

# Genome analysis of hybridisation between kakī and poaka

Natalie Forsdick



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#TeamKakī  
#ConSERTeam

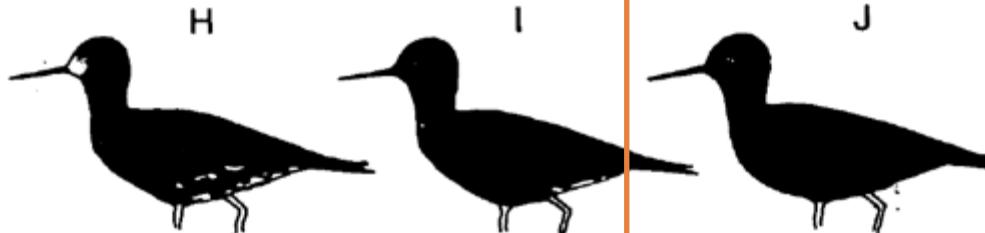
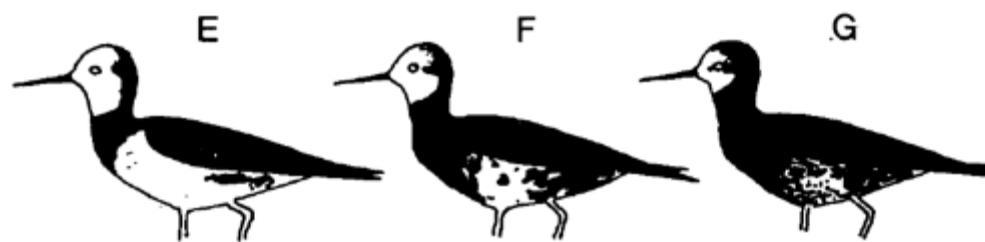
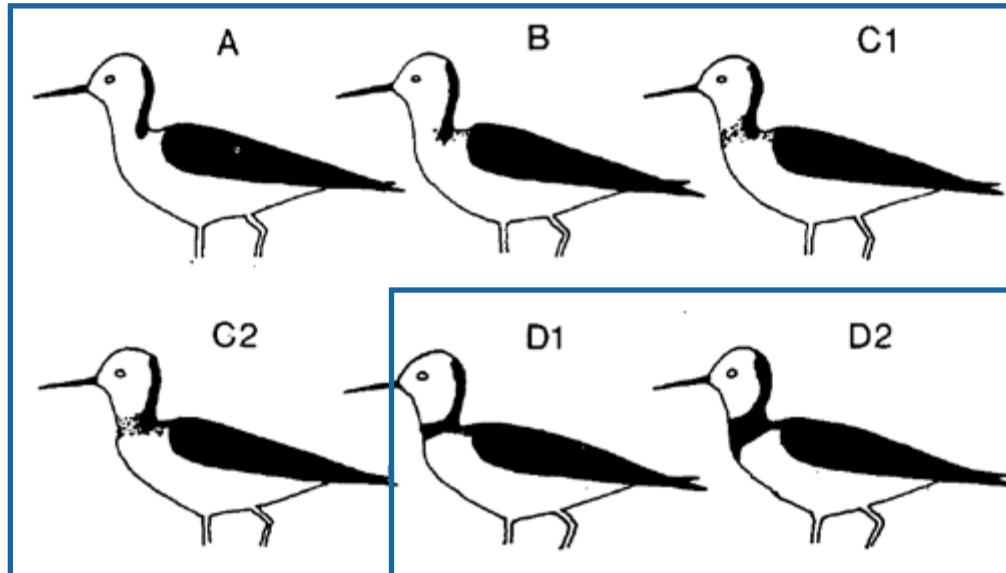
# Hybridisation with poaka



Edin Whitehead

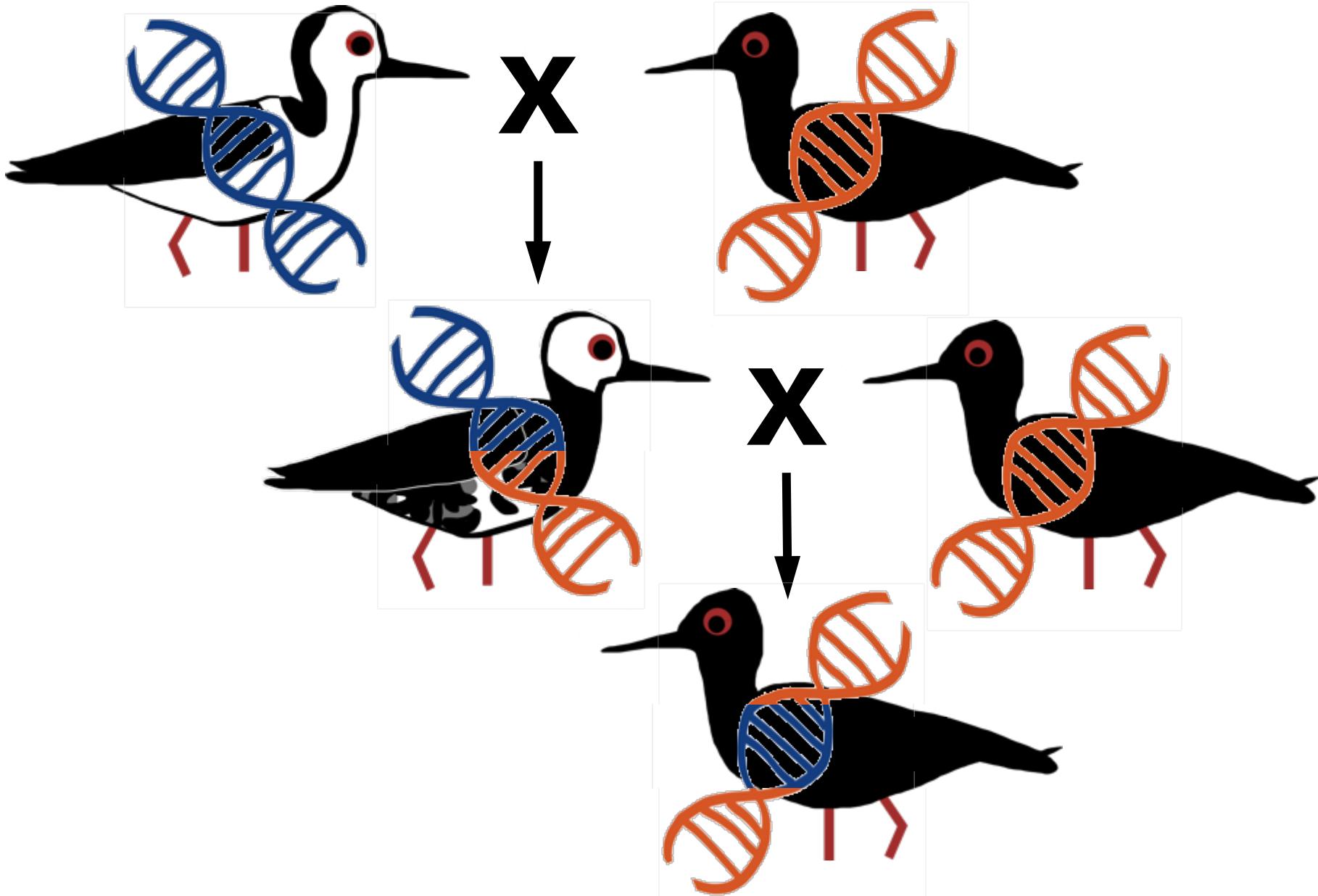
# Kakī x Poaka

Poaka



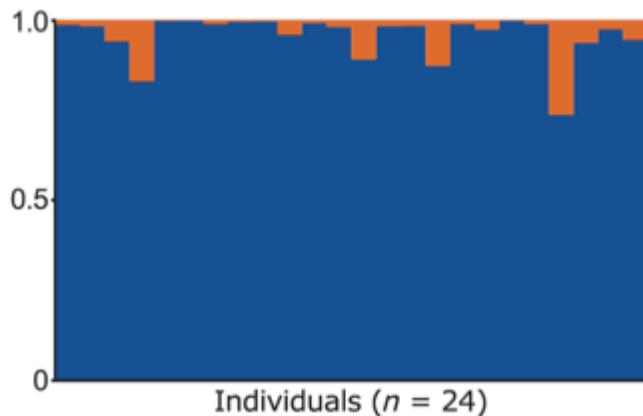
Kakī

# Genetic admixture

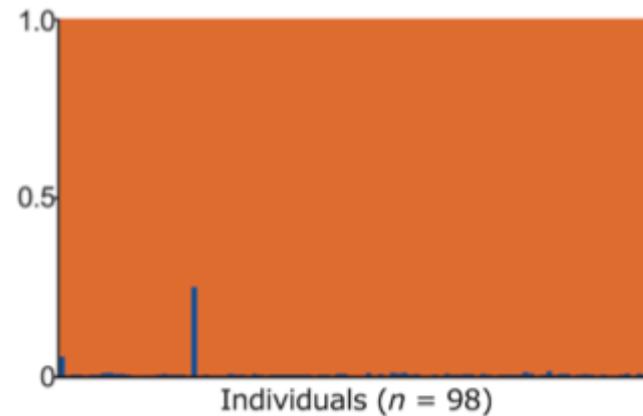


# Genetic analysis

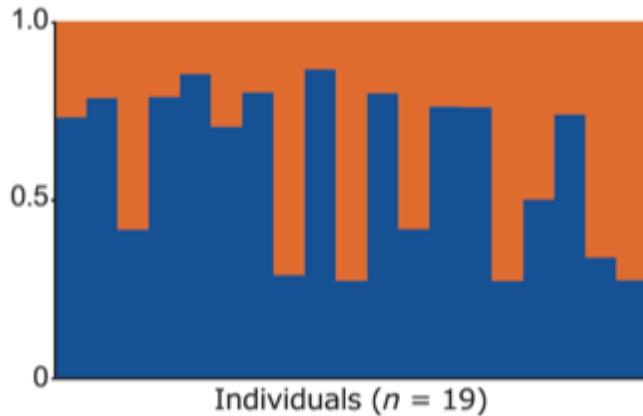
Poaka



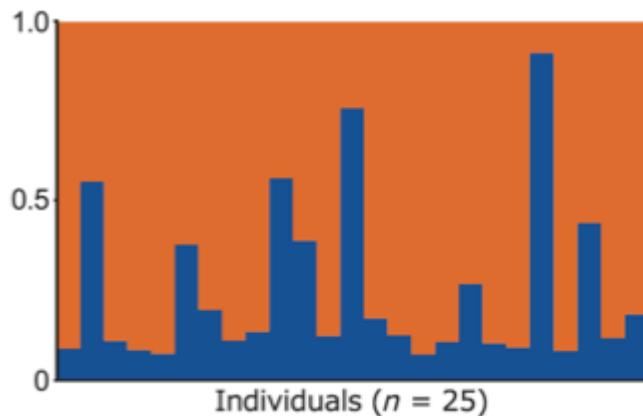
Kakī



Light hybrids



Dark hybrids



# Why use genomics?

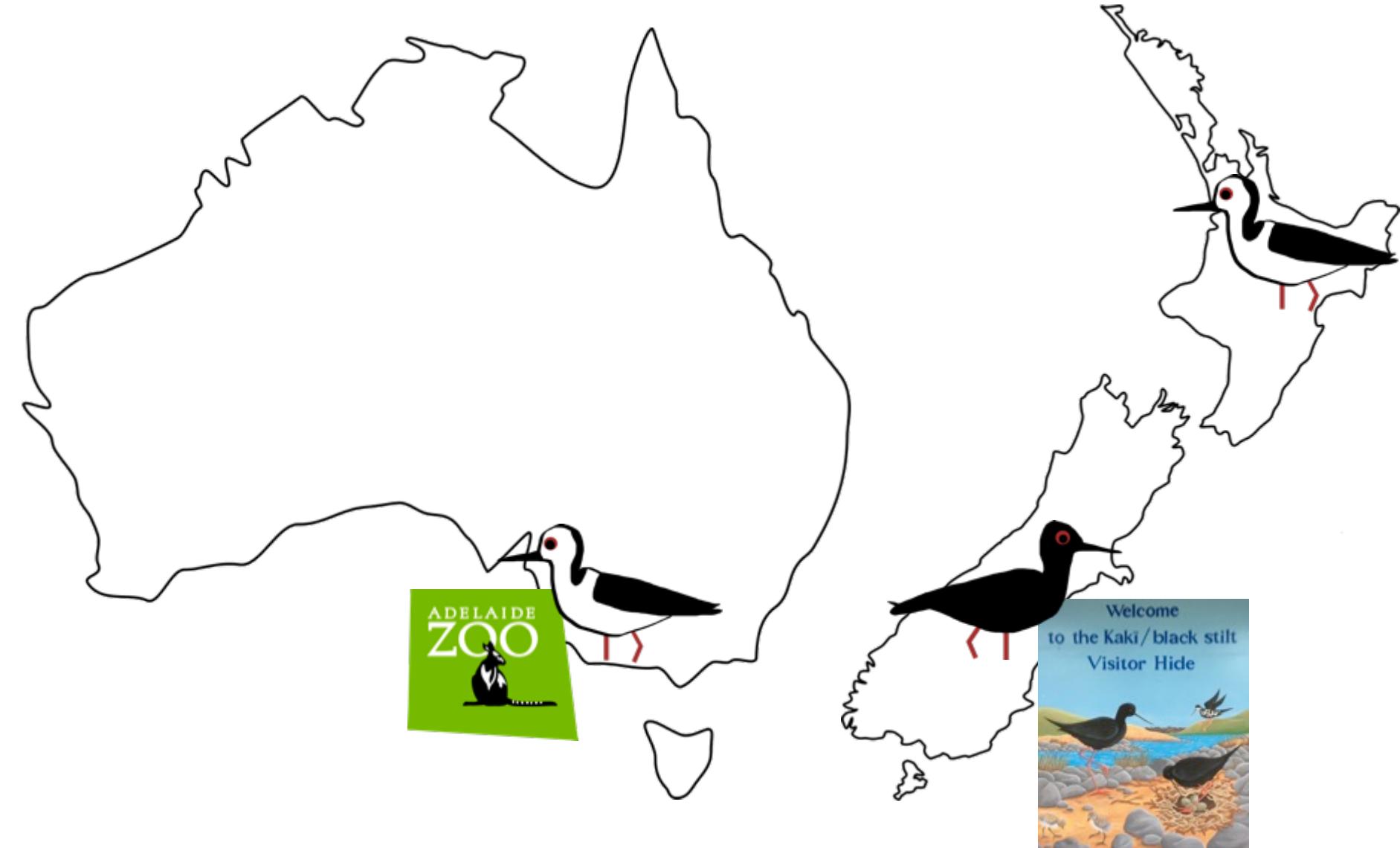
Whole genome sequencing

Step up from using ~10 markers to 1000s

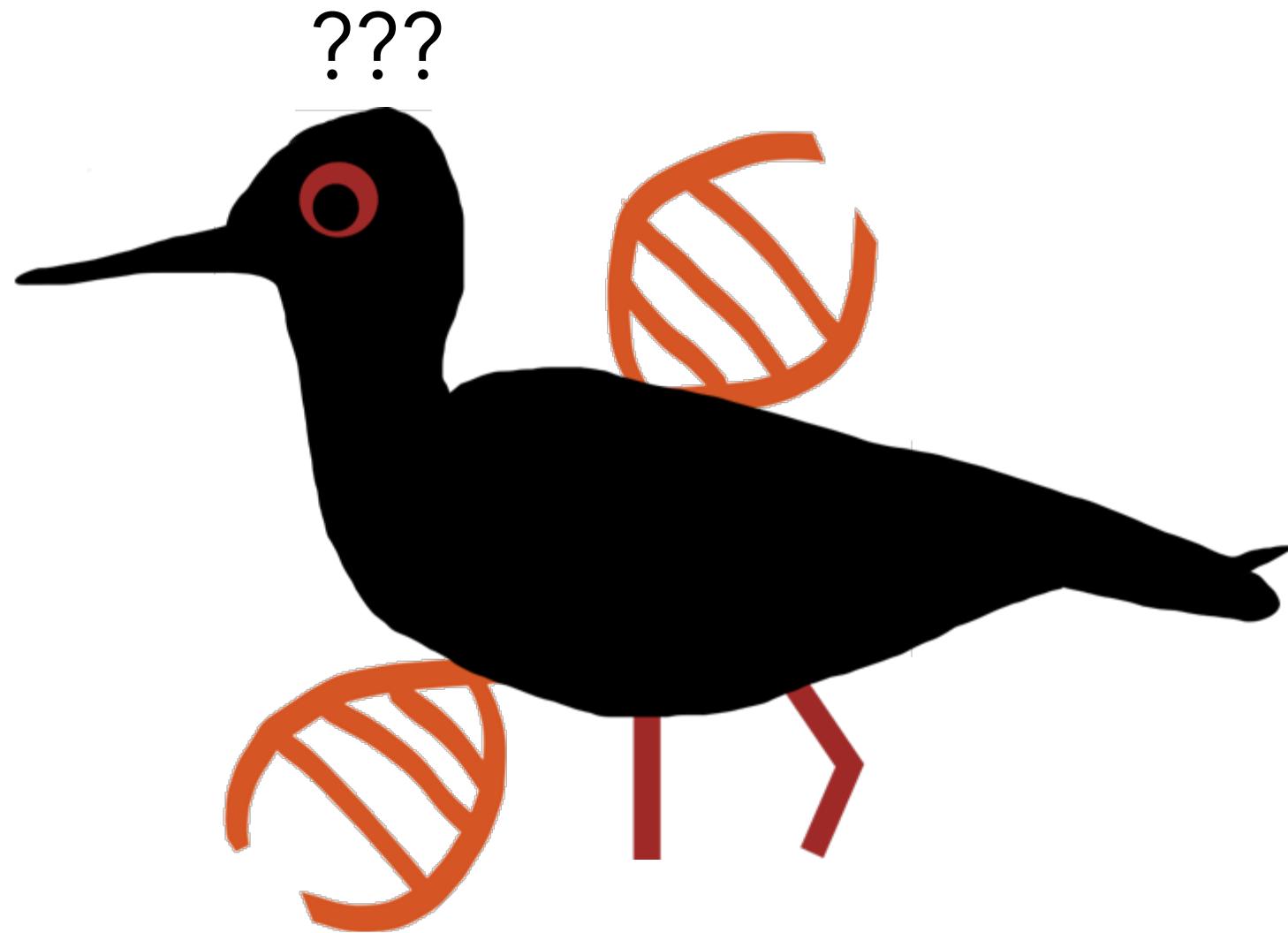
Greater power to detect differences/similarities

Ability to detect both neutral and adaptive variation

# Comparative approach

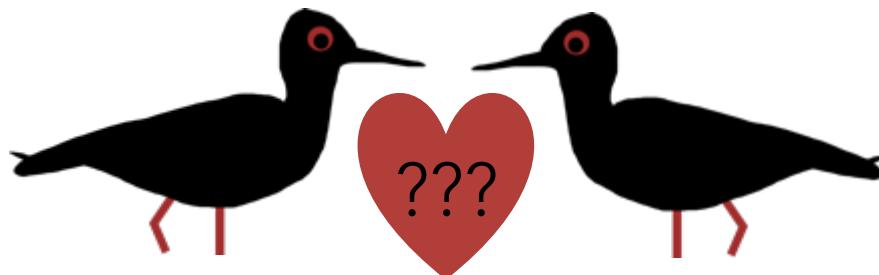


# No admixture?

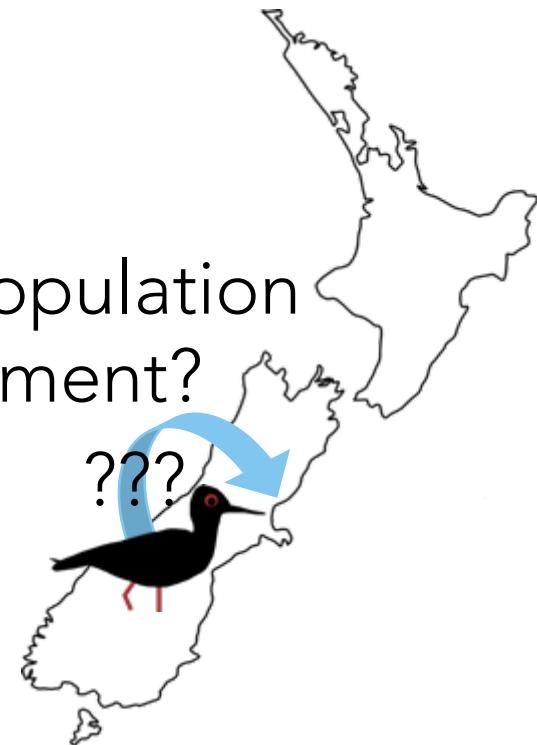


# Limited admixture?

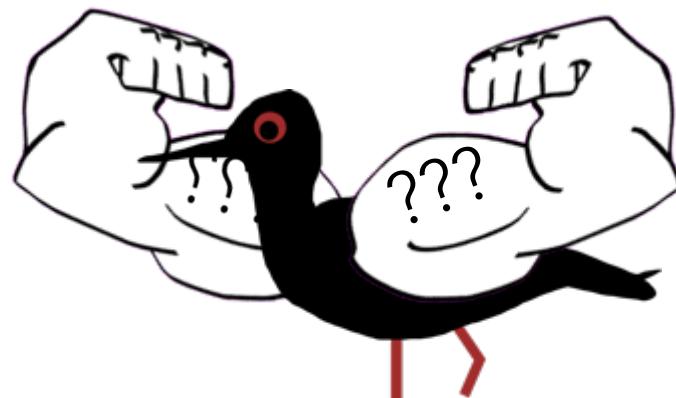
Captive breeding?



Future population establishment?



Benefits to recovery?



# Acknowledgements

- Michael Knapp, Tammy Steeves, Lisa Matisoo-Smith, Graham Wallis.
- Liz Brown, Simone Cleland, Richard Maloney, Cody Thyne at the Kakī Recovery Centre, Twizel.
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- Adelaide Zoo, Auckland Zoo, John Berry.
- Stephanie Galla, Olga Kardailsky, Denise Martini, Les McNoe.



Stephanie Galla

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