

OUR LAND AND WATER

Toitū te Whenua, Toiora te Wai

# Benign de-nitrification in the subsurface environment

NSC Our Land and Water Annual Symposium, 11-12 April, Lincoln University, Christchurch

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\*Presenting on behalf of our students, post-doc and colleagues Mn Aldrin Rivas, Mr. Ahmed Elwan, Mr. Pete McGowan, Mr. Stephen Collins, Ms. Genevieve Smith, Ms. Heather Martindale, Dr. Uwe Morgenstern, Dr. Neha Jha, Dr. Andrew McMillian, Dr. Andrew Manderson, Dr. Lucy Burkitt , A/Professor David Horne, Professor Mike Hedley, Ms. Abby Matthews, and Dr. Jon Roygard







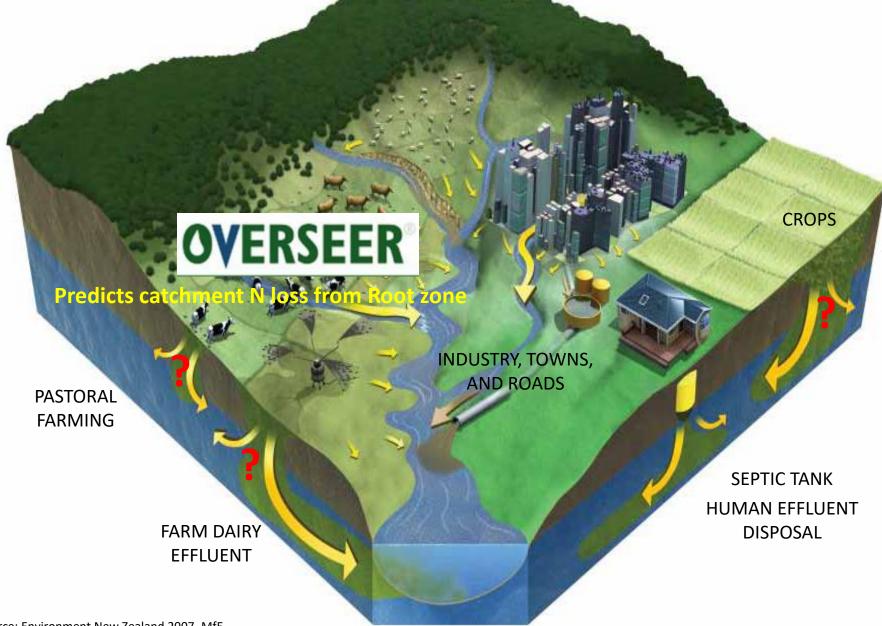


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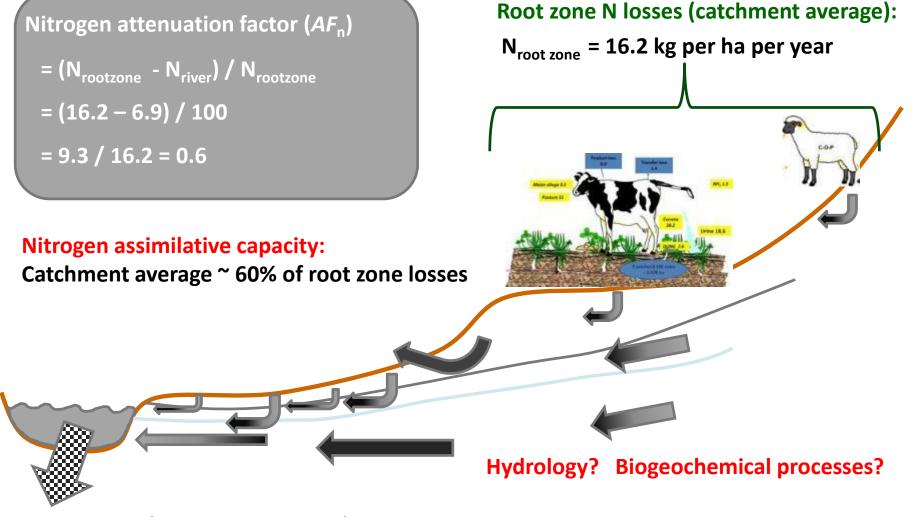
## Periphyton (benthic algae) Grows on the bed and on solid objects such as logs and stones in rivers

Associated with nutrient enrichment (excess of nutrients, nitrogen and phosphorus)

#### Sources and contributions to nutrient loadings?



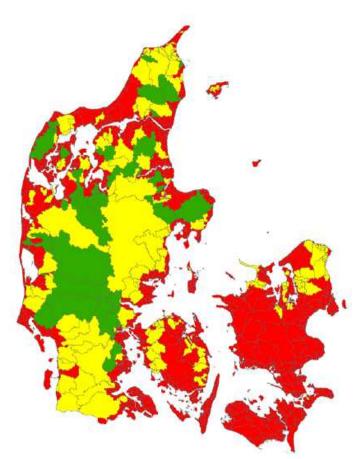
#### Manawatu Catchment @ Upper George



River N load (catchment average): N<sub>river</sub> = 6.9 kg per ha per year

Source: Elwan et al. (2015), Massey University

#### Sustainable primary production



Example: The Danish national map of nitrate reduction classes. (Source: Ernstsen et al., 2008)







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Nitrogen Attenuation Capacity

Green > 80 % N reduction

Yellow

Red

50 – 80 % N reduction Medium Capacity Areas: Reduce Nitrogen Leaching via Best Effluent and Nutrient Management Practices

**Targeted investment** 

in solutions, e.g.

**High Capacity Areas:** 

Sustainable Land Use

Intensification

N reductio

< 50 % N

reduction

Low Capacity Areas: Duration controlled grazing Sheep/Goat milking Cut and Carry Systems

#### **Programme Co-ordination**



Dr. Ranvir Singh



Assoc. Prof. Dave Horne



Dr. Uwe Morgenstern



Ms. Abby Matthews Dr. Jon Roygard





Prof. Mike Hedley

#### **Programme Partners**



UNIVERSITY OF NEW ZEALAND







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#### Developing techniques, methods and models

**Objective - Assess and map nutrient flow pathways and their potential attenuation** 

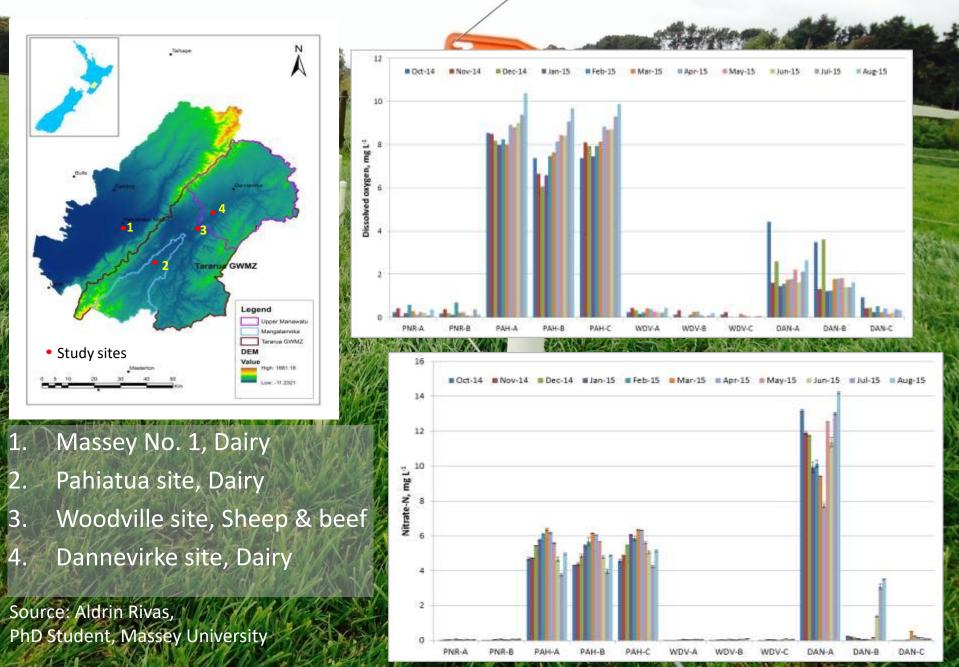
200 cm

Four piezometers at depth ranging from 5.8 To 8.7 m below ground level (bgl)

Suction cups (depth, bgl) 100 cm 60 cm

30 cm

#### In-field monitoring and observations

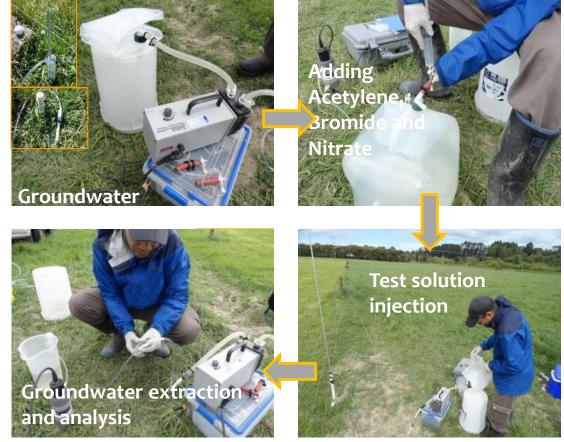


#### **Methods:**

- Lab incubations and in-field 'push-pull' tests
- Isotope tracer techniques
- Excess N<sub>2</sub> (being developed by GNS Sciences)
- Molecular techniques

   (being developed by Massey FLRC and Landcare Research )

#### Single Well 'Push-Pull' Test









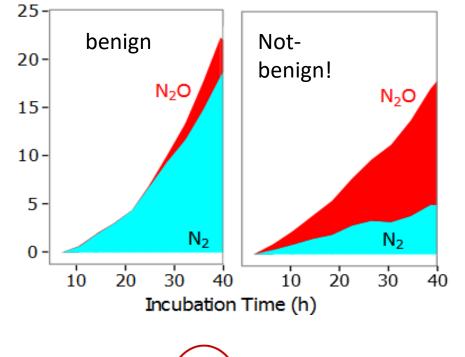
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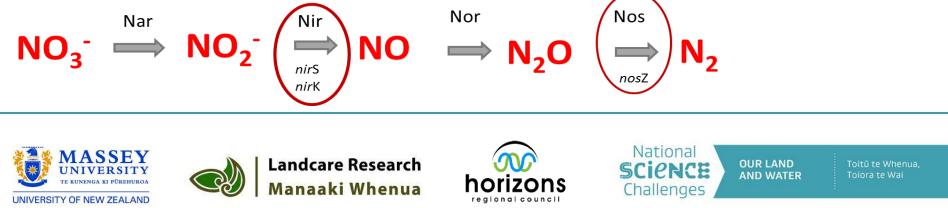


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#### Groundwater de-nitrification? benign or not-benign?

- Direct measurement of the terminal products of denitrification, N<sub>2</sub>O and N<sub>2</sub>
- Direct measurement of the de-nitrifiers, nirS, nirK and nosZ genes





# Relationships between nitrogen attenuation and catchment characteristics

Nitrogen attenuation factor  $(AF_n) = (N_{rootzone} - N_{river}) / N_{rootzone}$ 

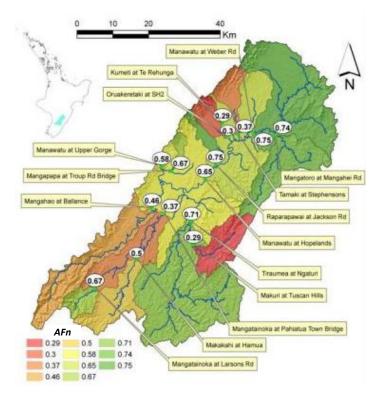
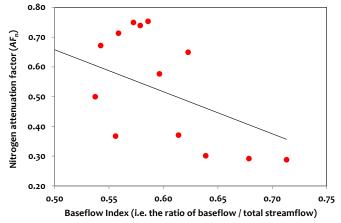


Table 1: Results of linear regression analysis between the  $AF_N$  values and catchments characteristics

| Catchment Characteristics                                 | AI             | N     |
|---|----------------|-------|
| 53 <del>-</del>   | R <sup>2</sup> | p     |
| Well-drained (e.g. soils with<br>drainage class 5) soils* | -0.35          | <0.05 |
| Fine textured (e.g. clay<br>oam) soils                    | 0.37           | <0.05 |
| Base Flow Index (BFI)                                     | -0.31          | <0.05 |

"Soils with drainage class 5, in the Fundamental Soil Layer "FSL", are well-drained soils.



Spatial distribution of the nitrogen attenuation factor for 15 sub-catchments in the Tararua Groundwater Management Zone (TGWMZ) (Elwan et al, 2015).







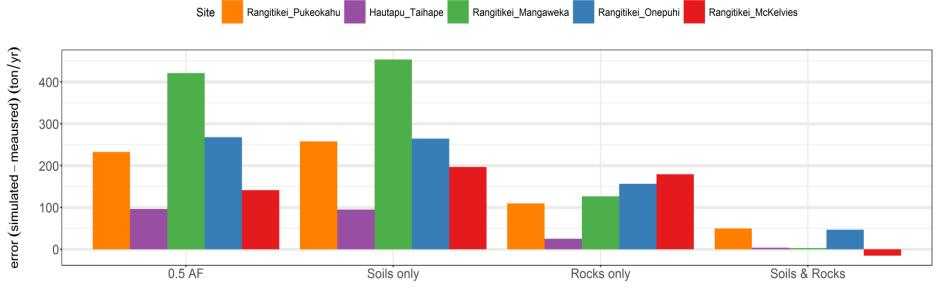
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#### Prediction of nitrogen loads in the Rangitikei River

Model - Variable nitrogen attenuation factor (based on soil and rock types – FSL and QMAP layers)

River N load (ton yr<sup>-1</sup>) = 
$$m \sum_{i=1}^{n} A_i * N_i * (1 - AF_{N_{RT}})(1 - AF_{N_{ST}})$$

Comparison of predicted vs. measured average annual soluble inorganic nitrogen (SIN) loads in different sub-catchments of the river

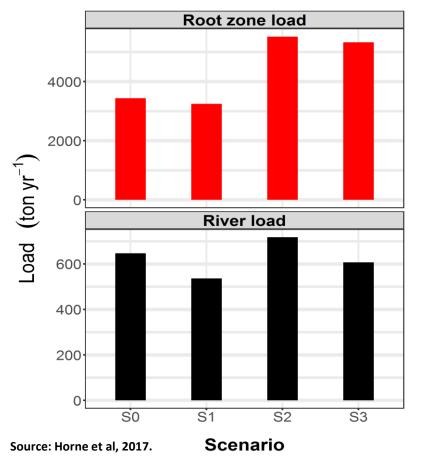


Source: Singh et al, 2017.



#### Prediction of nitrogen loads in the Rangitikei River

De-intensify (~9,800 ha) and Intensify (~83,000 ha) landuse (S3)



# Root zone N losses increase by 55%

#### River N load decreases by 6%







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## **Concluding Remarks**

- Opportunity to spatially align intensive high-value primary production with naturally high contaminant attenuation capacity areas
- Reduce water quality impacts, hence sustain and/or enhance cultural resources, mahinga kai, taonga species.
- Collaborative, co-developed, co-funded research programme
- Developing cost-effective practical techniques, methods and models
- Aligned with the OLW Challenge 'Sources and Flows' & 'Land Suitability' programmes for wider applications







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# Thank you – Questions and suggestions please!

## **Supporting Slides**





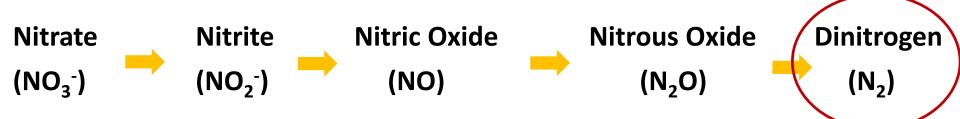


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De-nitrification is a biogeochemical process, which converts nitrate-nitrogen to gaseous forms of nitrogen; predominantly to dinitrogen in groundwater systems.



This capacity is mainly governed by the physical, chemical and biological characteristics, and importantly by nutrient and oxidisable carbon in flow pathways. It requires

- Low oxygen environment (influenced by hydrogeological settings);
- Carbon source (dissolved organic carbon); and
- Denitrify bacteria

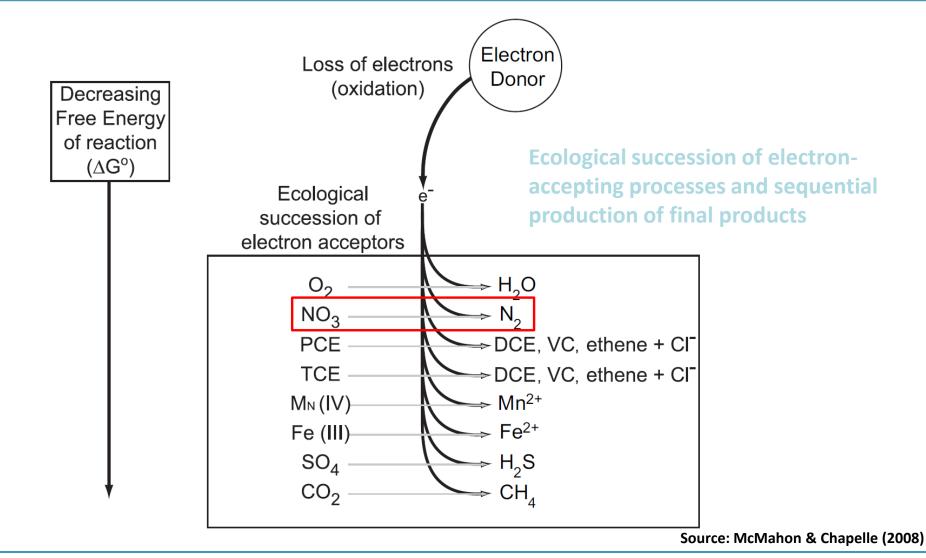




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|                               | Water Quality Criteria (mg/L) |                    |                  |                  |                   |  |
|-------------------------------|-------------------------------|--------------------|------------------|------------------|-------------------|--|
| Redox Process                 | 02                            | NO <sub>3</sub> -N | Mn <sup>2+</sup> | Fe <sup>2+</sup> | SO4 <sup>2-</sup> | Comments   |
| Oxic                          |                               |                    |                  |                  |                   |  |
| O <sub>2</sub> reduction      | $\geq 0.5$                    | <del></del>        | < 0.05           | < 0.1            | (                 | —  |
| Suboxic                       |                               |                    |                  |                  |                   |  |
| _                             | <0.5                          | <0.5               | <0.05            | <0.1             | _                 | Further definition<br>of redox processes<br>not possible |
| Anoxic                        | 401 - 2014 - C                | AND MADE MADE      |                  |                  |                   |  |
| $NO_3^-$ reduction            | < 0.5                         | $\geq 0.5$         | < 0.05           | < 0.1            | <u></u> 3         |  |
| Mn(IV) reduction              | < 0.5                         | < 0.5              | $\geq 0.05$      | < 0.1            |                   | —  |
| $Fe(III)/SO_4^{2-}$ reduction | < 0.5                         | < 0.5              | <u> </u>         | $\geq 0.1$       | $\geq 0.5$        | -  |
| Methanogenesis                | < 0.5                         | < 0.5              |                  | $\geq 0.1$       | < 0.5             | _  |
| Mixed                         |                               |                    |                  |                  |                   |  |
|                               | _                             | <u></u>            |                  |                  |                   | Criteria for more  |

#### Source: McMahon & Chapelle (2008)



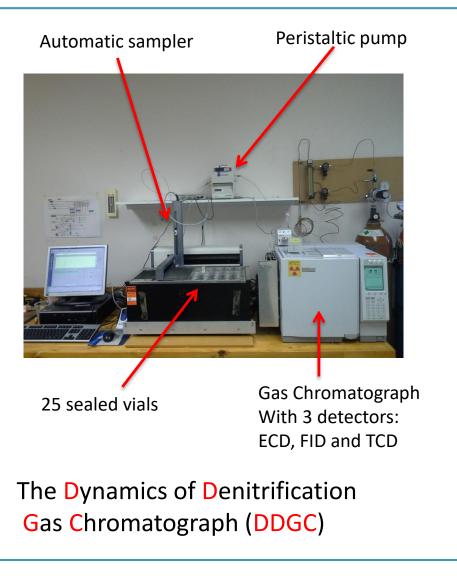


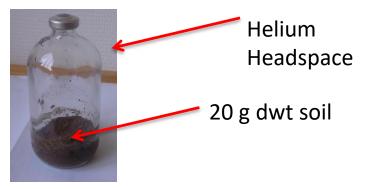
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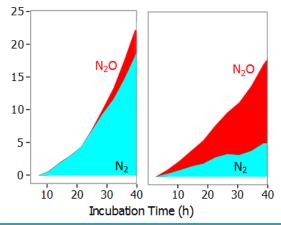
#### Direct measurement of the terminal products of denitrification, N<sub>2</sub>O and N<sub>2</sub>





125 mL Serum Vial

how "benign" the de-nitrification, i.e. calculate the  $N_2O:N_2$  ratio





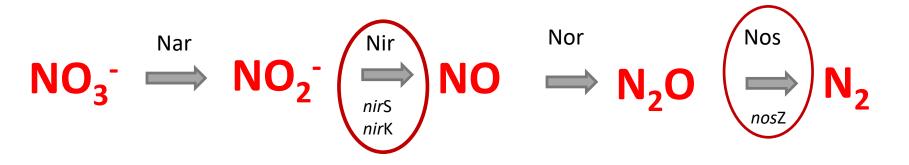


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#### Direct measurement of the de-nitrifiers, nirS, nirK and nosZ genes



- Anaerobic respiration
- Phylogenetically diverse: Bacteria, Archaea, Fungi, Protozoa







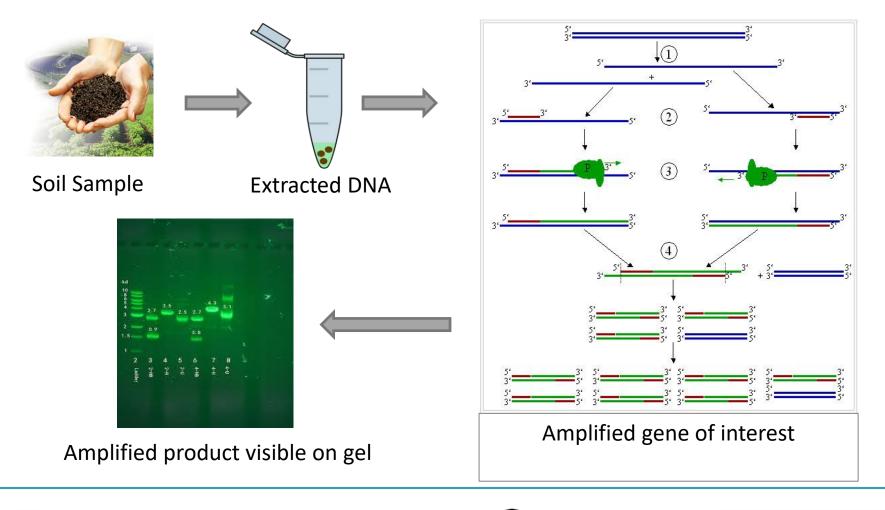
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#### Direct measurement of the de-nitrifiers, nirS, nirK and nosZ genes

#### **Polymerase Chain Reaction (PCR) technique**







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