## NCC SPRINGBOK (IMP ET) (AI) (ET) (H)

BREED:	Brahman
SOCIETY ID:	NCC591M
DOB:	2006 / 2 / 4
Sire:	MASBRA MAS 98 34 (IMP SAF)
Dam:	KLEIN COR MISS TARENTAAL (IMP SA)
DNA Case#:	UQ496450
Project use:	AI

## **REPRONOMICS PROGENY**

Cohort	#11	#12	#13	#14	#15	#16	#17	#18	#19	#20	TOTAL
N Brian Pastures							6	2			8
N Spyglass							7				7

	TRAIT		N PROG	Research EBV
	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	13	Average
	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:** 78 calves across 8 herds born 2008 to 2017 http://abri.une.edu.au/online/cgi-bin/i4.dll?1=22202F2F&2=2420&3=56&5=2B3C2B3C3A&6=595B59265823&9=5D505E

**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.