russia  
da6ik777@gmail.com

Victoria Mironova  
Institute of Cytology and Genetics SB RAS  
Novosibirsk, Russia  
kwiki@bionet.nsc.ru

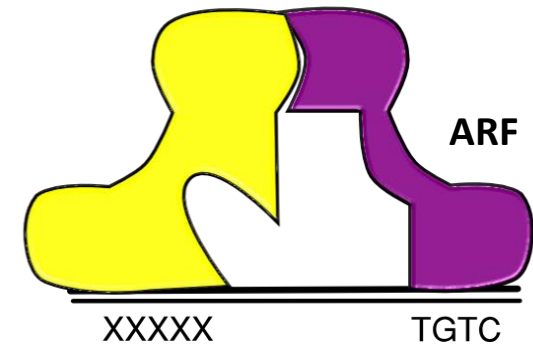
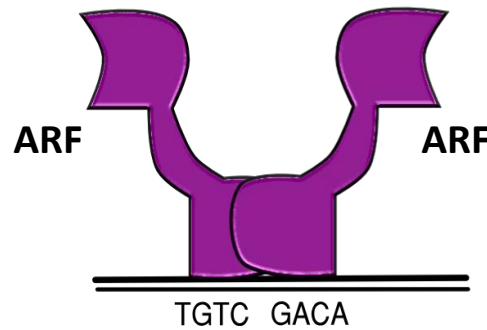
# MEMBRANE-ASSOCIATED KINASE REGULATOR4 is involved in the gene regulatory network of lateral root initiation in *Arabidopsis thaliana*

MAKR4 acts downstream of auxin oscillations to transform prebranch sites into lateral roots



Composite cis-regulatory element involved in the transcriptional response to auxin was detected in the promoter of the *MAKR4* gene with the consensus-based bioinformatics package *metaRE*

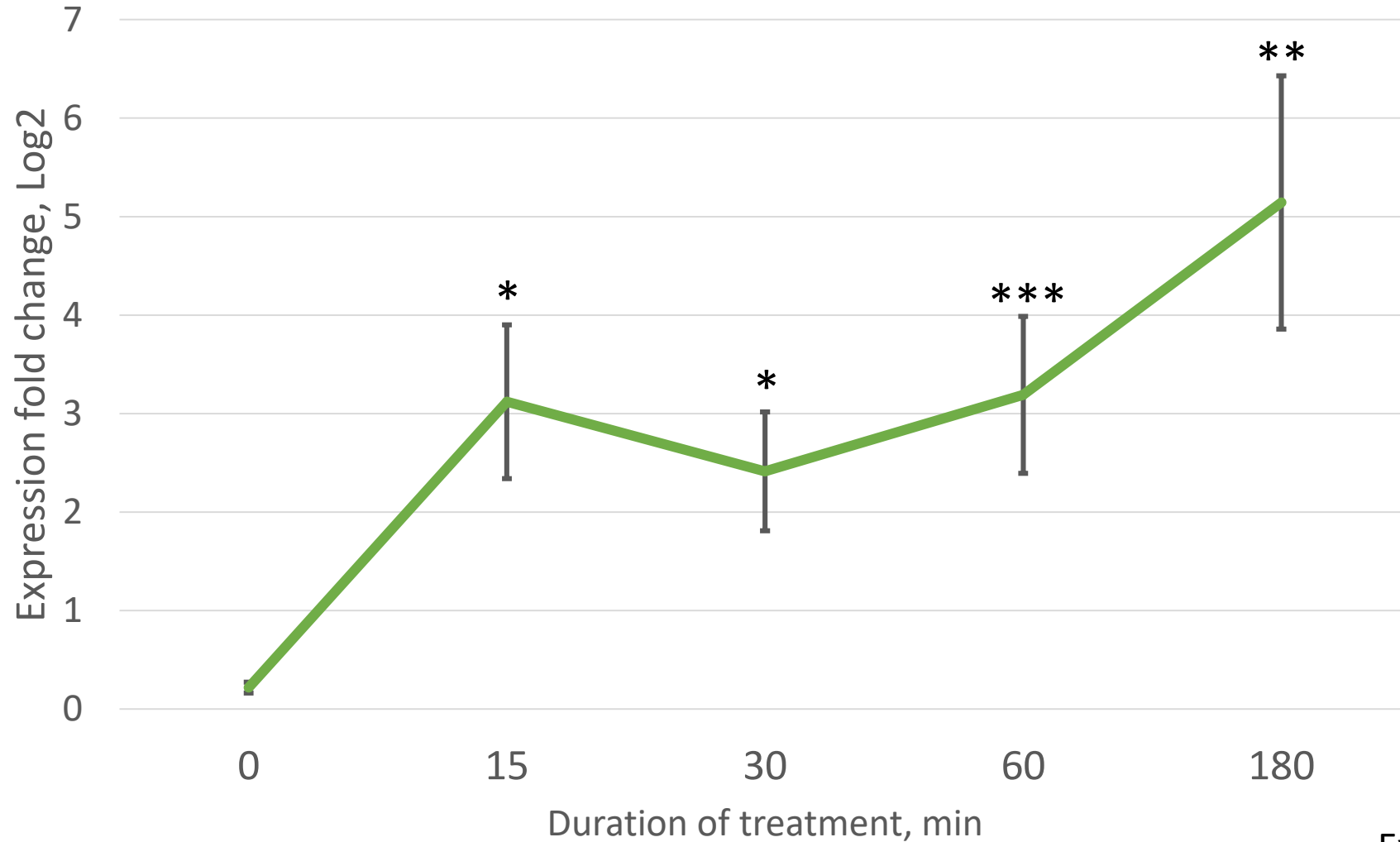
Auxin response factors (ARFs) bind to Auxin responsive elements (AuxRE) in upstream regions of auxin-sensitive genes as homo- or heterodimers



Predicted TF binding sites in the promoter of the *MAKR4* gene



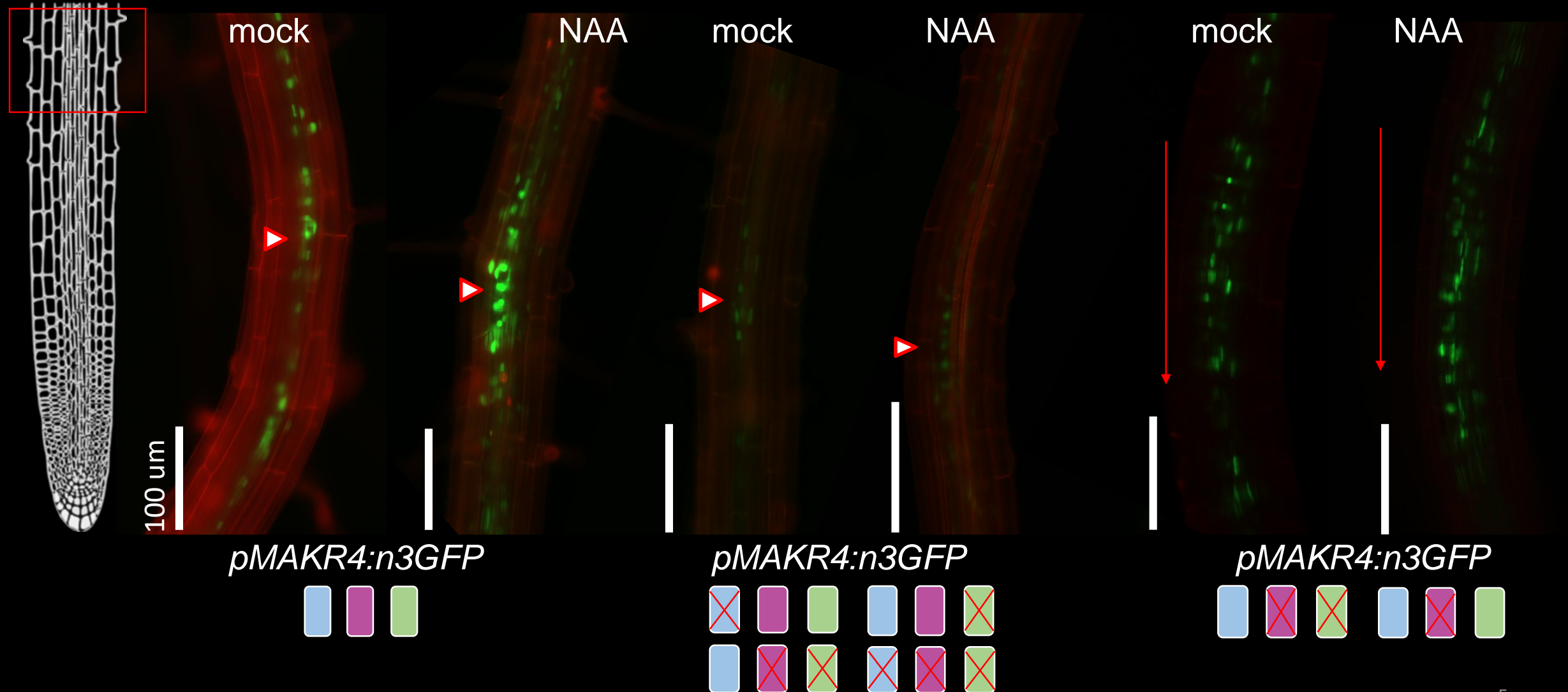
# Expression of the *MAKR4* in the *A.thaliana* root changes with different duration of auxin treatment



Student's *t*-test, significance level: \* – *p*-value < 0,05;  
\*\* – *p*-value < 0,005; \*\*\* – *p*-value < 0,001

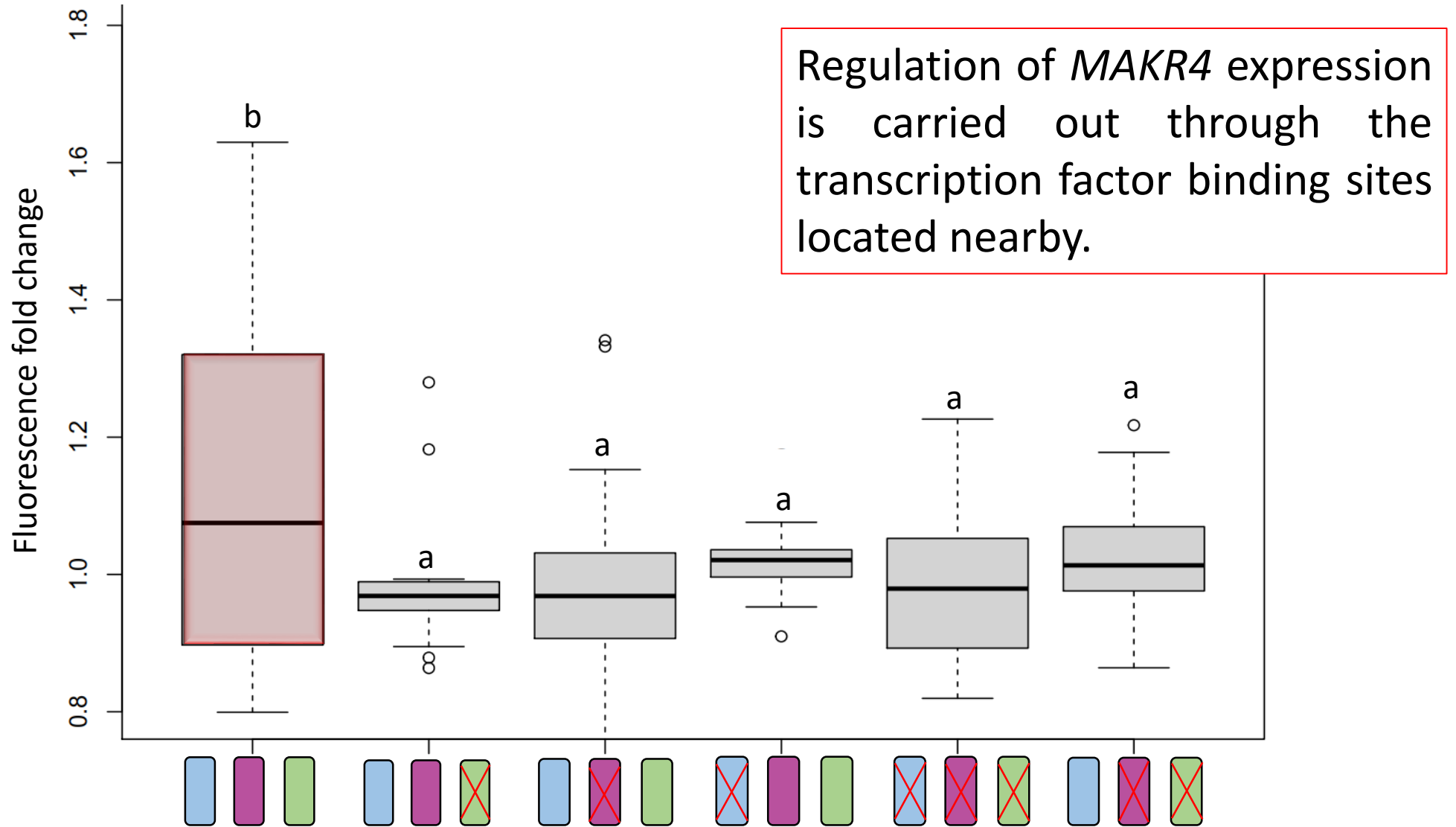
Experimental conditions:  
NAA treatment  
concentration is 1 μM.

# GFP expression under different variants of the *MAKR4* promoters varies



Cell walls stained with propidium iodide (PI)

# Functionality of the predicted sites of the composite cis-regulatory element confirmed quantitatively



ANOVA  
a differs from b,  $p < 0,05$