N PROG

Research EBV

CUNGELELLA FOUNDATION (P) D5

BREED: Droughtmaster SOCIETY ID: 6NN95104M DOB: 1994 / 11 / 4

Sire: GAYVIEW 828 (S) D4

Dam: CUNGELELLA 1/1 (P) D4

DNA Case#: UQ496454

Project use: Al

REPRONOMICS PROGENY

Cohort	#11	#12	#13	#14	#15	#16	#17	#18	#19	#20	TOTAL
N Brian Pastures							8				8
N Spyglass							4				4

TRAIT

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

PROGENY RECORDED WITH BREEDPLAN: 317 calves across 7 herds born 1998 to 2016 http://abri.une.edu.au/online/cgi-bin/i4.dll?1=2F3F2F3D&2=2420&5=2B3C2B3C3A&6=5D5A5B5B252526212D&9=5D5B5A58

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.