Characterization of the barley (*Hordeum vulgare*) phyllosphere

C. Gravouil^{1,3}, J.M. Fountaine², A.M. Macrae¹, I. Hein¹, M.J. Dickinson³, A.C. Newton¹ 'scottish Crop Research Institute, Invergowrie, Dundee, DD2 5DA, UK 'scottish Agricultural College, West Mains Read, Edinburgh, EHP 3JG, Scotland, UK: 'School of Biological Sciences, University of Nottingham, Sutton Bonington Campus, Loughborough, LE12 5RD, UK Email: clement_gravoull@sci.ac.uk

Microbial interactions in the phyllosphere are poorly characterized but could provide a valuable approach to directly or indirectly control diseases or improve tolerance[‡]. The aim of this work is to develop and use molecular techniques to characterise the dynamics of microbial populations on barley leaves. We focus pathogen, *Pectobacterium atrosepticum* (Pba) the causal agent of potato blackleg disease, on the barley on the effects of barley leaf-associated microbes and a particular non-host foliar disease *Rhynchosporium secalis* (Rs)^o.



phyllosphere at a molecular level.

 The composition of microbial communities on the barley phylloplane can be affected by location, cultivar and physicochemical properties.

 Various microbes live on the phylloplane and can assist or compete with Rs pathogenicity.

 Pba is present on leaves of most barley cultivars from the field and has the ability to control Rs. Further identify microbes (bacteria, yeasts and fungi)
present on the leaf surface with molecular tools and characterize the chemistry of leaf waxes and leachate.
Identify effects of treatments (fungicide, elicitor,

combination) on microbial dynamics in the field.

o Intensively characterise the Pba-Rs interaction using

confocal microscopy and micro-array.

Acknowledgement We would like to acknowledge Dr Ian Toth's team and Dr Tim Daniell from the SCRI for providing help with the Pba work and the T-RFLP analysis respectively. This work is funded by the Scottish Crop Research Institute (SCRI), the University of Nottingham, the British Society of Plant Pathology (BSPP), the Society of Experimental Biology (SEB) and the Scottish Government Rural and Environment Research and Analysis Directorate (RERAD) – Sustainable Agriculture–Barley Pathology Workpackage.

BSPP

SEB

livina technoloa

Nottingham

Material and methods • Barley was grown under field conditions with no treatment applied. Leaves were sampled at growth stage 30 and stored at -20°C. • The composition of bacterial communities were determined using Terminal-Restriction Fragment Length Polymorphism (T-RFLP) based on 16S

Terminal-Restriction Fragment Length Polymorphism (T-RFLP) based on 165 ribosomal DNA and analysed using Principal Component Analysis (PCA). • Culturable microbes were isolated on nutrient agar and CzV8CM supplemented with streptomycin and incubated at 18°C. • *in planta* effect of leaf-isolated micro-organisms was studied by pretreating barley with microbes two days before Rs infection. • Pba was detected using species specific primers^a and its effects on Rs were assessed *in vitro* and under field conditions, by growing barley on previously Pba-infected or non-infected fields.

The University of

References

[‡] Newton *et al.* (2010) Ann Appl Biol, in press.
⁶ Newton *et al.* (2004) New Phytol, 163: 133-138.
^a De Boer and Wart (1995) Phytopathology, 85: 854-858
[‡] Suda *et al.* (2008) Microbes Environ, 23: 248-252.