Identification of novel pathogenicity factors in the late blight pathogen, *Phytophthora infestans*

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🔊 Results



Conclusions

Transient RNAi is effective for identifying genes with a major role in pathogenicity. RNAi can inform hypotheses for downsteam analysis of pathogenicity in *P. infestans*.

Both formation of functional pre-infection structures and pathogenicity require the action of many genes.

🔰 <u>Future work</u> 賌 <u>Ackno</u>

Localisation of structural proteins by translational fusion to fluorescent proteins, and subsequent observation by confocal microscopy.

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