RS — ALUMY CREEK PRIME QUARTER S014 SV — NKE21S014

DOB: 18/07/2021 **Registration Status:** HBR **Mating Type:** AI **Genetic Status:** AMFU,CAFU,DDFU,NHFU

CONNEALY CONSENSUS 7229 SV	S A V FINAL ANSWER 0035 #
V A R GENERATION 2100 PV	S A V PIONEER 7301 #
SANDPOINT BLACKBIRD 8809 #	S A V BLACKBIRD 5297 #
Sire: USA18232879 EF PRIME QUARTER 5369 PV	**Dam:** NKEH68 ALUMY CREEK TRILOGY H68 #
G A R PROPHET SV	S A NEUTRON 377 #
EF RITA 3422 #	ALUMY CREEK TRILOGY E19 #
EF RITA 7328 #	ALUMY CREEK TRILOGY X10 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+2.3	+5.3	-7.3	+5.5	+67	+116	+137	+107	+0.21	+6.4	+13	+2.7
ACC	66%	57%	83%	82%	83%	81%	81%	78%	65%	67%	75%	44%
Perc	56	31	13	82	4	5	18	42	65	84	83	99

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+2.7	+13	+83	+7.1	+0.0	+0.9	+0.2	+1.6	+0.23	+1.12	+1.04	+1.16
ACC	79%	76%	71%	70%	70%	71%	62%	74%	61%	69%	69%	63%
Perc	30	82	15	42	50	30	59	69	51	92	67	86

Selection Indexes

$A	$A-L
$223	**$372**
33	38

Traits Observed: GL, BWT, 200WT, 400WT, 600WT, SC, Scan(EMA, Rib, Rump, IMF), Genomics

Statistics: Number of Herds: 1, Prog Analysed: 3, Genomic Prog: 3

RS — BALDRIDGE 38 SPECIAL PV — USA18229487

DOB: 13/01/2015 **Registration Status:** HBR **Mating Type:** Natural **Genetic Status:** AMF,CAF,DDF,NHF,MAF,OSF,RGF

BASIN FRANCHISE P142 #	SITZ UPWARD 307R SV
EF COMPLEMENT 8088 PV	STYLES UPGRADE J59 #
EF EVERELDA ENTENSE 6117 #	PLAINVIEW LASSIE 71B #
Sire: USA17082311 EF COMMANDO 1366 PV	**Dam:** USA17149410 BALDRIDGE ISABEL Y69 #
B/R AMBUSH 28 #	BALDRIDGE KABOOM K243 KCF #
RIVERBEND YOUNG LUCY W1470 #	BALDRIDGE ISABEL T935 #
RIVERBEND YOUNG LUCY T1080 #	BALDRIDGE ISABEL P4527 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+7.7	+5.7	-4.9	+2.6	+64	+111	+144	+109	+0.31	+8.2	+20	+2.6
ACC	94%	83%	99%	99%	98%	98%	98%	97%	90%	92%	97%	71%
Perc	10	27	44	22	7	11	10	40	36	53	27	25

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+2.6	+16	+79	+6.0	+1.4	-1.4	-0.4	+3.1	+0.15	+0.60	+0.76	+0.88
ACC	98%	99%	94%	92%	92%	92%	89%	92%	79%	99%	99%	96%
Perc	33	71	22	55	21	70	87	33	42	9	10	12

Selection Indexes

$A	$A-L
$253	**$426**
9	7

Traits Observed: Genomics

Statistics: Number of Herds: 137, Prog Analysed: 2476, Genomic Prog: 1647

RS — BALDRIDGE COMPASS C041 SV — USA18229488

DOB: 14/01/2015 **Registration Status:** HBR **Mating Type:** ET **Genetic Status:** AMF,CAF,DDF,NHF,MHF,OHF,OSF

BASIN FRANCHISE P142 #	SITZ UPWARD 307R SV
EF COMPLEMENT 8088 PV	STYLES UPGRADE J59 #
EF EVERELDA ENTENSE 6117 #	PLAINVIEW LASSIE 71B #
Sire: USA17082311 EF COMMANDO 1366 PV	**Dam:** USA17149410 BALDRIDGE ISABEL Y69 #
B/R AMBUSH 28 #	BALDRIDGE KABOOM K243 KCF #
RIVERBEND YOUNG LUCY W1470 #	BALDRIDGE ISABEL T935 #
RIVERBEND YOUNG LUCY T1080 #	BALDRIDGE ISABEL P4527 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+7.4	+5.8	-3.6	+3.0	+61	+107	+133	+85	+0.42	+3.4	+29	+1.8
ACC	93%	84%	99%	99%	98%	98%	98%	97%	92%	93%	97%	71%
Perc	12	26	65	29	14	17	24	77	14	99	2	64

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+1.8	+22	+70	+6.3	+0.6	-0.9	+0.2	+2.7	+0.37	+0.64	+0.64	+0.80
ACC	98%	97%	94%	93%	93%	93%	90%	93%	82%	98%	98%	96%
Perc	63	44	47	52	36	62	59	42	66	13	2	4

Selection Indexes

$A	$A-L
$253	**$400**
9	17

Traits Observed: Genomics

Statistics: Number of Herds: 95, Prog Analysed: 1300, Genomic Prog: 860

RS — CONNEALY BIG VALLEY ^{PV} — USA19249322

DOB: 30/01/2018 **Registration Status:** HBR **Mating Type:** Natural **Genetic Status:** AMF,CAF,DDF,NHF,DWF,MAF,MHF,OHF,OSF,

	Sire:	Dam:
	MOGCK BULLSEYE ^{PV}	CONNEALY CONFIDENCE 0100 [#]
	HOOVER NO DOUBT ^{PV}	CONNEALY GREELEY [#]
	MISS BLACKCAP ELLSTON J2 [#]	PRECIOUS OF CONANGA 0484 [#]

Sire: USA18533906 CONNEALY DRY VALLEY ^{PV} **Dam:** USA17921087 BUNTY LAY OF CONANGA 4930 [#]

	SUMMITCREST COMPLETE 1P55 [#]	G A R PREDESTINED [#]
	BECCA OF CONANGA 1617 [#]	BUNTY LANA OF CONANGA 4102 [#]
	BRISHA OF CONANGA 746 1237 [#]	BUNTY LEAH OF CONANGA 551 [#]

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	-6.2	+3.8	-3.1	+7.1	+71	+128	+153	+135	+0.14	+7.5	+11	+2.5
ACC	66%	52%	91%	88%	87%	85%	84%	82%	61%	62%	81%	38%
Perc	95	48	73	97	2	1	5	10	81	67	91	82

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+2.5	+29	+93	+6.6	-0.7	-3.3	+0.5	+1.8	+0.01	+1.10	+1.02	+0.80
ACC	83%	72%	79%	78%	74%	71%	67%	80%	59%	95%	94%	57%
Perc	37	21	4	48	67	92	40	64	27	91	63	4

Selection Indexes

$A	$A-L
$223	$385
33	27

Traits Observed: Genomics

Statistics: Number of Herds: 4, Prog Analysed: 38, Genomic Prog: 19

RS — EF PRIME QUARTER 5369 ^{PV} — USA18232879

DOB: 26/09/2015 **Registration Status:** HBR **Mating Type:** Natural **Genetic Status:** AMF,CAF,DDF,NHF,MHF,OHF,OSF

	Sire:	Dam:
	CONNEALY CONSENSUS [#]	C R A BEXTOR 872 5205 608 [#]
	CONNEALY CONSENSUS 7229 ^{SV}	G A R PROPHET ^{SV}
	BLUE LILLY OF CONANGA 16 [#]	G A R OBJECTIVE 1885 [#]

Sire: USA17171587 V A R GENERATION 2100 ^{PV} **Dam:** USA17756888 EF RITA 3422 [#]

	CONNEALY ONWARD [#]	G A R PREDESTINED [#]
	SANDPOINT BLACKBIRD 8809 [#]	EF RITA 7328 [#]
	RIVERBEND BLACKBIRD 4301 [#]	A PLUS RITA 5H11 8009 [#]

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+2.3	+8.0	-7.3	+3.2	+59	+95	+119	+71	+0.09	+5.5	+11	+2.1
ACC	81%	70%	96%	96%	95%	95%	94%	90%	77%	78%	87%	59%
Perc	56	8	13	33	19	44	56	91	89	92	89	79

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+2.1	+36	+72	+6.1	+1.1	+0.1	-0.2	+2.8	+0.40	+1.00	+0.82	+1.00
ACC	93%	89%	86%	85%	85%	84%	79%	86%	71%	85%	85%	72%
Perc	52	7	42	54	26	44	80	40	69	79	17	42

Selection Indexes

$A	$A-L
$234	$361
22	47

Traits Observed: Genomics

Statistics: Number of Herds: 13, Prog Analysed: 154, Genomic Prog: 82

RS — FERGUSON TRAILBLAZER 239E ^{SV} — USA18996007

DOB: 18/02/2017 **Registration Status:** HBR **Mating Type:** Natural **Genetic Status:** AMF,CAF,DDF,NHF,MHF,OHF,OSF

	Sire:	Dam:
	MYTTY IN FOCUS [#]	O C C EMBLAZON 854E [#]
	A A R TEN X 7008 S A ^{SV}	LD EMBLAZON 999 ^{PV}
	A A R LADY KELTON 5551 [#]	SH FOREVER LADY 3124 5118 [#]

Sire: USA17262835 V A R DISCOVERY 2240 ^{PV} **Dam:** USA17717153 MOLITOR999 BARBELLA 940-3012 [#]

	SITZ UPWARD 307R ^{SV}	S A V FINAL ANSWER 0035 [#]
	DEER VALLEY RITA 0308 [#]	MOLITOR FA BARBELLA 389-940 [#]
	G A R OBJECTIVE 2345 [#]	MOLITOR5321 BARBELLA9027 389 [#]

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+3.4	+7.3	-7.3	+3.1	+72	+131	+174	+158	+0.26	+6.3	+13	+2.2
ACC	80%	67%	98%	97%	95%	95%	95%	90%	71%	76%	85%	55%
Perc	46	13	13	31	1	1	1	2	51	85	79	13

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+2.2	+39	+97	+3.0	+1.9	+1.0	-1.3	+4.7	+0.42	+1.24	+1.14	+0.88
ACC	94%	88%	86%	86%	85%	83%	79%	86%	68%	87%	84%	67%
Perc	48	4	3	87	14	28	99	8	71	98	85	12

Selection Indexes

$A	$A-L
$275	$492
2	1

Traits Observed: Genomics

Statistics: Number of Herds: 35, Prog Analysed: 422, Genomic Prog: 286

RS   G A R PROACTIVE ^{SV}   USA18333424

DOB: 08/09/2015   Registration Status: **HBR**   Mating Type: **Natural**   Genetic Status: **AMF,CAF,DDF,NHF,DWF,MHF,OHF,OSF,RGF**

B A R EXT TRAVELER 205 #		BOYD NEW DAY 8005 #
C R A BEXTOR 872 5205 608 #		MCC DAYBREAK #
CRA LADY JAYE 608 498 S EASY #		MCC MISS FOCUS 134 #
Sire: USA16295688 G A R PROPHET ^{SV}		**Dam: USA17056864 G A R DAYBREAK 1521** #
S S OBJECTIVE T510 0T26 #		G A R NEW DESIGN 5050 #
G A R OBJECTIVE 1885 #		G A R 5050 NEW DESIGN 1039 #
G A R 1407 NEW DESIGN 2232 #		G A R OBJECTIVE 2345 #

Mid January 2025 TransTasman Angus Cattle Evaluation

Selection Indexes

	$A	$A-L
	$256	$432
	7	5

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+2.7	+7.5	-2.5	+2.8	+66	+111	+144	+121	+0.19	+7.4	+17	+1.6
ACC	80%	70%	96%	94%	94%	94%	93%	92%	75%	86%	89%	60%
Perc	52	11	80	25	5	10	11	23	70	68	54	10

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+1.6	+13	+81	+7.4	-3.5	-5.8	+1.1	+1.6	-0.02	+1.06	+1.16	+1.12
ACC	91%	89%	87%	86%	85%	84%	80%	87%	72%	93%	92%	70%
Perc	71	82	19	38	98	99	12	69	24	87	88	78

Traits Observed: Genomics

Statistics: Number of Herds: 9, Prog Analysed: 86, Genomic Prog: 77

RS   HOFFMAN THEDFORD ^{PV}   USA19820180

DOB: 28/01/2020   Registration Status: **HBR**   Mating Type: **Natural**   Genetic Status: **AMF,CAF,DDF,NHF,DWF,MAF,MHF,OHF,OSF,**

CONNEALY CONSENSUS 7229 ^{SV}		MOGCK SURE SHOT #
CONNEALY BLACK GRANITE #		KG SOLUTION 0018 #
EURA ELGA OF CONANGA 9109 #		KG RITO LADY 8724 #
Sire: USA18389838 BAR R JET BLACK 5063 ^{PV}		**Dam: USA17651108 HA RITO LADY 3839** #
SITZ UPWARD 307R ^{SV}		HA PROGRAM 5652 #
BAR R IRIS ANITA 0113 #		HA RITO LADY 0622 #
BAR R ANITA 7081 #		HA RITO LADY 8395 #

Mid January 2025 TransTasman Angus Cattle Evaluation

Selection Indexes

	$A	$A-L
	$255	$417
	8	9

	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+7.1	+7.5	-5.1	+2.2	+63	+111	+141	+96	+0.02	+8.3	+24	+3.4
ACC	70%	54%	91%	93%	88%	86%	85%	83%	62%	63%	80%	40%
Perc	14	11	41	16	10	10	13	61	95	51	8	49

	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+3.4	+24	+91	+10.0	-1.0	-3.2	+0.6	+1.9	-0.31	+1.14	+0.76	+0.86
ACC	83%	74%	80%	78%	75%	74%	67%	80%	61%	96%	95%	57%
Perc	14	37	6	15	73	91	34	62	7	94	10	9

Traits Observed: Genomics

Statistics: Number of Herds: 14, Prog Analysed: 65, Genomic Prog: 25

RS   K C F BENNETT EXPONENTIAL ^{PV}   USA19507801

DOB: 17/02/2019   Registration Status: **HBR**   Mating Type: **Natural**   Genetic Status: **AMF,CAF,DDF,NHF,DWF,MAF,MHF,OHF,OSF,**

SUMMITCREST COMPLETE 1P55 #		NICHOLS EXTRA K205 #
KM BROKEN BOW 002 ^{PV}		K C F BENNETT SOUTHSIDE ^{PV}
SUMMITCREST PRINCESS 0P12 #		K C F MISS 208 S11 #
Sire: USA18658677 CASINO BOMBER N33 #		**Dam: USA17863237 K C F MISS SOUTHSIDE B226** #
DPL UPWARD L70 #		CONNEALY TRUSTMARK #
CASINO ANNIE K48 #		K C F MISS TRUSTMARK W236 #
CASINO ANNIE G64 #		K C F MISS 589 M359 #

Mid January 2025 TransTasman Angus Cattle Evaluation

Selection Indexes

	$A	$A-L
	$258	$437
	7	4

	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+4.2	+3.0	-3.5	+3.3	+75	+126	+152	+134	+0.24	+6.1	+19	+2.3
ACC	70%	55%	95%	94%	91%	91%	89%	85%	63%	64%	81%	43%
Perc	38	57	67	35	1	2	5	11	56	88	37	71

	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+2.3	+16	+99	+8.7	-4.3	-8.3	+1.4	+2.2	-0.45	+0.86	+0.82	+1.12
ACC	89%	85%	82%	81%	79%	77%	72%	83%	61%	98%	98%	80%
Perc	44	73	2	25	99	99	6	54	4	54	17	78

Traits Observed: Structure(Claw Set x 1, Foot Angle x 1), Genomics

Statistics: Number of Herds: 9, Prog Analysed: 82, Genomic Prog: 49

RS — K C F BENNETT THEROCK A473 PV — USA17849954

DOB: 08/10/2013 Registration Status: **HBR** Mating Type: **Natural** Genetic Status: **AMF,CAF,DDF,NHF**

	Sire	Dam
	NICHOLS EXTRA H6 #	RITO 1I2 OF 2536 RITO 6I6 #
	NICHOLS EXTRA K205 #	GAR-EGL PROTEGE #
	NICHOLS BLACK HEIRESS F346 #	L B 6807 ISABEL 339 #
Sire: USA16430862 K C F BENNETT SOUTHSIDE PV	Dam: USA16430913 K C F MISS PROTEGE W148 #
	BON VIEW NEW DESIGN 208 SV	C A FUTURE DIRECTION 5321 SV
	K C F MISS 208 S11 #	BREELA OF CONANGA 3991 #
	K C F MISS 338 P14 #	BREESHA OF CONANGA 1251 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	-2.0	+3.5	-8.4	+4.6	+62	+108	+134	+142	+0.22	+10.6	+15	+3.1
ACC	81%	68%	97%	96%	94%	94%	94%	91%	94%	95%	91%	52%
Perc	84	52	6	66	12	14	23	7	62	13	70	29

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+3.1	+17	+83	+2.6	-2.6	-5.1	-0.1	+2.7	+0.10	+1.34	+1.26	+1.08
ACC	91%	86%	87%	85%	85%	83%	78%	86%	67%	96%	96%	41%
Perc	19	67	15	89	94	99	75	42	36	99	96	67

Selection Indexes

$A	$A-L
$185	$358
75	50

Traits Observed: Genomics

Statistics: Number of Herds: 19, Prog Analysed: 198, Genomic Prog: 107

RS — LD CAPITALIST 316 PV — USA17666102

DOB: 26/01/2013 Registration Status: **HBR** Mating Type: **Natural** Genetic Status: **AMF,CAF,DDF,NHF,DWF,MAF,MHF,OHF,OSF,**

	Sire	Dam
	SITZ TRAVELER 8180 #	G A R PRECISION 1680 #
	S A V FINAL ANSWER 0035 #	C A FUTURE DIRECTION 5321 SV
	S A V EMULOUS 8145 #	C A MISS POWER FIX 308 #
Sire: USA16752262 CONNEALY CAPITALIST 028 #	Dam: USA14407230 LD DIXIE ERICA 2053 #
	C R A BEXTOR 872 5205 608 #	LD ROYCE ONAROLL 810 #
	PRIDES PITA OF CONANGA 8821 #	LD DIXIE ERICA OAR 0853 #
	PRIDES TRAV OF CONANGA 6499 #	DIXIE ERICA OF R R 8553 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+9.4	+9.8	-3.5	+2.0	+50	+89	+107	+84	+0.35	+6.3	+14	+1.0
ACC	98%	92%	99%	99%	99%	99%	99%	98%	96%	97%	99%	88%
Perc	3	2	67	14	58	65	79	79	27	86	77	77

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+1.0	+8	+76	+8.3	+1.1	+1.8	+0.2	+1.3	+0.48	+0.90	+0.88	+0.88
ACC	99%	99%	98%	97%	97%	97%	96%	96%	90%	99%	99%	98%
Perc	87	93	30	29	26	18	59	76	76	62	29	12

Selection Indexes

$A	$A-L
$206	$350
53	56

Traits Observed: Genomics

Statistics: Number of Herds: 227, Prog Analysed: 3737, Genomic Prog: 1831

RS — LT REVERED SV — USA19548516

DOB: 17/02/2019 Registration Status: **HBR** Mating Type: **Natural** Genetic Status: **AMF,CAF,DDF,NHF,DWF,MAF,MHF,OHF,OSF,**

	Sire	Dam
	BASIN RAINMAKER P175 #	VISION UNANIMOUS 1418 PV
	BASIN RAINMAKER 2704 #	S FOUNDATION 514 PV
	BASIN ERICA 7520 BV #	S LADY ANN 8384 #
Sire: USA17913751 BASIN RAINMAKER 4404 PV	Dam: USA18953479 LT ASHLEY 7078 #
	BASIN PAYWEIGHT 107S #	K C F BENNETT TOTAL #
	BASIN JOY 1036 #	LT ASHLEY 8263 #
	BASIN JOY 566T #	LT FOREVER LADY 6124 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+8.2	+10.4	-6.5	+2.9	+60	+114	+133	+98	+0.05	+5.9	+23	+2.9
ACC	73%	55%	96%	96%	92%	92%	91%	86%	61%	62%	81%	45%
Perc	8	1	21	27	15	7	25	57	93	90	11	13

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+2.9	+42	+84	+5.3	+1.2	+0.1	+0.5	+1.5	+0.21	+0.88	+0.82	+1.06
ACC	89%	74%	82%	80%	79%	78%	72%	82%	62%	89%	89%	61%
Perc	24	3	14	64	24	44	40	72	48	58	17	61

Selection Indexes

$A	$A-L
$268	$444
4	3

Traits Observed: Genomics

Statistics: Number of Herds: 16, Prog Analysed: 163, Genomic Prog: 74

RS — MOGCK ENTICE ^{SV} — USA18952921

DOB: 31/01/2017 **Registration Status:** HBR **Mating Type:** Natural **Genetic Status:** AMF,CAF,DDF,NHF,OHF,OSF

SYDGEN GOOGOL #
SYDGEN EXCEED 3223 PV
SYDGEN FOREVER LADY 1255 #
Sire: USA18170041 SYDGEN ENHANCE ^{SV}
SYDGEN LIBERTY GA 8627 #
SYDGEN RITA 2618 #
FOX RUN RITA 9308 #

MOGCK SURE SHOT #
MOGCK SURE SHOT 253 #
MOGCK MISS 61 #
Dam: USA18334720 MOGCK ERICA 2255 #
CONNEALY 5050 611B #
MOGCK ERICA 2162 #
MOGCK ERICA 08 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+1.6	+2.5	-7.5	+5.3	+72	+133	+178	+160	+0.33	+8.7	+26	+5.1
ACC	85%	71%	98%	98%	96%	97%	96%	92%	77%	84%	89%	54%
Perc	62	62	12	79	1	1	1	2	31	43	5	54

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+5.1	+37	+97	+8.8	-3.7	-5.4	+0.6	+2.0	-0.57	+0.64	+0.94	+0.90
ACC	95%	93%	88%	87%	86%	84%	80%	87%	68%	99%	99%	73%
Perc	1	6	3	24	99	99	34	59	2	13	43	16

Selection Indexes

$A	$A-L
$238	$438
19	4

Traits Observed: BWT, Genomics

Statistics: Number of Herds: 33, Prog Analysed: 518, Genomic Prog: 373

RS — MUSGRAVE APACHE ^{SV} — USA18194405

DOB: 04/01/2015 **Registration Status:** HBR **Mating Type:** Natural **Genetic Status:** AMF,CAF,DDF,NHF,DWF,MHF,OHF,OSF,RGF

SITZ UPWARD 307R ^{SV}
KOUPALS B&B IDENTITY ^{SV}
B&B ERICA 605 [#]
Sire: USA17264774 MUSGRAVE AVIATOR ^{SV}
S A V FINAL ANSWER 0035 #
MCATL FOREVER LADY 1429-138 #
ALC FOREVER LADY R02S #

HOOVER DAM #
MUSGRAVE BOULDER PV
MILL BRAE SA JAUNTY 3079 #
Dam: USA17606917 MUSGRAVE CAROLINE 1304-189 #
S A V NET WORTH 4200 #
MCATL LADY CAROLINE 189-1615 #
M A LADY CAROLINE 1615-3106 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+9.5	+8.3	-3.1	+1.1	+45	+77	+99	+50	+0.05	+5.5	+28	+1.8
ACC	88%	73%	98%	98%	97%	97%	97%	94%	65%	79%	95%	58%
Perc	3	7	73	6	81	91	89	99	93	92	2	31

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+1.8	+0	+65	+5.7	+0.7	-0.2	+0.2	+1.8	+0.10	+0.92	+1.12	+1.10
ACC	96%	93%	91%	88%	89%	88%	83%	88%	71%	89%	89%	83%
Perc	63	99	62	59	34	49	59	64	36	66	82	73

Selection Indexes

$A	$A-L
$212	$330
46	71

Traits Observed: Genomics

Statistics: Number of Herds: 28, Prog Analysed: 482, Genomic Prog: 133

RS — POSS ELEMENT 215 # — USA17383988

DOB: 29/01/2012 **Registration Status:** HBR **Mating Type:** Natural **Genetic Status:** AMFU,CAFU,DDF,NHFU

TC TOTAL 410 #
POSS TOTAL IMPACT 745 #
POSS BLACKCAP 5116 #
Sire: USA16750909 POSS EASY IMPACT 0119 #
HYLINE RIGHT TIME 338 #
POSS ELMARETTA 736 #
POSS ELMARETTA 025 #

C A FUTURE DIRECTION 5321 ^{SV}
ALC BIG EYE D09N #
ALC HAZEL L12L #
Dam: USA16784565 POSS ERICA 004 #
RITO 1I2 OF 2536 RITO 6I6 #
POSS ERICA 814 #
POSS ERICA 627 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+4.2	+7.0	-7.0	+0.9	+55	+91	+119	+115	+0.34	+7.1	+16	+1.4
ACC	83%	67%	97%	96%	95%	95%	94%	91%	91%	91%	92%	52%
Perc	38	15	16	5	34	59	55	30	29	73	59	7

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+1.4	+14	+63	+6.7	+1.6	+2.9	-0.1	+1.5	+0.27	+0.92	+0.96	+0.96
ACC	93%	86%	88%	87%	86%	86%	81%	87%	67%	90%	89%	41%
Perc	77	79	69	47	18	9	75	72	55	66	48	30

Selection Indexes

$A	$A-L
$226	$399
30	18

Traits Observed: Genomics

Statistics: Number of Herds: 16, Prog Analysed: 187, Genomic Prog: 45

DOB: **14/01/2019**   Registration Status: **HBR**   Mating Type: **Natural**   Genetic Status: **AMF,CAF,DDF,NHF,DWF,MAF,MHF,OHF,OSF,**

S A V FINAL ANSWER 0035 #
CONNEALY CAPITALIST 028 #
PRIDES PITA OF CONANGA 8821 #
Sire: **USA17666102 LD CAPITALIST 316** ^{PV}
C A FUTURE DIRECTION 5321 ^{SV}
LD DIXIE ERICA 2053 #
LD DIXIE ERICA OAR 0853 #

PA POWER TOOL 9108 ^{SV}
RAVEN POWERBALL 53 ^{PV}
RAVEN EMMA E 4241 #
Dam: **USA19014827 ROLLIN ROCK BLACKBIRD 7059** #
E&B 878 NEW DESIGN 435
ROLLIN ROCK BLACKBIRD 9080 #
ROLLIN ROCK BLACKBIRD 7225 #

Mid January 2025 TransTasman Angus Cattle Evaluation

TACE	Dir	Dtrs	GL	BW	200 W	400 W	600 W	MCW	MBC	MCH	Milk	DTC
EBV	+12.0	+10.1	-9.2	-0.4	+65	+121	+154	+125	+0.28	+6.6	+21	+3.2
ACC	78%	66%	98%	97%	94%	95%	92%	88%	72%	74%	82%	56%
Perc	1	2	4	1	7	3	4	18	45	82	23	79

TACE	SS	Doc	CWT	EMA	Rib	Rump	RBY	IMF	NFI-F	CS	FA	LA
EBV	+3.2	+6	+93	+6.1	+0.1	-1.3	-0.7	+3.4	+0.82	+0.96	+1.08	+0.94
ACC	90%	85%	84%	83%	82%	81%	76%	84%	68%	85%	85%	70%
Perc	17	96	5	54	48	69	94	27	95	73	76	24

Selection Indexes

$A	$A-L
$235	**$423**
21	**8**

Traits Observed: Genomics

Statistics: Number of Herds: 25, Prog Analysed: 323, Genomic Prog: 190

Understanding the
TransTasman
Angus Cattle Evaluation (TACE)

TACE
TransTasman Angus Cattle Evaluation

What is the TransTasman Angus Cattle Evaluation?

The TransTasman Angus Cattle Evaluation is the genetic evaluation program adopted by Angus Australia for Angus and Angus influenced beef cattle. The TransTasman Angus Cattle Evaluation uses Best Linear Unbiased Prediction (BLUP) technology to produce Estimated Breeding Values (EBVs) of recorded cattle for a range of important production traits (e.g. weight, carcase, fertility).

The TransTasman Angus Cattle Evaluation is an international genetic evaluation and includes pedigree, performance and genomic information from the Angus Australia and Angus New Zealand databases, along with selected information from the American and Canadian Angus Associations.

The TransTasman Angus Cattle Evaluation utilises a range of genetic evaluation software, including the internationally recognised BLUPF90 family of programs, and BREEDPLAN® beef genetic evaluation analytical software, as developed by the Animal Genetics and Breeding Unit (AGBU), a joint institute of NSW Agriculture and the University of New England, and Meat and Livestock Australia Limited (MLA).

What is an EBV?

An animal's breeding value can be defined as its genetic merit for each trait. While it is not possible to determine an animal's true breeding value, it is possible to estimate it. These estimates of an animal's true breeding value are called EBVs (Estimated Breeding Values).

EBVs are expressed as the difference between an individual animal's genetics and a historical genetic level (i.e. group of animals) within the TACE genetic evaluation, and are reported in the units in which the measurements are taken.

Using EBVs to Compare the Genetics of Two Animals

TACE EBVs can be used to estimate the expected difference in the genetics of two animals, with the expected difference equating to half the difference in the EBVs of the animals, all other things being equal (e.g. they are joined to the same animal/s).

For example, a bull with a 200 Day Growth EBV of +60 would be expected to produce progeny that are, on average, 10 kg heavier at 200 days of age than a bull with a 200 Day Growth EBV of +40 kg (i.e. 20 kg difference between the sire's EBVs, then halved as the sire only contributes half the genetics).

Or similarly, a bull with an IMF EBV of +3.0 would be expected to produce progeny with on average, 1% more intramuscular fat in a 400 kg carcase than a bull with a IMF EBV of +1.0 (i.e. 2% difference between the sire's EBVs, then halved as the sire only contributes half the genetics).

Using EBVs to Benchmark an Animal's Genetics with the Breed

EBVs can also be used to benchmark an animal's genetics relative to the genetics of other Angus or Angus infused animals recorded with Angus Australia.

To benchmark an animal's genetics relative to other Angus animals, an animal's EBV can be compared to the EBV reference tables, which provide:

- the breed average EBV
- the percentile bands table

The current breed average EBV is listed on the bottom of each page in this publication, while the current EBV reference tables are included at the end of these introductory notes.

For easy reference, the percentile band in which an animal's EBV ranks is also published in association with the EBV.

Considering Accuracy

An accuracy value is published with each EBV, and is usually displayed as a percentage value immediately below the EBV.

The accuracy value provides an indication of the reliability of the EBV in estimating the animal's genetics (or true breeding value), and is an indication of the amount of information that has been used in the calculation of the EBV.

EBVs with accuracy values below 50% should be considered as preliminary or of low accuracy, 50-74% as of medium accuracy, 75-90% of medium to high accuracy, and 90% or greater as high accuracy.

Description of TACE EBVs

EBVs are calculated for a range of traits within TACE, covering calving ease, growth, fertility, maternal performance, carcase merit, feed efficiency and structural soundness. A description of each EBV included in this publication is provided on the following page.

UNDERSTANDING ESTIMATED BREEDING VALUES (EBVS)

Category	Trait	Units	Description	Interpretation
Calving Ease/Birth	CEDir	%	Genetic differences in the ability of a sire's calves to be born unassisted from 2 year old heifers.	Higher EBVs indicate fewer calving difficulties in 2 year old heifers.
Calving Ease/Birth	CEDtrs	%	Genetic differences in the ability of a sire's daughters to calve unassisted at 2 years of age.	Higher EBVs indicate fewer calving difficulties in 2 year old heifers.
Calving Ease/Birth	GL	days	Genetic differences between animals in the length of time from the date of conception to the birth of the calf.	Lower EBVs indicate shorter gestation length.
Calving Ease/Birth	BW	kg	Genetic differences between animals in calf weight at birth.	Lower EBVs indicate lighter birth weight.
Growth	200 Day	kg	Genetic differences between animals in live weight at 200 days of age due to genetics for growth.	Higher EBVs indicate heavier live weight.
Growth	400 Day	kg	Genetic differences between animals in live weight at 400 days of age.	Higher EBVs indicate heavier live weight.
Growth	600 Day	kg	Genetic differences between animals in live weight at 600 days of age.	Higher EBVs indicate heavier live weight.
Maternal	MCH	cm	Genetic differences between animals in the height of mature females.	Higher EBVs indicate taller mature females.
Maternal	MBC	score	Genetic differences between animals in the body condition of mature females.	Higher EBVs indicate more body condition of mature females.
Maternal	MCW	kg	Genetic differences between animals in live weight of cows at 5 years of age.	Higher EBVs indicate heavier mature weight.
Maternal	Milk	kg	Genetic differences between animals in live weight at 200 days of age due to the maternal contribution of its dam.	Higher EBVs indicate heavier live weight.
Fertility	DtC	days	Genetic differences between animals in the time from the start of the joining period (i.e. when the female is introduced to a bull) until subsequent calving.	Lower EBVs indicate shorter time to calving.
Fertility	SS	cm	Genetic differences between animals in scrotal circumference at 400 days of age.	Higher EBVs indicate larger scrotal circumference.
Carcase	CWT	kg	Genetic differences between animals in hot standard carcase weight at 750 days of age.	Higher EBVs indicate heavier carcase weight.
Carcase	EMA	cm^2	Genetic differences between animals in eye muscle area at the 12/13th rib site in a 400 kg carcase.	Higher EBVs indicate larger eye muscle area.
Carcase	Rib Fat	mm	Genetic differences between animals in fat depth at the 12/13th rib site in a 400 kg carcase.	Higher EBVs indicate more fat.
Carcase	P8 Fat	mm	Genetic differences between animals in fat depth at the P8 rump site in a 400 kg carcase.	Higher EBVs indicate more fat.
Carcase	RBY	%	Genetic differences between animals in boned out saleable meat from a 400 kg carcase.	Higher EBVs indicate higher yield.
Carcase	IMF	%	Genetic differences between animals in intramuscular fat (marbling) at the 12/13th rib site in a 400 kg carcase.	Higher EBVs indicate more intramuscular fat.
Feed/Temp.	NFI-F	kg/day	Genetic differences between animals in feed intake at a standard weight and rate of weight gain when animals are in a feedlot finishing phase.	Lower EBVs indicate more feed efficiency.
Feed/Temp.	Doc	%	Genetic differences between animals in temperament.	Higher EBVs indicate better temperament.
Structure	Claw Set	score	Genetic differences in claw set structure (shape and evenness of claws).	Lower EBVs indicate less curl of the claw set.
Structure	Foot Angle	score	Genetic differences in foot angle (strength of pastern, depth of heel).	Lower EBVs indicate more heel depth.
Structure	Leg Angle	score	Genetic differences in rear leg structure when viewed from the side (angle at front of the hock).	Lower EBVs indicate a less angular leg angle.
Selection Index	$A	$	Genetic differences between animals in net profitability per cow joined in a typical commercial self replacing herd using Angus bulls. This selection index is not specific to a particular market end-point, but identifies animals that will improve overall net profitability in the majority of commercial, self replacing, grass and grain finishing beef production systems.	Higher selection indexes indicate greater profitability.
Selection Index	$A-L	$	The $A-L index is similar to the $A index but is modelled on a production system where feed is surplus to requirements for the majority of the year, or the cost of supplying additional feed when animal feed requirements increase is low. While the $A aims to maintain mature cow weight, the $A-L does not aim to limit the increase in mature cow weight as there is minimal cost incurred if the feed maintenance requirements of the female breeding herd increase as a result of selection decisions.	Higher selection indexes indicate greater profitability.

UNDERSTANDING ESTIMATED BREEDING VALUES (EBVS)

Selection Indexes				
$D	$	Genetic differences between animals in net profitability per cow joined in a commercial self replacing herd targeting the domestic supermarket trade. Steers are either finished using pasture, pasture supplemented by grain, or grain (e.g. 50 -70 days) with steers assumed to be slaughtered at 510kg live weight (280kg carcase weight with 12mm P8 fat depth) at 16 months of age.		Higher selection indexes indicate greater profitability.
$D-L	$	The $D-L index is similar to the $D index but is modelled on a production system where feed is surplus to requirements for the majority of the year, or the cost of supplying additional feed when animal feed requirements increase is low. While the $D aims to maintain mature cow weight, the $D-L does not aim to limit the increase in mature cow weight as there is minimal cost incurred if the feed maintenance requirements of the female breeding herd increase as a result of selection decisions.		Higher selection indexes indicate greater profitability.
$GN	$	Genetic differences between animals in net profitability per cow joined in a commercial self replacing herd targeting pasture grown steers with a 250 day feedlot finishing period for the grain fed high quality, highly marbled markets. Steers are assumed to be slaughtered at 800 kg live weight (455 kg carcase weight with 30 mm P8 fat depth) at 24 months of age, with a significant premium for steers that exhibit superior marbling.		Higher selection indexes indicate greater profitability.
$GN-L	$	The $GN-L index is similar to the $GN index but is modelled on a production system where feed is surplus to requirements for the majority of the year, or the cost of supplying additional feed when animal feed requirements increase is low. While the $GN aims to maintain mature cow weight, the $GN-L does not aim to limit the increase in mature cow weight as there is minimal cost incurred if the feed maintenance requirements of the female breeding herd increase as a result of selection decisions.		Higher selection indexes indicate greater profitability.
$GS	$	Genetic differences between animals in net profitability per cow joined in a commercial self replacing herd targeting pasture finished steers. Steers are assumed to be slaughtered at 650 kg live weight (350 kg carcase weight with 12 mm P8 fat depth) at 22 months of age. Emphasis has been placed on eating quality and tenderness to favour animals that are suited to MSA requirements.		Higher selection indexes indicate greater profitability.
$GS-L	$	The $GS-L index is similar to the $GS index but is modelled on a production system where feed is surplus to requirements for the majority of the year, or the cost of supplying additional feed when animal feed requirements increase is low. While the $GS aims to maintain mature cow weight, the $GS-L does not aim to limit the increase in mature cow weight as there is minimal cost incurred if the feed maintenance requirements of the female breeding herd increase as a result of selection decisions.		Higher selection indexes indicate greater profitability.
$PRO	$	Genetic differences between animals in net profitability per cow joined in a commercial self replacing herd based in New Zealand that targets the production of grass finished steers for the AngusPure programme. Steers are assumed marketed at approximately 530 kg live weight (290 kg carcase weight with 10 mm P8 fat depth) at 20 months of age, with a significant premium for steers that exhibit superior marbling.		Higher selection indexes indicate greater profitability.
$T	$	Genetic difference between animals in net profitability per cow joined in a situation where Angus bulls are being used as a terminal sire over mature breeding females and all progeny, both male and female, are slaughtered. The Angus Terminal Sire Index focusses on increasing growth, carcase yield and eating quality. Daughters are not retained for breeding and therefore no emphasis is given to female fertility or maternal traits.		Higher selection indexes indicate greater profitability.