



**For now &
our future**

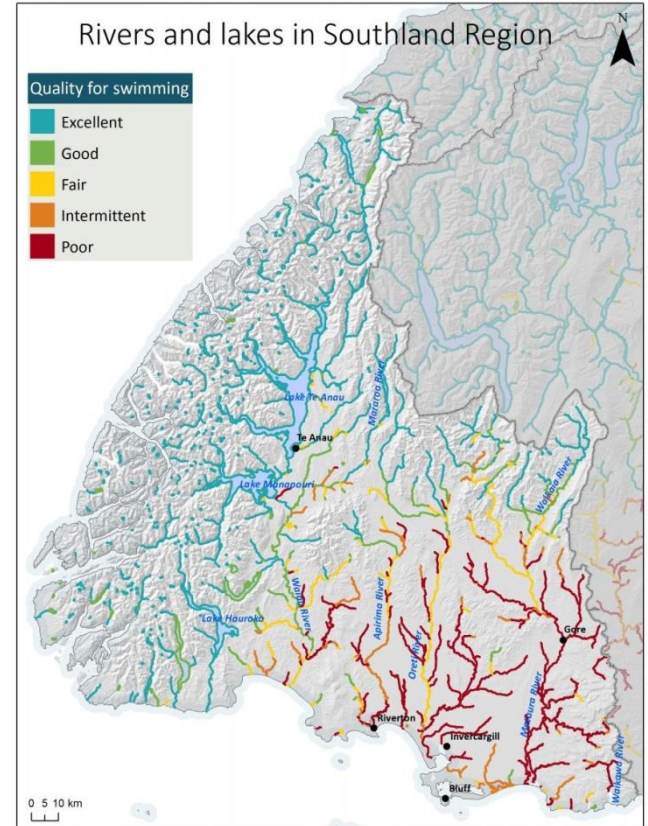
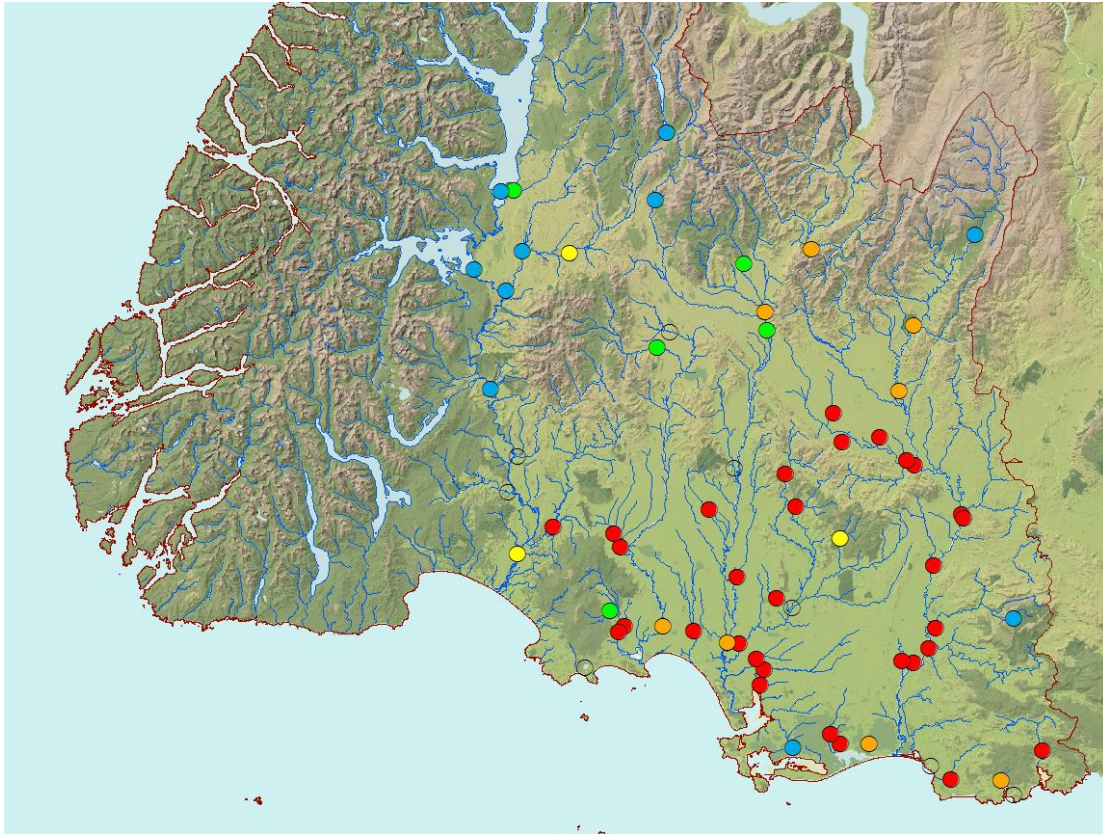
Where does microbial contamination come from?

Nick Ward Team leader – ecosystem response

Acknowledgements

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- Chris Owen (Contractor Southern Waterways)
- Graham McBride (NIWA – Retired)
- Richard Muirhead (Ag Research)
- Chris Palliser (NIWA)
- Rob Williamson (ES Science Assistant)

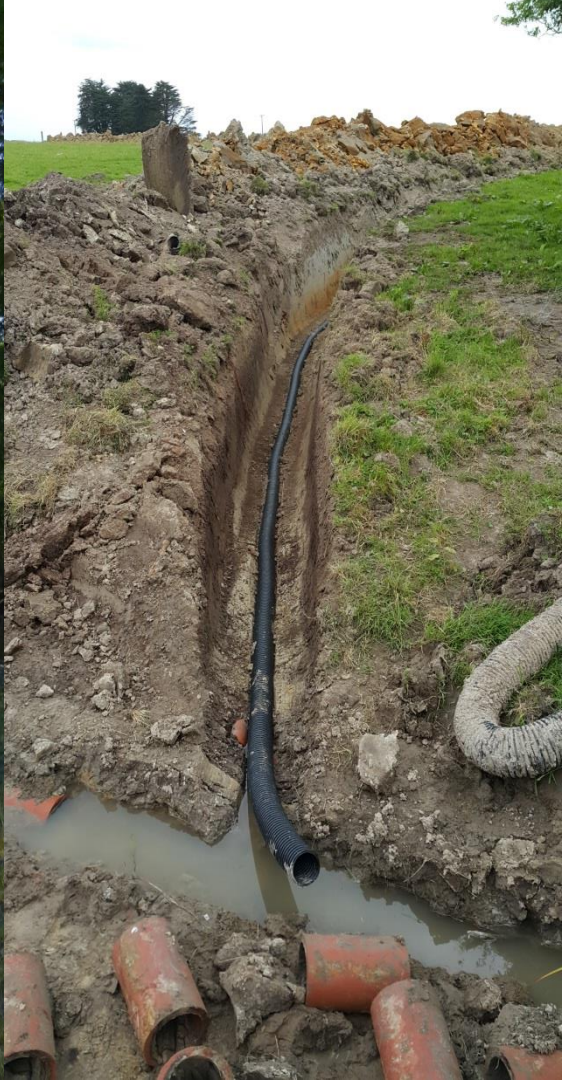
What's the problem?



Sources



Pathways

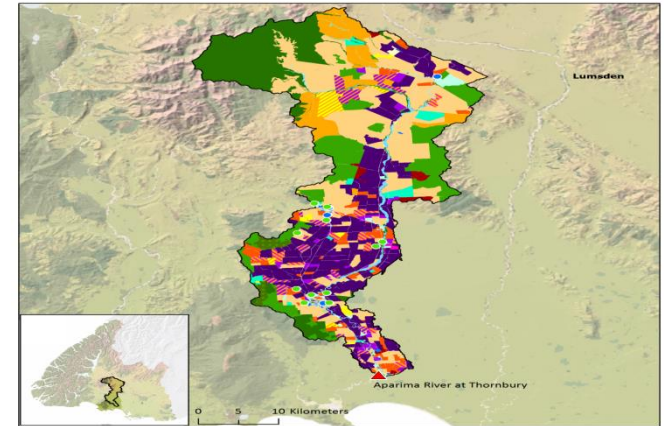


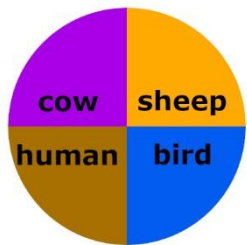
Sources of faecal pollution

- Over 250 samples analysed
- PCR Markers, Campylobacter
- Human, ruminant (sheep or cow), wildfowl

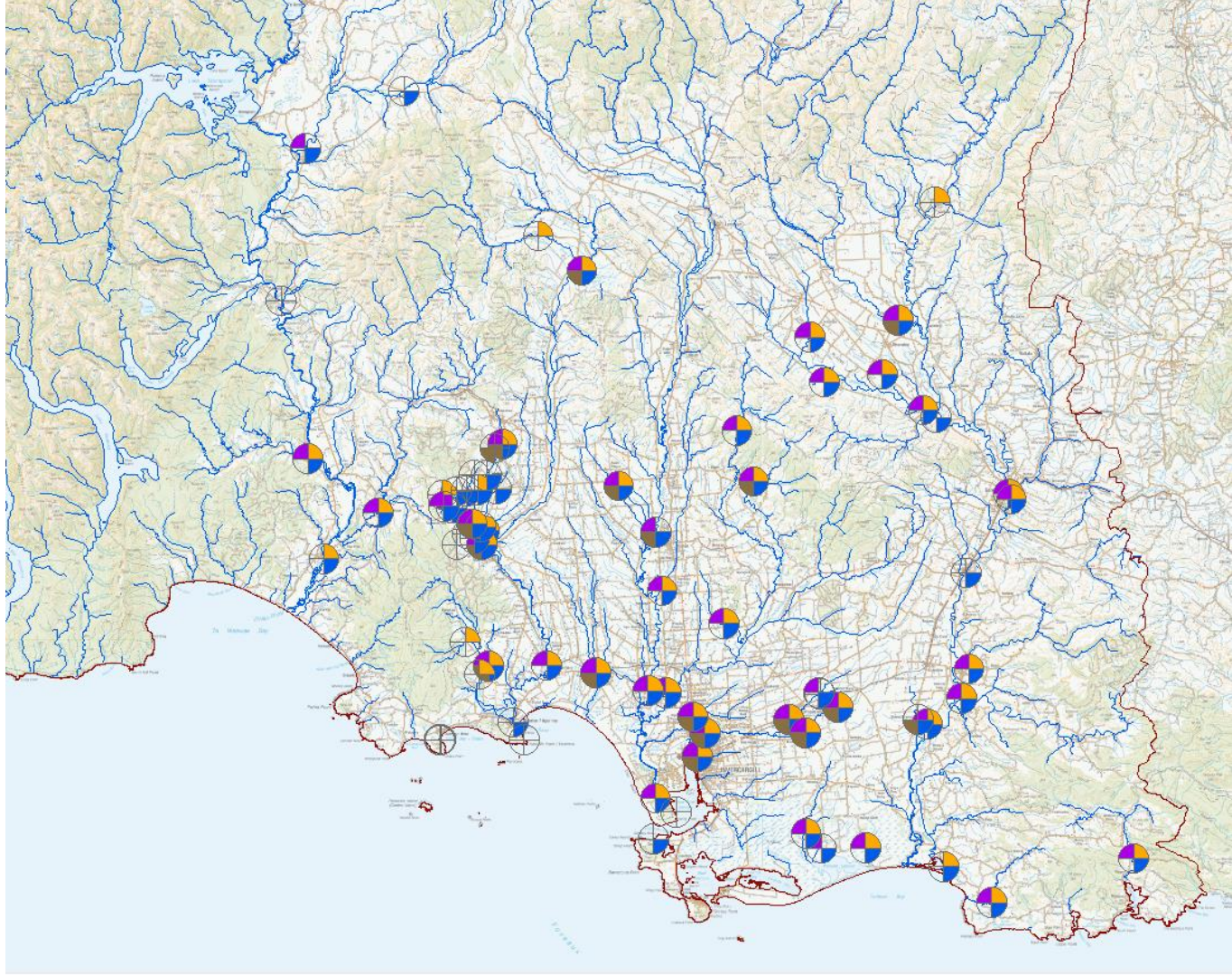


simple
vs
complicated

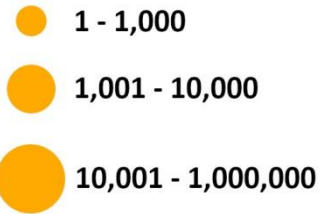




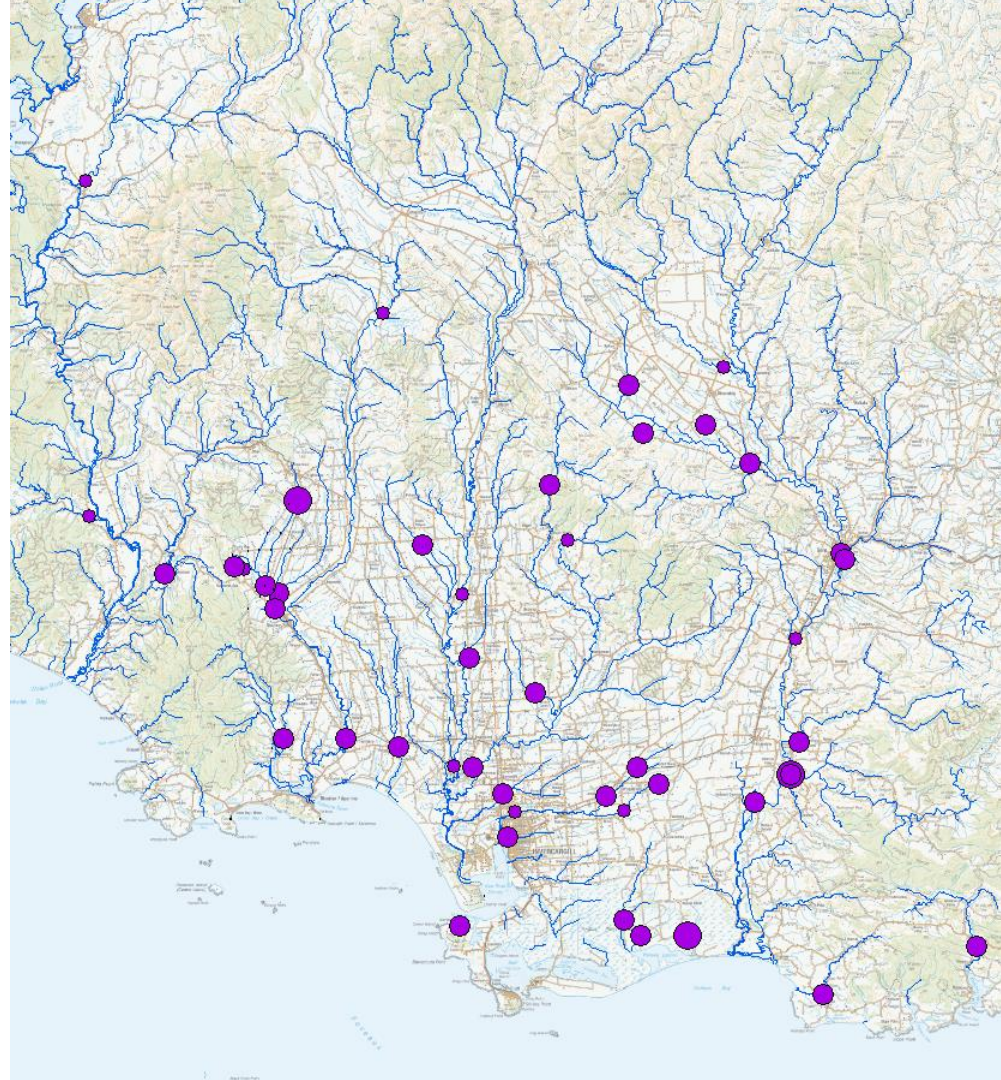
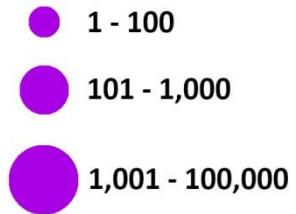
Absence/Presence



Maximum sheep PCR copies



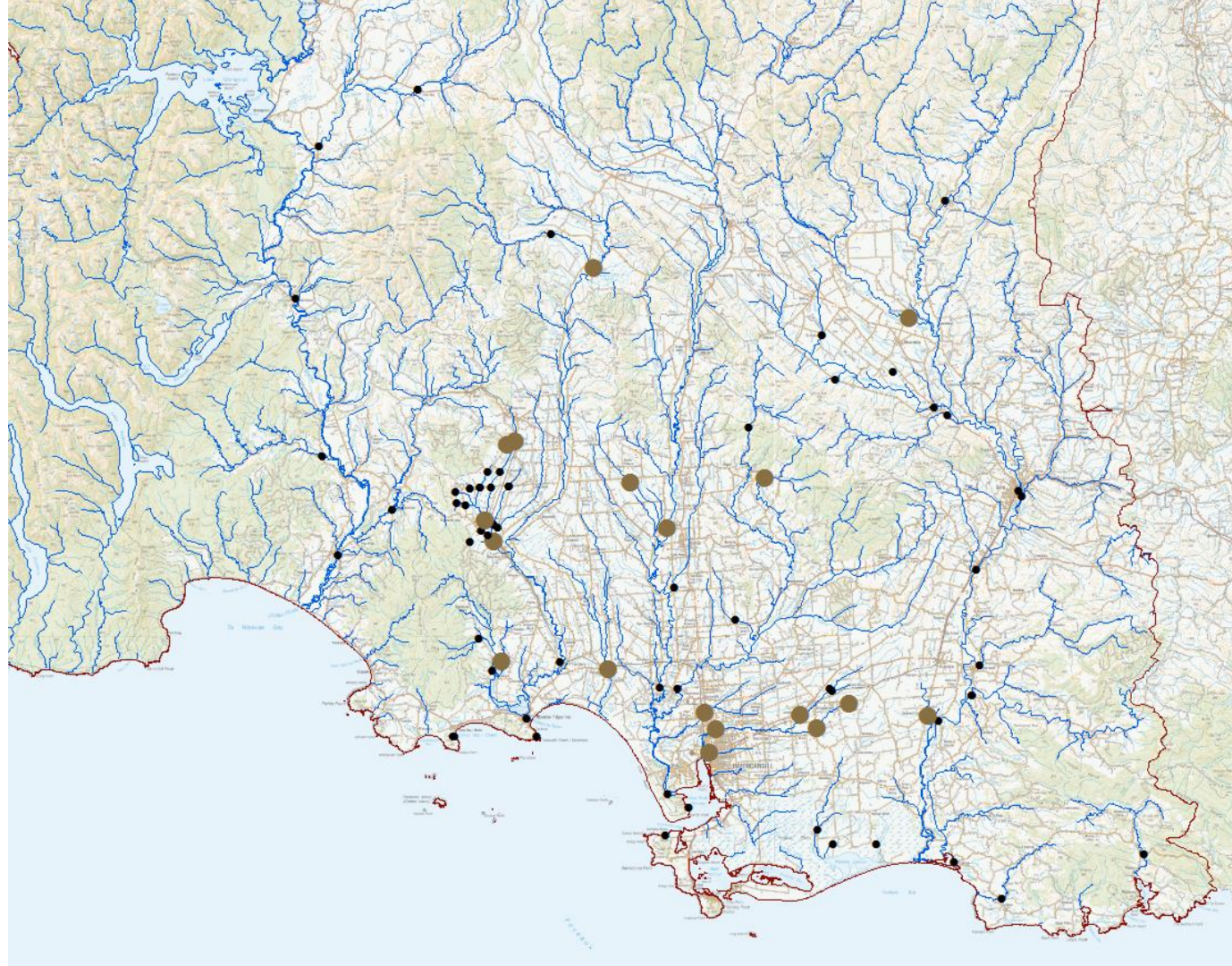
Maximum cow PCR copies



Human source

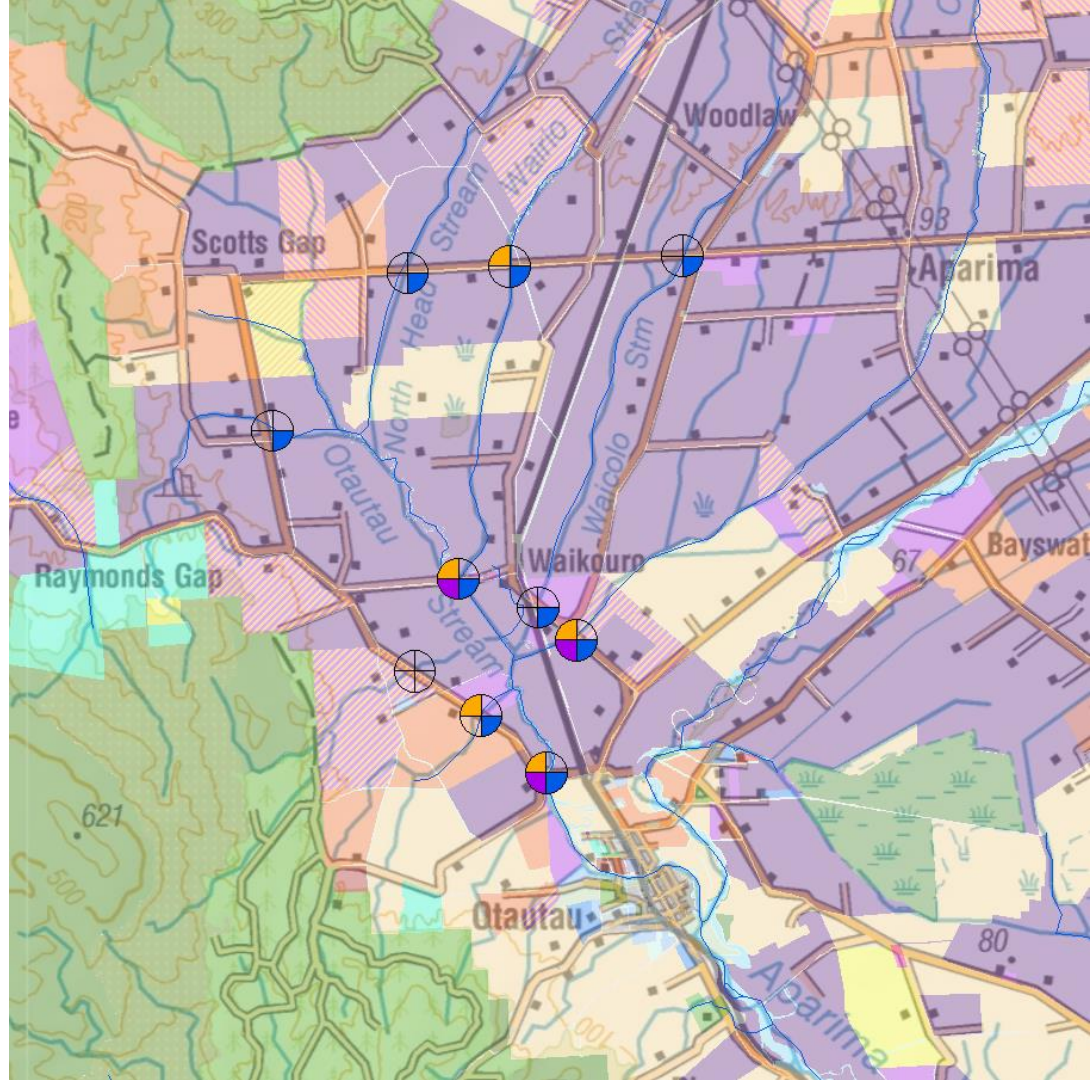
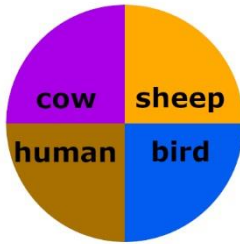
● Absence

● Presence



Otautau Stream

Sub-catchment study



Thoughts

- Can you target policy spatially?
- What about microbial antibiotic resistance?
- Flow separation and geochemical tracing

Integration

- Every site is different – use context
- Microbial is just the tip of the iceberg
..... but great segway

