



Does including genomic information improve breeding values for production and health traits?

K Kaseja, E Smith, N Lambe, J Conington

Showcasing Smarter Sheep Science

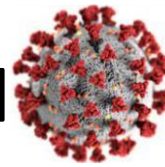
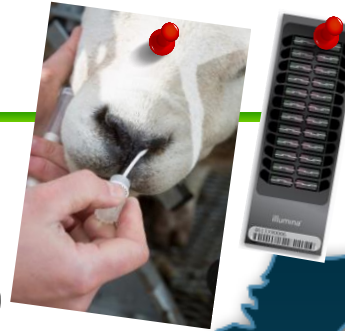
7th December 2022



This project has received funding from the European Union's Horizon 2020 research and innovation programme under the Grant Agreement n°772787

Available data

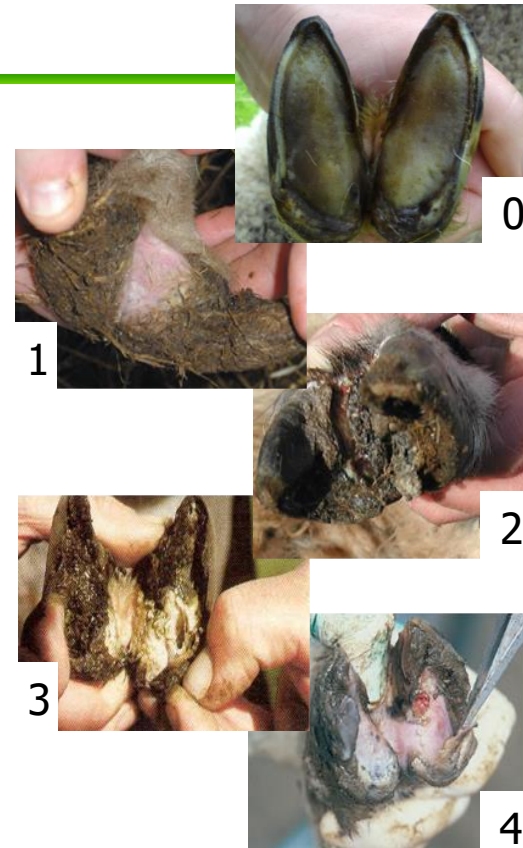
- Pre-existing data
 - Genotyped population, $n > 8,000$
 - Footrot phenotypes, $n = 8,572$
 - Mastitis phenotypes, $n = 6,676$
- Year 1: data collection [2019]
 - Footrot phenotypes, $n = 426$
 - Mastitis phenotypes, $n = 473$
- Year 2: no data collection [2020]
- Year 3: data collection [2021]
 - Footrot phenotypes, $n = 762$
 - Mastitis phenotypes, $n = 784$



Health traits scoring

- Footrot

- Each hoof scored: range 0 to 4
- All four scores summed together: range 0 to 16



- Mastitis – as proxy trait: California Mastitis Test (CMT)

- Milk sample from each side of the udder scored: range 0 to 4
- Two scores from two sides summed together: range 0 to 8



Is there potential for improvement?

N animals with phenotype

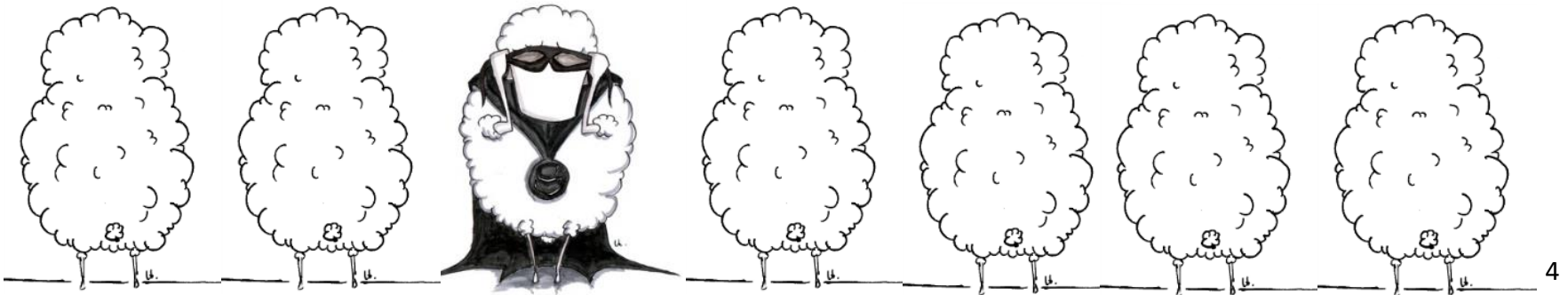
Footrot	6,216
California Mastitis Test	3,346

heritability

Footrot	0.12 (0.02)
California Mastitis Test	0.07 (0.03)

a measure of how well differences in genes account for differences in particular traits

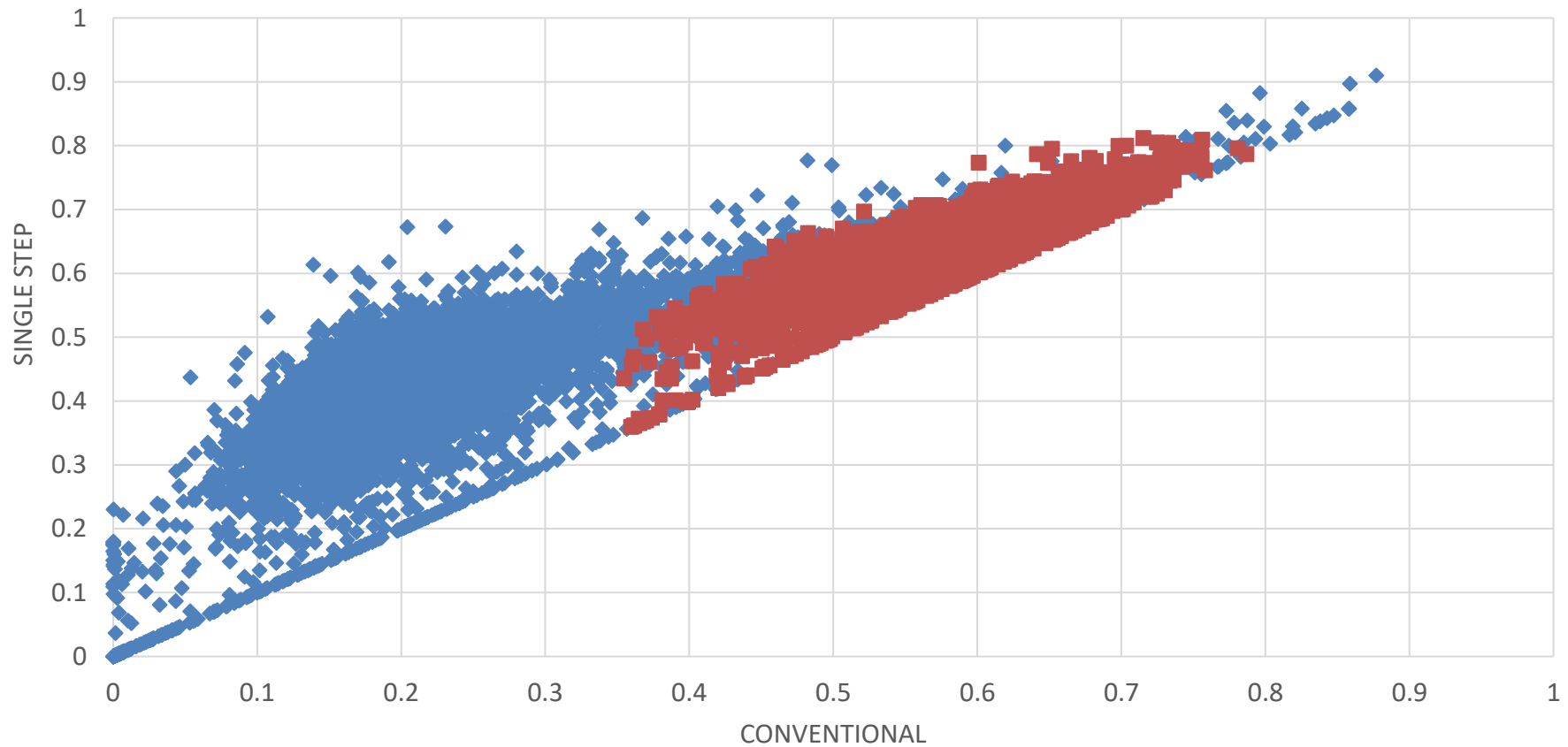
close to zero indicates that almost all the variability in a trait is due to environmental factors, with **very little influence** from genes





FOOTROT

◆ no phenotype ■ phenotype available

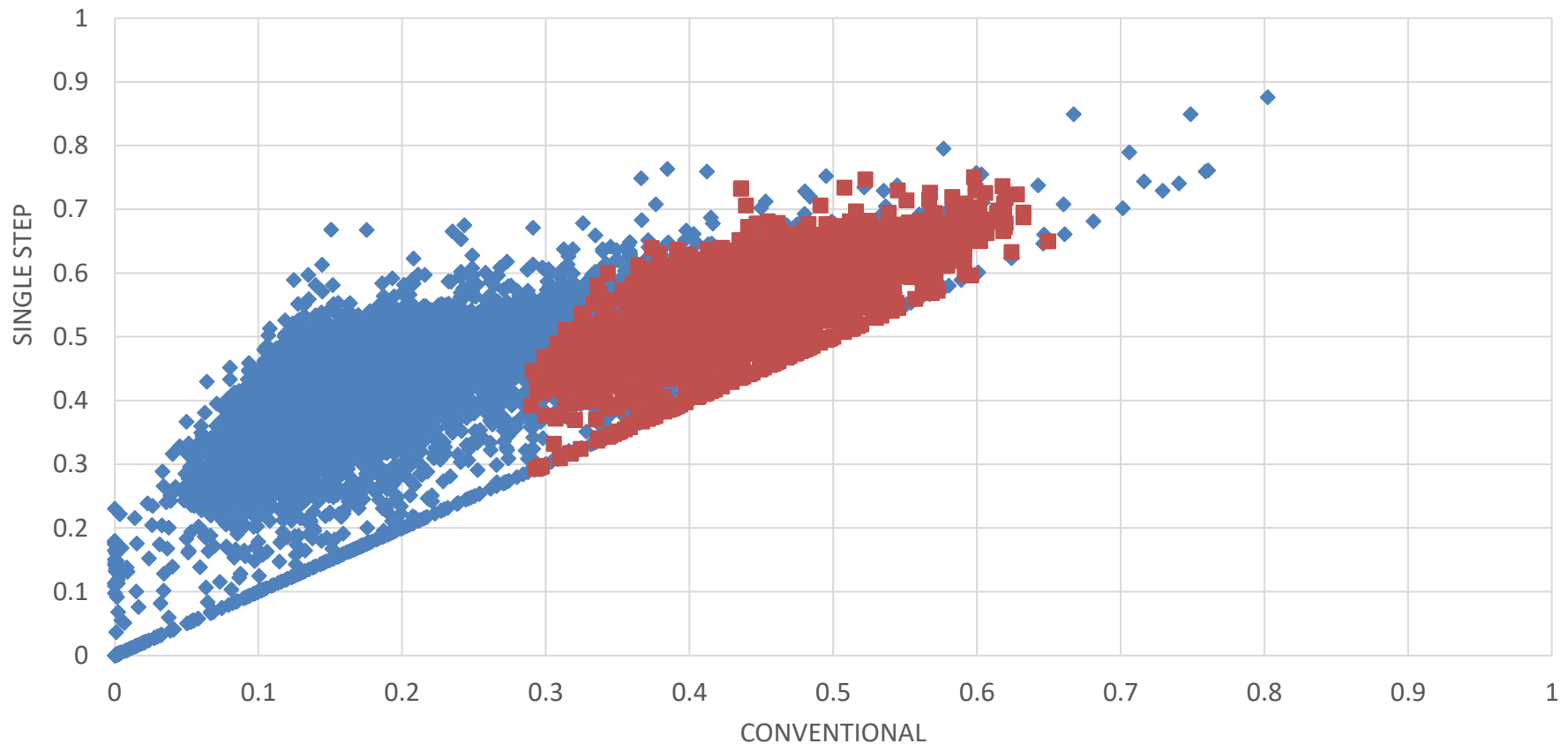


genotyped = 9,391

phenotyped = 4,506

CALIFORNIA MASTITIS TEST

◆ no phenotype ■ phenotype available

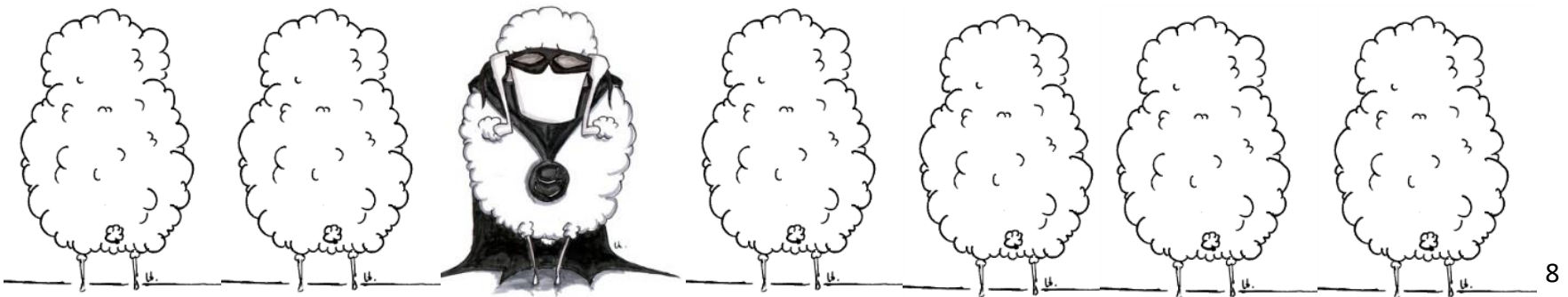


genotyped = 9,391

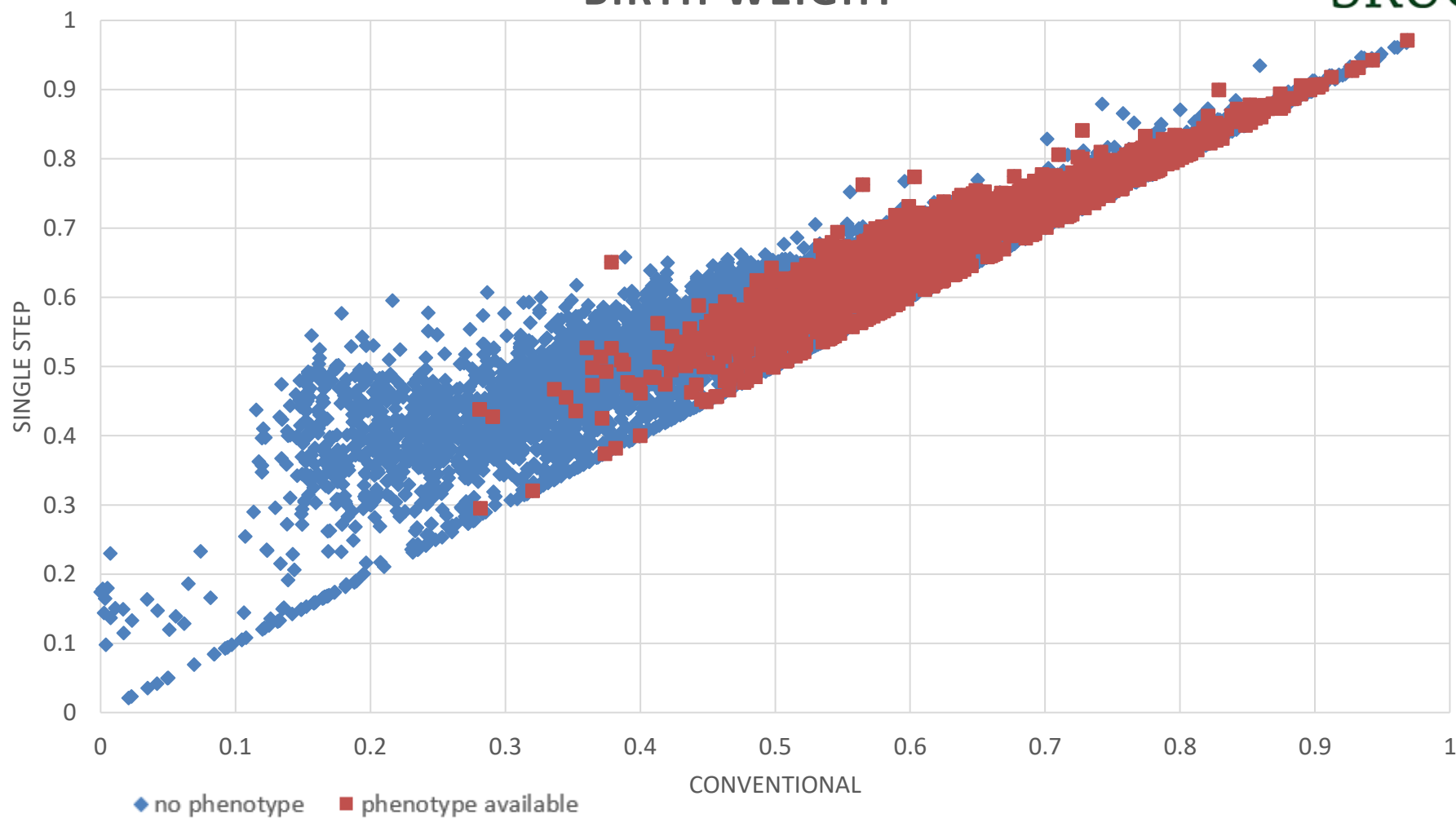
phenotyped = 3,434 (females only)

How about production traits?

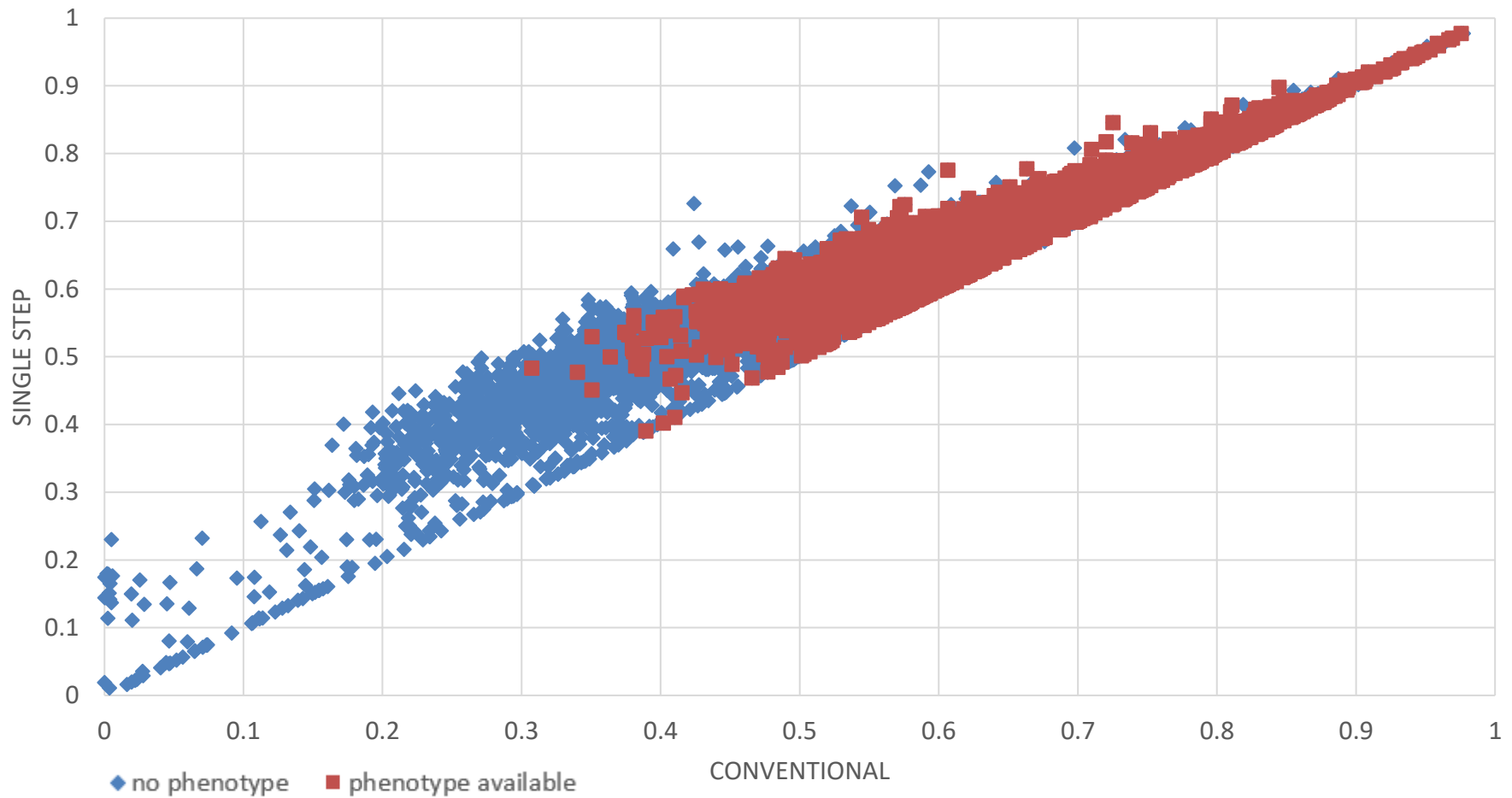
	N animals with phenotype	heritability
birth weight	33,857	0.10 (0.01)
weaning weight	72,845	0.09 (0.01)
scan weight	70,492	0.33 (0.01)
muscle depth	67,865	0.36 (0.01)
fat depth	67,867	0.36 (0.01)



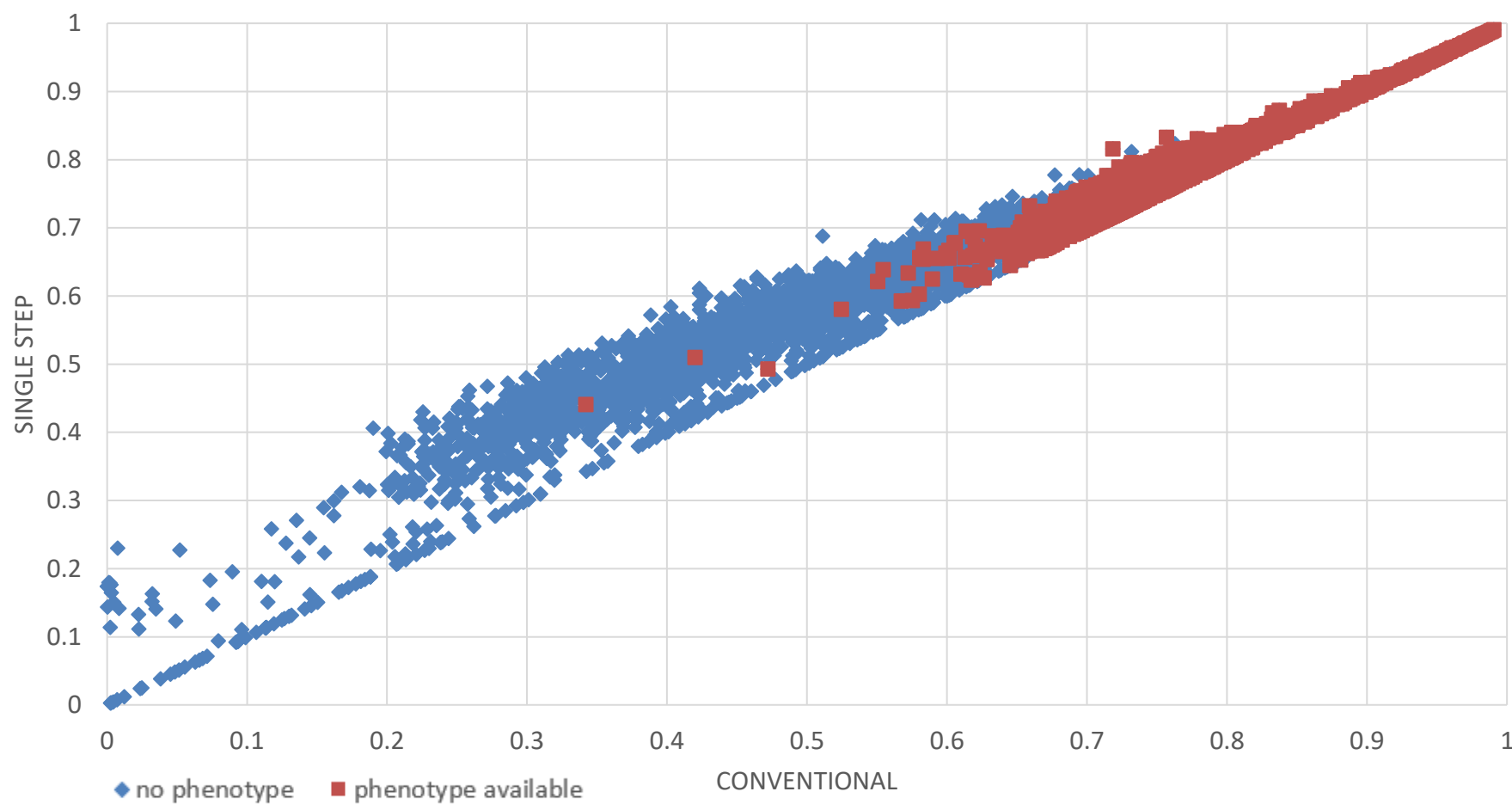
BIRTH WEIGHT



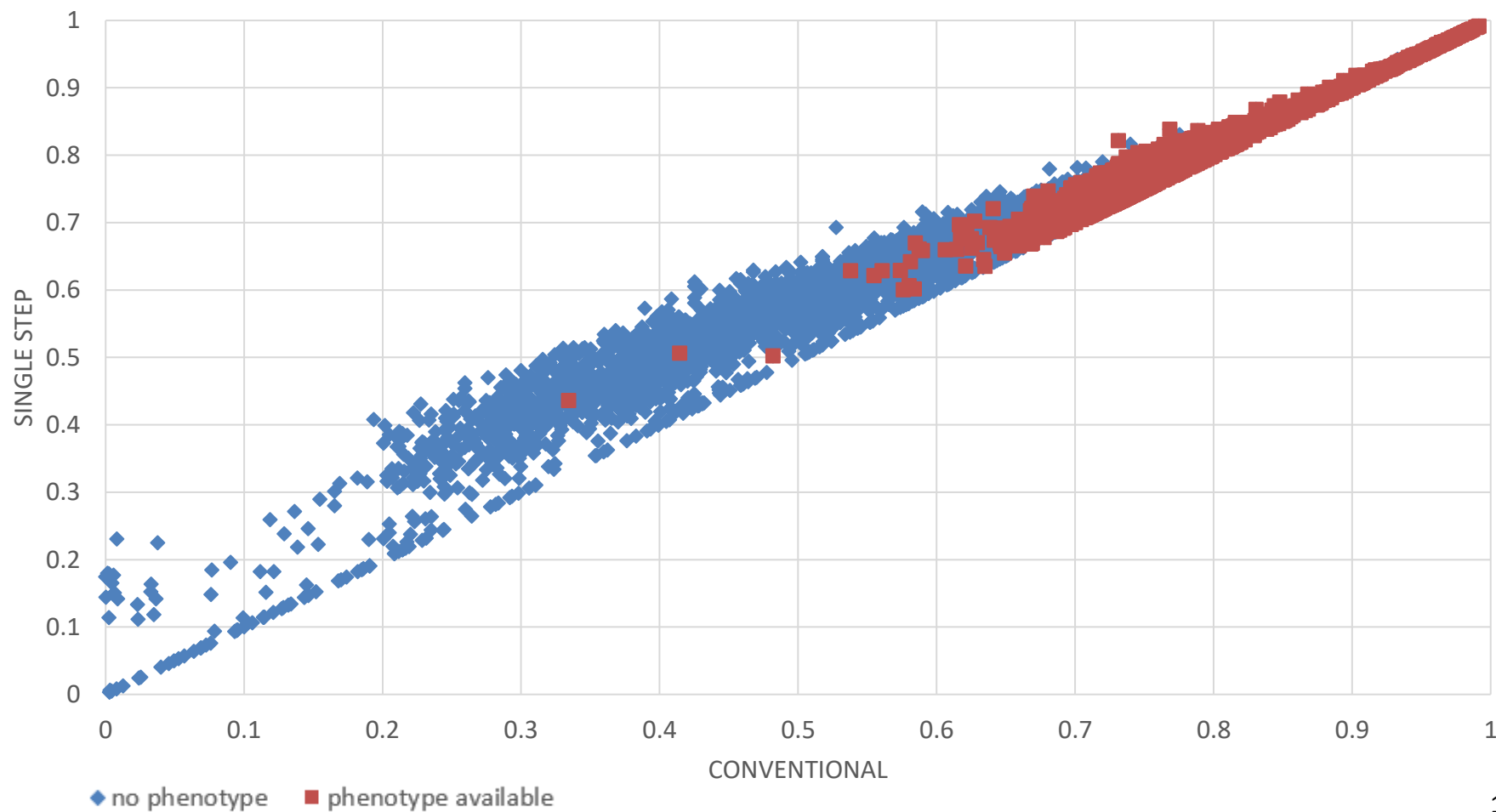
EIGHT WEEK WEIGHT



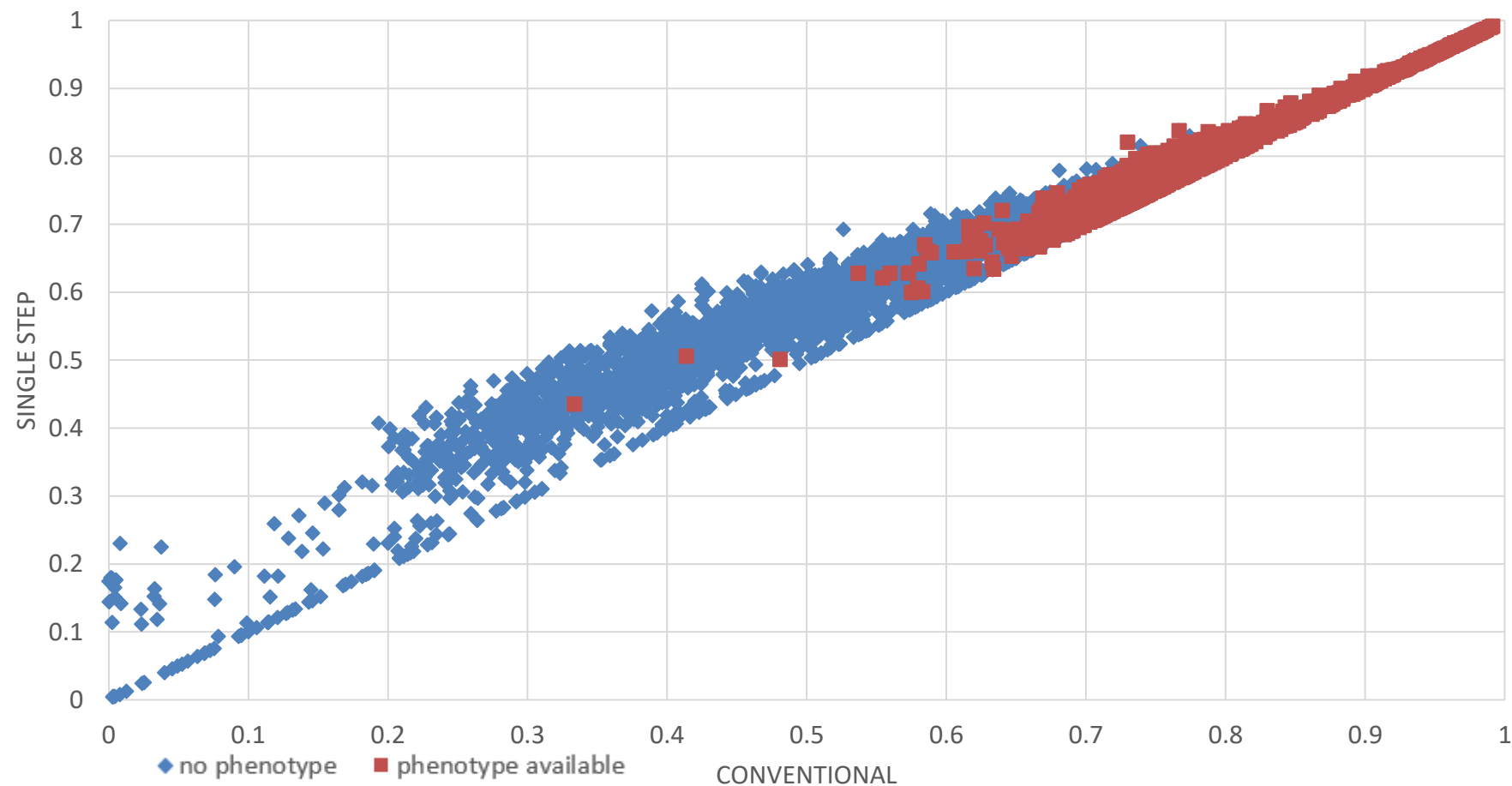
SCAN WEIGHT



MUSCLE DEPTH



FAT DEPTH



Does including genomic information improve breeding values for production and health traits?



Animals without phenotypic data gain the most in EBV accuracy when genotypic information is included

Both phenotyped and un-phenotyped can gain accuracy, hence their estimated breeding values will be closer to real breeding values

SMARTER PARTNERS



Thank you for your attention

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