Diversity of the nitrite reductase (NirK) gene in an upland pasture system T.J. Daniell, J. N. Squires, L. Cardle*, K. Ritz~, R.E. Wheatley and B.S. Griffiths. Plant Soil Interface, SCRI, Invergowrie, Dundee, DD2 5DA, UK. Scottish Crop *Computational Biology, SCRI, Invergowrie, Dundee, DD2 5DA, UK Research Institute

Current address: National Soil Resources Institute, Cranfield University, UK.

Introduction

NO₃ Denitrification can be economically and environmentally undesirable ᡟ Serious contribution to global warming and ozone depletion NO₂ 2 nitrite reductase enzymes \rightarrow NO \rightarrow N₂O \rightarrow N₂O nflavoprotein Cu (Nir K)

Materials and methods

Experiment 1

Soil samples were taken from plots which had been rotovated and reseeded with either Lolium perenne or Agrostis capillaris or left fallow (fig 1). Total DNA was extracted and amplified with nirK primers. Clones were screened by RFLP and selected examples were sequenced and placed on a tree (fig 2). Types from this experiment are labelled A-V.



Experiment 2

Intact soil cores were sampled as shown in fig 1 (red circles), individual root fragments were isolated from the cores, DNA was isolated and plant species identified utilising molecular methods (University of York). From aliquots of the same DNA extract nirK PCR was performed and high throughput sequencing methods were applied. 96 clones from each sample were sequenced. Sequences were analysed and examples with 5 or more clones of each type are displayed in fig 2, types from this experiment are labelled 1-24 and single clones. Percentages of clones representing each sequence type and diversity indices are shown in table 1.Experiment 1 Soil samples were taken from



Fig 2: A Neighbor joining tree of *nirK* sequences. Database sequences are shown in black, sequences from experiment 1 are represented by circles and those from experiment 2 are represented by squares. Sequence groups with a total of less than 5 clones have been removed for clarity. Bootstrap values above 70% are shown.

| Plant species | | Replicate 1 2 | | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 | 24 | Singles Clones Shanno | | |
|---------------|-------------|---------------|------|------|------|------|------|------|------|------|-----|------|------|-----|------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----------------------|----|------|
| Agrostis ca | pillaris | 1 | 80.6 | 9.0 | 4.5 | | - | 1.5 | - | | | 1.1 | 1.0 | | 1.0 | - | - | | | | | - | 1.5 | 1.5 | | - | 3 | 67 | 0.78 |
| Agrostis ca | pillaris | 3 | 93.8 | 3.8 | - | 1.3 | | - | - | 1.3 | - | 1.0 | 1.1 | 1.1 | | - | - | - | | - | - | - | - | - | - | - | - | 80 | 0.29 |
| Agrostis ca | pillaris | 4 | 41.7 | 22.2 | 31.9 | - | - | - | - | - | 1.4 | 1.1 | 1.0 | | | - | - | - | | - | 2.8 | - | | - | | - | - | 72 | 1.22 |
| Agrostis ca | inina | 1 | 65.9 | 25.6 | 1.2 | | - | - | - | | | 1.1 | 1.0 | | 1.0 | - | - | 4.9 | | | | - | | 1.2 | | 1.2 | | 82 | 0.93 |
| Festuca rub | ora | 1 | 64.0 | - | 3.4 | 3.4 | 6.7 | - | - | - | 5.6 | 11.2 | 1.1 | 1.1 | | - | - | - | | - | - | - | - | - | 2.2 | - | 4.5 | 89 | 1.34 |
| Festuca rul | bra | 2 | 96.4 | 1.8 | | 1.8 | - | - | - | | | 1.1 | 1.0 | | 1.0 | - | - | | | | | - | | | | - | | 56 | 0.18 |
| Festuca rul | ora | 3 | 85.7 | 1.6 | 9.5 | - | - | - | - | 1.6 | | 1.1 | 1.0 | | 1.0 | - | - | | | | | - | | | | - | 1.6 | 63 | 0.55 |
| Festuca iur | ncifolia | 1 | 60.9 | 4.7 | 3.1 | 6.3 | | - | - | 6.3 | 1.6 | 1.0 | 1.0 | 3.1 | 12.5 | | - | | | - | | - | | - | | - | 3.1 | 64 | 1.40 |
| Festuca jur | ncifolia | 2 | 71.4 | - | - | 11.1 | | - | - | - | - | 1.0 | 15.9 | - | | - | - | - | | - | - | - | - | - | - | - | 3.2 | 63 | 0.84 |
| Festuca jur | ncifolia | 3 | 39.7 | | 2.7 | 1.4 | 42.5 | - | - | 1.4 | | 4.1 | - | 1.1 | 1.1 | | - | | | - | | 1.4 | | - | | - | 5.5 | 73 | 1.43 |
| Festuca jur | ncifolia | 5 | 71.4 | - | 6.3 | | - | - | - | 11.1 | 4.8 | 1.1 | 1.0 | | 1.0 | - | - | | 4.8 | | | - | | | | - | 1.6 | 63 | 1.02 |
| Poa pratent | sis | 2 | 33.3 | - | 14.8 | 18.5 | - | - | 3.7 | 18.5 | 9.3 | 1.1 | 1.0 | | 1.0 | - | - | | | | | - | | | | - | - | 54 | 1.69 |
| Poa praten: | sis | 3 | 72.4 | 17.2 | 4.6 | 2.3 | - | - | - | - | 3.4 | 1.1 | 1.0 | | 1.0 | - | - | | | | | - | | | | - | | 87 | 0.88 |
| Poa pratent | sis | 5 | 56.5 | 17.4 | 5.8 | 1.4 | - | - | 13.0 | - | - | 1.1 | 1.0 | | 1.0 | - | - | | 1.4 | | 1.4 | - | | | | - | | 69 | 1.36 |
| Holcus lana | atus | 1 | 47.5 | | 2.5 | - | | 11.3 | 15.0 | - | 3.8 | 1.0 | 1.0 | 1.1 | 1.1 | 8.8 | 6.3 | | | 3.8 | | - | | - | | - | 1.3 | 80 | 1.60 |
| Holcus lana | atus | 2 | 68.5 | 22.2 | 1.9 | 1.9 | - | 3.7 | - | 1.9 | - | 1.1 | 1.0 | | 1.0 | - | - | | | | | - | | | | - | - | 54 | 0.94 |
| Holcus Jana | atus | 3 | 50.8 | 6.8 | 5.1 | - | | 22.0 | 5.1 | - | 1.7 | 1.1 | 1.7 | 5.1 | 1.1 | | - | | | | | - | 1.7 | | | - | | 59 | 1.53 |
| Deschamps | sia cespito | sa 1 | 92.2 | 3.9 | 1.3 | - | | 2.6 | - | - | - | 1.0 | | | 1.1 | | - | | | - | | - | | - | | - | | 77 | 0.35 |
| Deschamps | a cesnito | sa 3 | 79.8 | | 67 | 9.0 | - | - | | | | 1.1 | 1.1 | 3.4 | 1.1 | | - | | | | | - | | | | 1.1 | | 89 | 0.74 |

Results and discussion

Samples from experiment 2 (rhizoplane) are dominated by one sequence type. This sequence type was not observed in experiment 1. Very few types are

shared between experiment 1 and 2.

This could be due to a number of reasons including:

- 1. Differences in rhizoplane and bulk soil communities of denitrifiers
- 2. Spatial differences in community structure
- 3. Differences in N input (experiment 1 was fenced, experiment 2 was open)

4. Effects of harsh physical disturbance

5 Effects of monoculture

NirK diversity is higher in the rhizoplane of sub-dominant arass species Samples from

Acknowledgements

This work is part of the Scottish Executive Environment and Rural affairs Department MICRONET project, funded under the Flexible Funding Scheme. Plant Identification was undertaken by Karyn Ridgway and Peter Young, University of York.