

Abstract

Immune-related hybrid incompatibility (HI) is one form of post-zygotic reproductive isolation, frequently caused by allelic mismatches of genes involved in plant immunity.

Immune-related HI between the North European Landsberg *erecta* (*Ler*) and Central Asian Kashmir-2 (*Kas-2*) *Arabidopsis thaliana* (*A.t.*) accessions likely arose as a by-product of adaptation to the biotic environment. It is due to the recessive genetic interaction between a highly variable *RECOGNITION OF PERONOSPORA PARASITICA1* (*RPP1*)-like resistance gene cluster in *Ler* and *Kas-2* alleles of the receptor-like kinase-encoding gene *STRUBBELIG RECEPTOR FAMILY 3* (*SRF3*). Similar to *A.t.* autoimmune-mutants, this epistasis exposes a temperature-dependent trade-off between immunity and growth. Some *RPP1* receptors are known to be involved in pathogen effector recognition, culminating in effector-triggered immunity. However, the precise sub-cellular localization and molecular function of *SRF3* in *A.t.* was still undefined, when this work was initiated. Previous work showed that compatible *SRF3*-forms (*SRF3^{Ler}*) in *Kas-2* backgrounds dampen activation of defensive *MPK4* and *MPK6* upon perception of the bacterial PAMP-epitope *flg22* and enhance susceptibility to *Pseudomonas syringae*.

The work presented integrates *SRF3* in the context of PAMP-triggered immunity (PTI). *SRF3*-GFP localizes to the plasma membrane and plasmodesmata (PD). *SRF3^{Ler}*-GFP-associated proteins identified in a proteomics approach are enriched for plasmodesmal and plasma membrane proteins involved in PTI and callose deposition. Contrary to MAPK activation, presence of *SRF3^{Ler}* in a *Kas-2* genetic background leads to enhanced ROS burst and restriction of PD-flux upon *flg22* treatment, but renders *Kas-2* almost insensitive to seedling growth inhibition in the presence of *flg22*. The results presented, suggest that *SRF3^{Ler}* differentially affects independent PTI signalling branches downstream of the PRR complex and provide an example how studying immune-related HI can be used to identify molecular components involved in plant defence.