

# Breeding better working dogs with genomics

Claire Wade University of Sydney



# TOPICS (for today)

- Farm dogs
- Labrador Retrievers (Field Trial)
  - Genetic mapping of “drive”

# The Farm Dog Project

- Identify valuable phenotypes (from a user perspective)
- Assess phenotypes
- Use phenotypes to inform better dog breeding and reduce wastage

Team members:

Paul McGreevy  
Elizabeth Arnott  
Jonathan Early  
Lisa Mascord



# Identifying valuable phenotypes:

- Discussion with stakeholders
- Observation of trialling metrics
- Analysis of relevant training literature



Fig 1. Top 10 terms for General Behavioural attributes

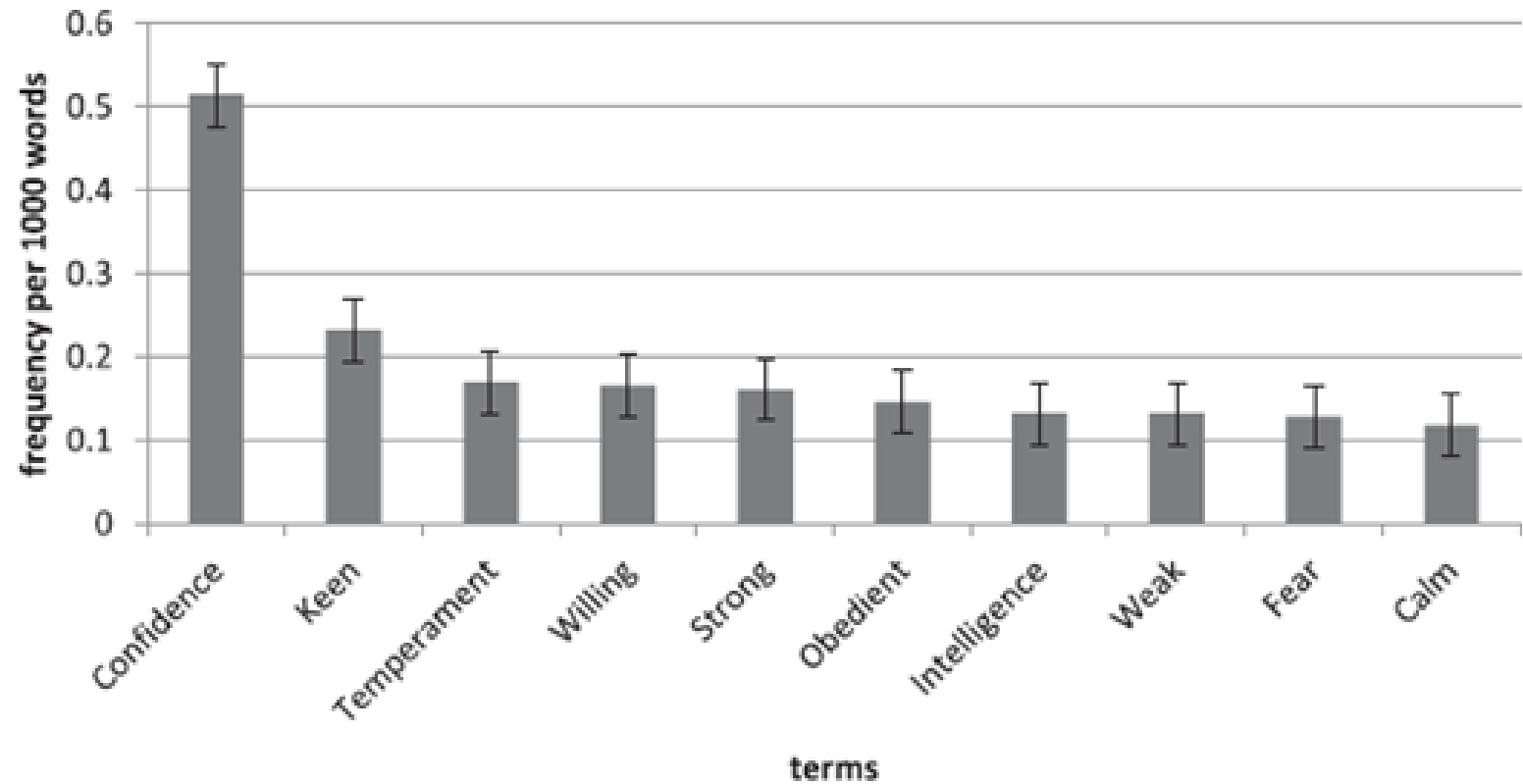
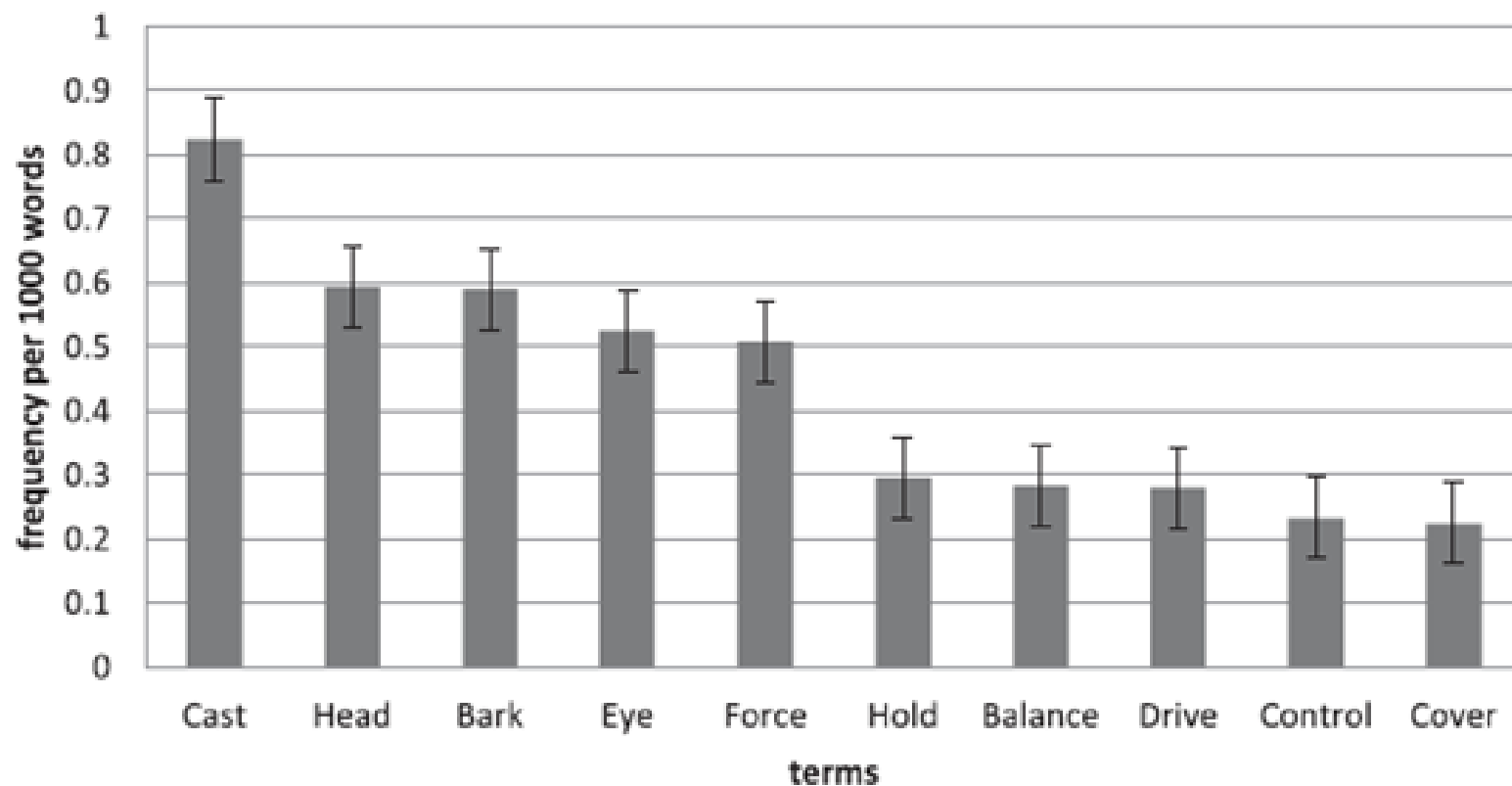
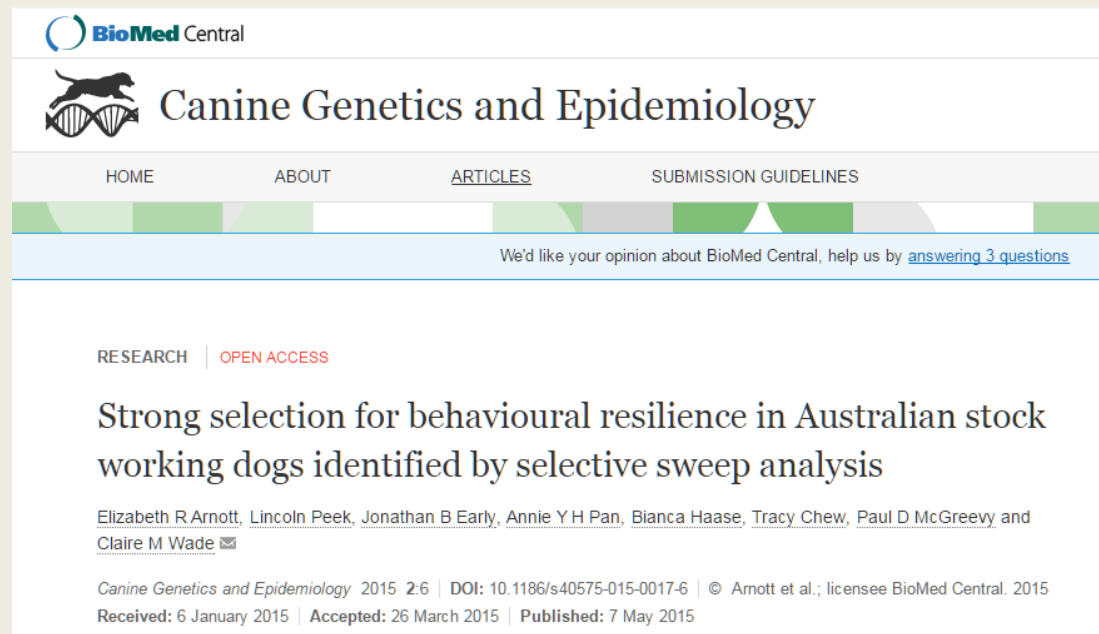


Fig 2. Top 10 terms for specialist working attributes of farm dogs



# Identifying valuable phenotypes:

- ...or looking at what breeders value without realizing it
- We do this by assessing the footprints of selection in the DNA



# Livestock working (Herding) Dog Assessment Form

This form is produced by The University of Sydney Farm Dog Project to collect trait information on Australian stock herding dogs. For participant information on the project please go to:  
[http://sydney.edu.au/vetscience/research/animal\\_behaviour/farmdog/](http://sydney.edu.au/vetscience/research/animal_behaviour/farmdog/).

This form should take about 5 minutes to complete. It is important that all questions are answered. You may complete the form anonymously as you are not obliged to leave your personal details. Your personal details will not be linked to your responses and your contact details will not be passed on to any third party. By completing this form you will be contributing to our knowledge of stock herding dogs. The information you provide will assist with ongoing research at the University of Sydney. We believe this will benefit working farm dogs and their owners/handlers/breeders.

This project is supported by a grant from the Australian government's Rural Industries Research and Development Corporation, Meat and Livestock Australia and in kind, support from the Working Kelpie Council of Australia.

\*Required



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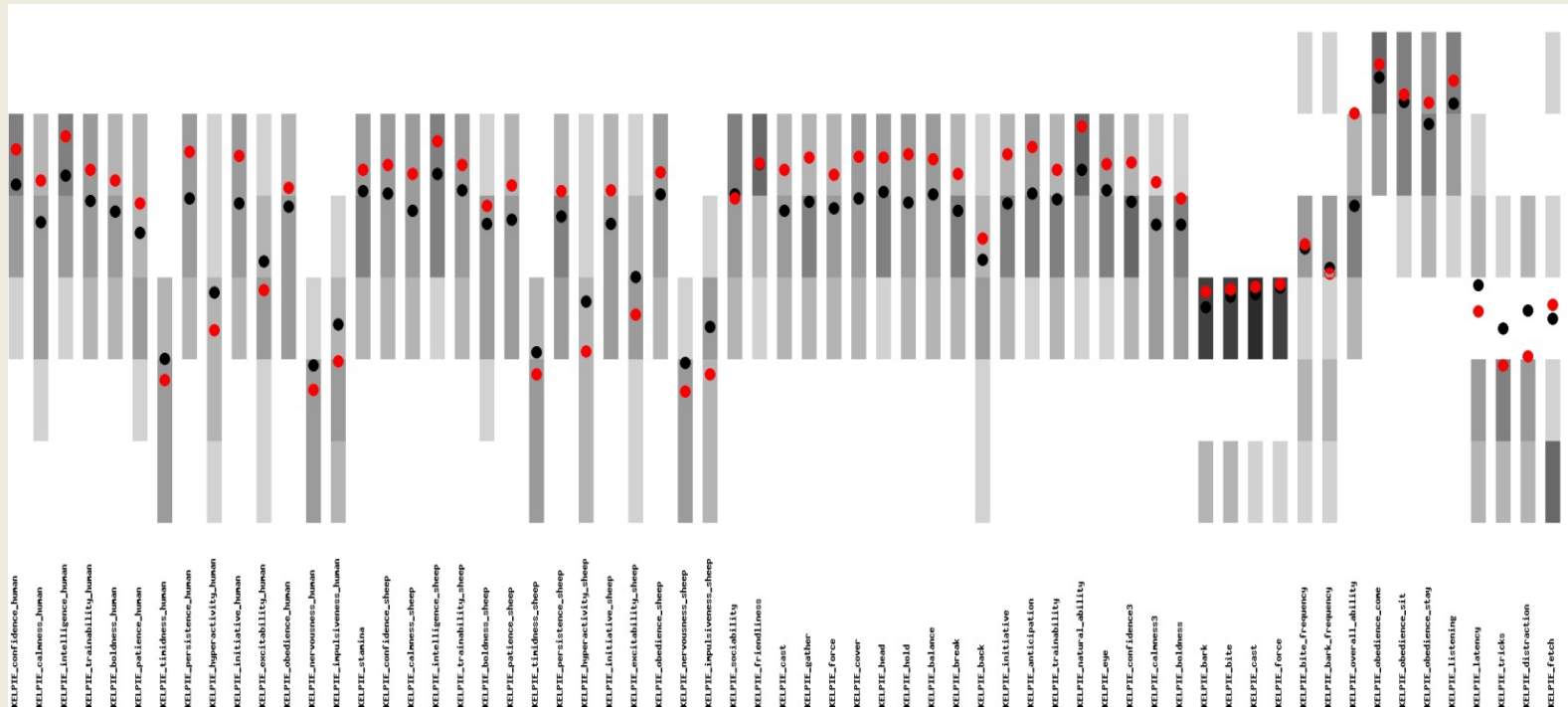


# Livestock working (Herding) Dog Assessment Form

- Targeted owner-driven phenotyping of 62 traits
  - As of March 2017 359 participants
- DNA samples collected from phenotyped dogs
  - As of March 2017 we have collected DNA from and genotyped 210 Kelpies we have samples from a further 100 Kelpies
  - We have whole genome sequence from 12 Kelpies including some that are competitive and successful trialling dogs
- Pedigree information and registration information collected
  - The entire Working Kelpie Council pedigree database was provided in collaboration with project

# Giving back

- Profiling the average dog (●) within a breed
- Profiling elite working dogs (●) within a breed



# Giving back

- If we choose the animals with the highest overall ability and compare these with the mean, we have a powerful insight into what breeders value
  - Scores relate to a Likert scale
  - These small computationally derived charts show scores in the range of 1-5 or 1-6 (most/highest at the top)
  - Darker hues describe a higher proportion of dogs at that score
- Elite dogs
- Mean



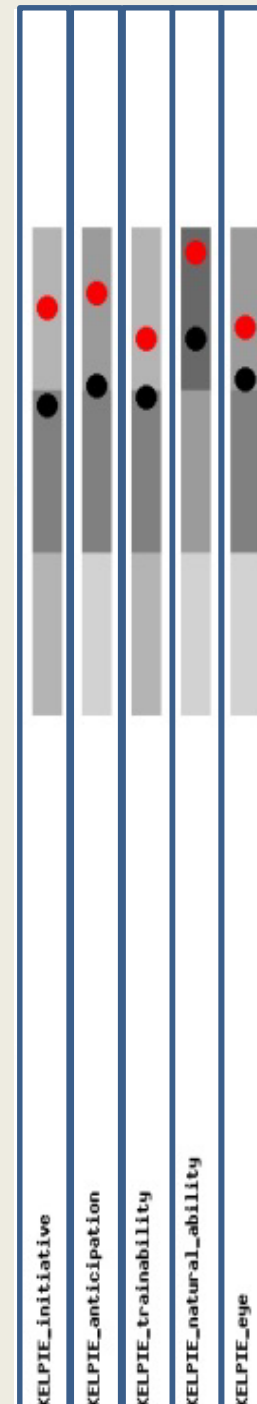
# Giving back

Elite Kelpies (●) have more than average (●):

- Initiative
- Anticipation
- Trainability
- Natural ability
- Eye

● Elite dogs

● Mean

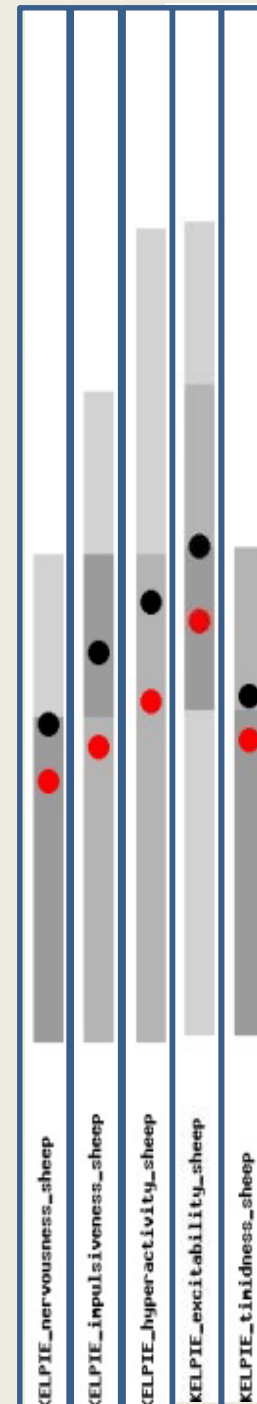


# Giving back

And elite Kelpies have less:

- Nervousness
- Impulsiveness
- Hyperactivity
- Excitability
- Timidness

● Elite dogs  
● Mean



# Giving back

Kelpie breeders are readily able to fix:

- Bark
- Bite
- Cast
- Force

...although different work types have different requirements

e.g. dogs that work on trucks and in yards should bark and bite versus those that work in the field should not



● Elite dogs

● Mean

# Giving back

...and nobody judging a kelpie as an elite worker cares too much about:

- Sociability
- Friendliness
- Come
- Sit
- Stay

● Elite dogs

● Mean

...since most kelpies are in the appropriate zone for those things



# Next steps:

- Add Functionality:
  - Identify the best breeders
  - Provide a means for users to find breeders producing the kind of dogs they need for their work type
  - Map traits
  - Generate Estimated Breeding Values
  - Overcome data curation challenges!



# Farm Dog Project Summary

- This is a long-term project and we are still collecting data
- We have clearly identified traits that describe elite working dogs
- We are actively collecting DNA and pedigrees for genetic and genomic analysis







# Elite Retrievers



- Comparison of elite field trial Labrador retrievers with other Labradors
- Can we gain insight into the genetic characteristics that underpin elite retrieving?



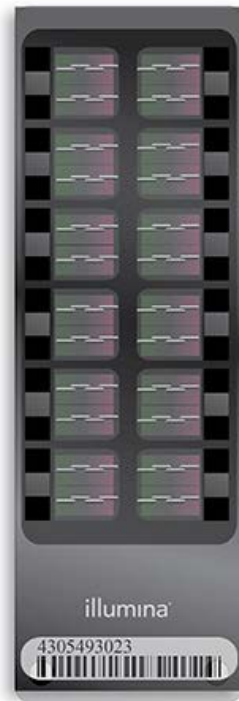
# Background

- Dogs with elite retrieving ability in Australian trials are admixed USA Labrador bloodlines
  - Very high drive
  - Very high energy
  - Lean
  - Muscular (which may equate with low subcutaneous fat)
  - Skeletal differences
    - Taller/Finer than English/Show type



# Data (population screening)

- High density and Ultra-high density DNA genotyping arrays (Illumina Canine HD and UHD)





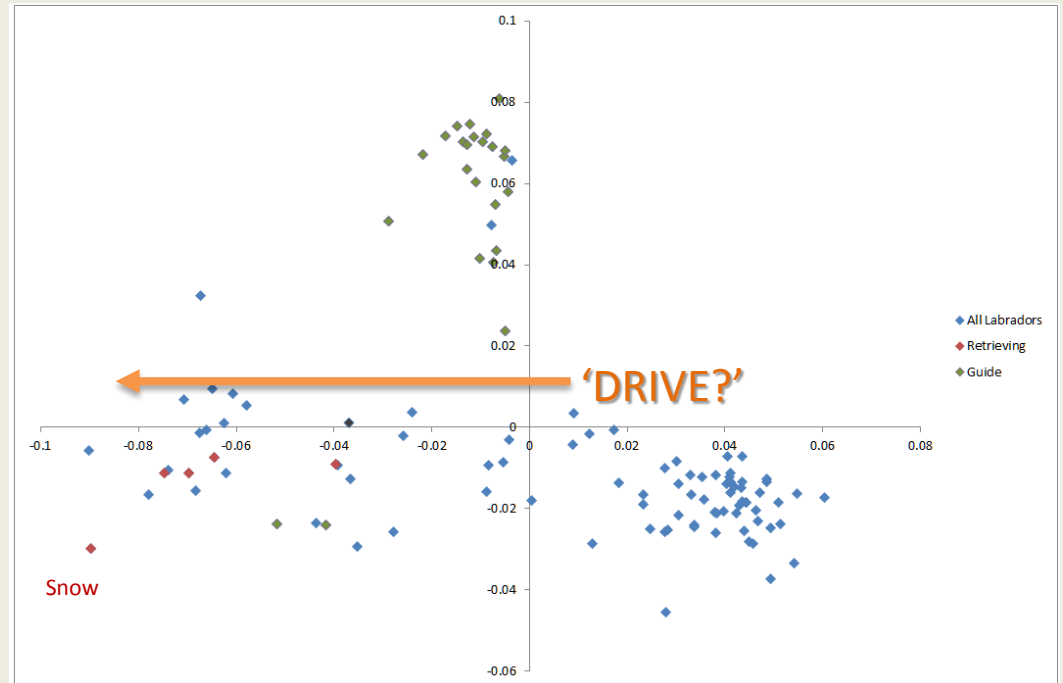
# High resolution data (mutation level)

- Whole genome sequencing (Illumina HiSeq2K and Illumina X10)



# Samples

- Elite trialling families ascertained by successes in National and State trials
- Admixed population
  - (USA retrieving trial champion semen imports by Australian working dog lines)



“Extremely high drive. No quit. Athletic. Outstanding work ethic. Missy keeps everyone on their toes!”

“Snow has a high work ethic, loves training, great memory, loves water, excellent fluid movement, hunt all day, sociable.”



# Admixture mapping

- Identifies regions according to population of origin
- Using known pedigree this can include heterozygous regions
- Hypothesis that USA-type affords skill





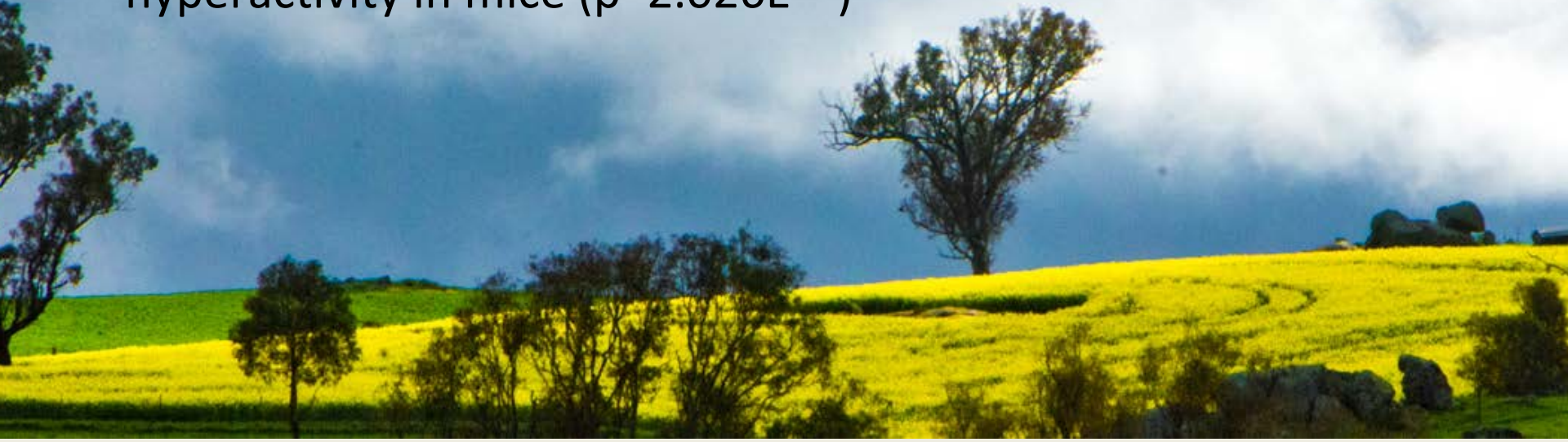
# Admixture mapping

- Admixture of individuals in our study is known by their pedigree
- We identify variants that are enriched in dogs with USA type ancestry
- We identify variants with homozygosity/heterozygosity according to pedigree expectation for individuals
- Probabilities are typically stronger than those from association



# Preliminary signals from admixture analysis

- On chromosome 20 in a region where there are several genes with a strong influence on adiposity ( $p=4.04E^{-25}$ )
  - Haplotype is shared with whippet and boxer
- A signal on chromosome 9 near a gene that is associated with hyperactivity in mice ( $p=2.026E^{-22}$ )





# Association mapping



- Cuts the study cohort down to include only retrievers of USA “type” by the MDS plot
- Looks at characteristics enriched in the retrieving elite dogs
- Cut down cohort only includes 25 USA-type dogs so p-values are more modest

# Association mapping



- Mechanosensitivity

$$(p_{\text{raw}} = 1.502\text{e-}08) | (p_{\text{genome}} = 0.03)$$

The haplotype includes a coding variant predicted to be damaging to the protein in the underlying gene

- Stress response

$$(p_{\text{raw}} = 2.772\text{e-}08) | (p_{\text{genome}} = 0.09)$$

In particular there is a gene at this locus that shares a stress response pathway with one identified in the admixture analysis.



# Association mapping

Mice that have been bred for different levels of acoustic startle and depression have different expression of the associated gene.



JOURNAL OF MOLECULAR NEUROSCIENCE

► [springer.com](http://springer.com)

*J Mol Neurosci.* 2013 May; 50(1): 33–57.

PMCID: PMC3622021

Published online 2012 Jul 27. doi: [10.1007/s12031-012-9850-1](https://doi.org/10.1007/s12031-012-9850-1)

## Effects of Chronic Stress on Prefrontal Cortex Transcriptome in Mice Displaying Different Genetic Backgrounds

Pawel Lisowski,  Marek Wieczorek, Joanna Goscik, Grzegorz R. Juszczak, Adrian M. Stankiewicz, Lech Zwierzchowski, and Artur H. Swiergiel



# Retrieving summary

- Variants uncovered depend on the hypothesis
- Admixture (USA-type superiority) compares USA-ancestral alleles with Australian derived
- Association looks for elite family specific differences relative to other USA-type dogs
- Work is ongoing/preliminary

# Retrieving summary

- Admixture mapping reveals regions associated with:
  - Adiposity
  - Stress response
  - Olfactory axon guidance
- Association (analysis restricted to dogs of USA-type) reveals regions associated with:
  - Mechanosensitivity
  - Stress sensitivity including acoustic startle (one gene in the same family as a gene in the admixture analysis)

# Retriever project summary

- We are expanding the families to be sampled
- We are excited by the findings thus far
- We are mutation screening in the implicated regions and will validate these in a wider cohort before trying to publish



# Please vote for our international competition entry to sequence the Australian Dingo

<http://www.pacb.com/smrt-science/smrt-grant/pag2017/dancing-dingos/>

The screenshot shows the PacBio website on an iPad. The top navigation bar includes links for Blog, Resource Library, DevNet, Customer Login, Investors, and Contact Us. The main navigation menu features PRODUCTS + SERVICES, RESEARCH FOCUS, APPLICATIONS, **SMRT SCIENCE** (highlighted), SUPPORT, and COMPANY. The header image displays 'DANCING DINGOES' in large text next to an owl. A sidebar on the left lists categories: SMRT SEQUENCING, SMRT RESOURCES, and SMRT GRANTS. Under SMRT GRANTS, the '2017 Plant and Animal SMRT Grant Program' is expanded, showing 'Dancing Dingoes' as a selected entry. The main content area for 'Dancing Dingoes' identifies it as a 2017 SMRT Grant Finalist and lists the Principal Investigators: Professor Bill Ballard and Professor Claire Wade, both from the University of New South Wales, Sydney, Australia. A photograph of a young dingo pup is shown. The 'About This Project' section describes the goal to sequence the genome of a pure desert dingo named Sandy, who was rescued from a harsh environment. A 'VOTE NOW' button with a right arrow is located at the bottom right.

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## DANCING DINGOES

SMRT SEQUENCING

SMRT RESOURCES


SMRT GRANTS

- 2017 Cancer SMRT Grant Program
- 2017 Plant and Animal SMRT Grant Program
  - Dancing Dingoes
  - Explosive Beetle
  - Pink Pigeon
  - Solar-powered Slug
  - Tamara Dingo

2017 SMRT Grant Finalist

### DANCING WITH DINGOES

Principal Investigators:  
**Professor Bill Ballard**, University of New South Wales, Sydney, Australia  
**Professor Claire Wade**, University of Sydney, Sydney, Australia



**About This Project**

We propose to sequence the genome of a pure desert dingo called Sandy. Sandy and her two siblings were just three weeks old when they were inadvertently stumbled upon in the harsh central Australian desert not far from the Strzelecki Track. They were dehydrated, could barely walk, and were close to death. Fortunately, the three amigos were saved and raised under idyllic surrounds by Barry Eggleton – a man who has now devoted his life to them.

**VOTE NOW**



# Acknowledgements

**Dog community**

**University of Sydney:**

**FARM DOG PROJECT:**

**Prof Paul McGreevy**

**Liz Arnott**

**Jonathan Early**

**Lisa Mascord**

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**Diane van Rooy**

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**Retrieving Photos: Al Dodge**



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