

Figure S3. Maximum-likehood phylogeny of Criollo and Arabidopsis PR-4 family members. Node labels represent bootstrap supportf rom 100 replicates. Branch lengths represent genetic distance in substitutions per site. AT5G05460 and Tc02\_g032300, two  $\alpha$ -expansins that are part members oft he PR-4 superfamily, were included as an outgroup.