N PROG

heifer

2nd mating

1st calving

Research EBV

LANCEFIELD S MCNEIL

 BREED:
 Brahman

 SOCIETY ID:
 LAS5339/1M

 DOB:
 2015 / 12 / 20

Sire: LANCEFIELD S DUTTON 4666/1 (PS)

Dam: LANCEFIELD S CHEEKY 5052/1 (PS)

DNA Case#: UQ660105
Project use: Natural Service

REPRONOMICS PROGENY

Cohort	#11	#12	#13	#14	#15	#16	#17	#18	#19	#20	TOTAL
N Brian Pastures									5	16	21
N Spyglass											0

TRAIT

Heifer age at puberty

Lactation anoestrous interval

Days to calving 1st mating 2nd mating **Body Condition Score** heifer <10 1st mating Repronomics research EBVs from the 2nd mating Hip Height heifer <10 April 2021 evaluation identified 1st mating the sire's EBV to be above average, 2nd mating average or below average compared Ultrasound EMA 1st mating with other sires with 10 or progeny 2nd mating for the trait. Each category contains Ultrasound P8 Fat heifer <10 approximately a third of the sires (from 1st mating all breeds), i.e. an above average sire 2nd mating Ultrasound Rib Fat 1st mating is in the top 30% of Repronomics sires 2nd mating recorded for that trait. ** indicates Live weight 1st mating the sire is in the top/bottom 3 of 2nd mating Repronomics sires Coat length score weaning 20 Sleeker Naval size score 1st mating Cow mothering score 1st calving Udder size score 1st calving

PROGENY RECORDED WITH BREEDPLAN: 21 calves across 1 herds born 2018 to 2019

Teat size score

http://abri.une.edu.au/online/cgi-bin/i4.dll?1=22202F2F&2=2420&3=56&5=2B3C2B3C3A&6=5F5C255D232623&9=5F5F5B5Da3C3A&6=5F5C255D232623&9=5F5C255D232623&9=5F5C255D232623&9=5F5C255D232623&9=5F5F5C255D232623&9=5F5F5C255D232625&9=5F5F5C255D232625&9=5F5F5C255D232625&9=5F5F5F5B5D25A%25254A%25254A%25254A%25254A%25254A%25254A%252544A%252544A%252544A%252544A%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%2525444%254444%254444%25444%25444%254444%25444%254444%25444%25444%25444%25444%25444%25444%25444%25444%25444%25444%254

Disclaimer: The results contained in this sheet have been obtained as part of the MLA funded Repronomics project.

These results are expected to change as more data is collected, or as models of analyses are refined.

Therefore, at this stage this sheet should NOT be reproduced or published.