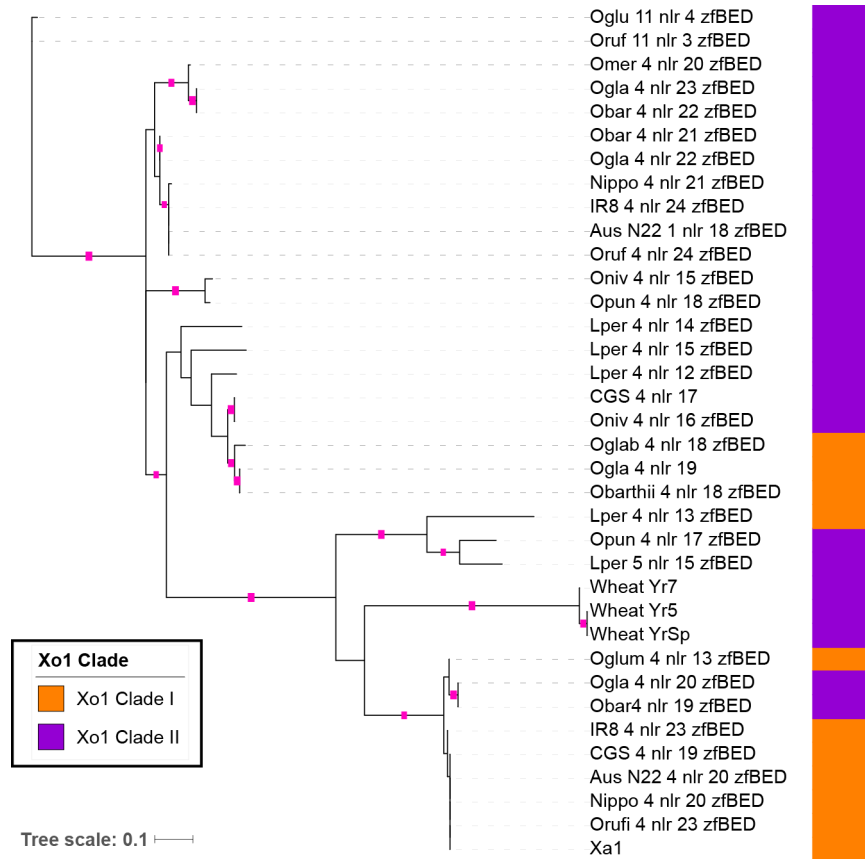


S3 Figure

Maximum likelihood tree of NB-ARC domain from Oryzeae NLRs



Maximum likelihood tree of zfBED domain nucleotide sequences from Xo1 clade I and II NLRs. Branches with bootstrap support greater than 80 percent are indicated with pink squares. Interactive tree available at <http://itol.embl.de/shared/acr242>.