

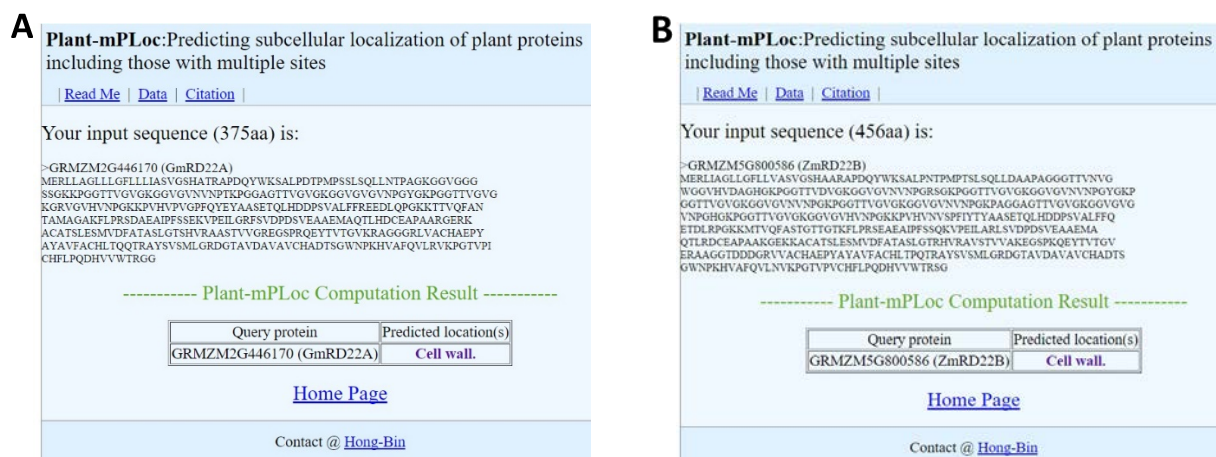
**Title:** Drought and exogenous abscisic acid alter hydrogen peroxide accumulation and differentially regulate the expression of two maize RD22-like genes

**Authors:** Kyle Phillips<sup>1</sup> & Ndiko Ludidi<sup>1,2\*</sup>

**Affiliations:** <sup>1</sup>Plant Biotechnology Research Group  
Department of Biotechnology  
Life Sciences Building  
University of the Western Cape  
Robert Sobukwe Road  
Bellville, 7530  
South Africa

<sup>2</sup>DST-NRF Centre of Excellence in Food Security  
University of the Western Cape  
Robert Sobukwe Road  
Bellville, 7530  
South Africa

**\*Corresponding Author:** [nludidi@uwc.ac.za](mailto:nludidi@uwc.ac.za)



**Figure S1: Prediction of the subcellular localization of maize RD22 proteins. (a)** Subcellular localization of ZmRD22A based on Plant-mPLOC prediction. **(b)** Subcellular localization of ZmRD22B based on Plant-mPLOC prediction.

**Supplementary Table S1: Primer sequences used in semi-quantitative and quantitative PCR analysis.**

Gene name	Forward primer	Reverse primer
ZmRD22A	5'-GCG GGC GGG CGG CGG GCG CCT G-3'	5'-TCA GCCGCC GCG GGT CCAGAC GAC G-3'
ZmRD22B	5'- CGA CGA CGA CGG CCG GGT CGT G -3'	5'- TCA GCC GCT GCG GGT CCA GAC GAC GTG -3'
Zm18s rRNA	5'-CCA TCC CTC CGT AGT TAG CTT CT -3'	5'-CCT GTC GGC CAA GGC TAT ATA C-3'
Zm $\beta$ -tubulin	5'- AGC CCG ATG GCA CCA TGC CCA GTG ATA CCT -3'	5'- AAC ACC AAG AAT CCC TGC AGC CCA GTG C -3'
Zm Actin	5'- GTG ACA ATG GCA CTG GAA TG -3'	5'- GAC CTG ACC ATC AGG CAT CT -3'