

TOTAL

LIVESTOCK GENETICS



SIRE DIRECTORY

AUGUST 2025

FOCUSED ON YOU,
YOUR HERD & YOUR FUTURE

Focused on You, Your Herd and Your Future Welcome to the Total Livestock Genetics 2025 Dairy Sire Catalogue.

At Total Livestock Genetics, our focus has always been simple — deliver the genetics that move your herd forward. That means we listen to what breeders want not only today but also with an eye towards tomorrow. Being part of the Urus group means we can source the very best sires from around the world, regardless of postcode, with one clear goal — to meet the demands of Australian dairy farmers and help them achieve lasting success. By combining global reach with a sharp focus on local priorities, we ensure our line-up reflects not just the latest in genetic progress, but also the traits and profiles that matter most in your milking sheds.

Leading the way this year are two outstanding Excitement sons. Peak **ENTERPRISE** from Genex delivers on all fronts — 553 BPI, 3286 GTPI and an elite conformation profile backed by strength, balance and production. **ENTERPRISE** is the kind of sire that truly ticks all the boxes for progressive breeding programs. Alongside him is **ELVIN**, another Excitement son bred in Australia, who brings elite mammary credentials (+1.05 UDC), strong udder balance, and the kind of modern production profile highlighted by +0.83% Fat and +159 ASI. With his combination of type and components, **ELVIN** is a young sire with genuine international appeal.

Also taking centre stage is **APOLLO-PP**, one of the most complete homozygous polled sires on the market. Canada's #1 GPA-LPI Homozygous Polled bull, **APOLLO PP** offers excellent type, outstanding mammary traits, and the added benefits of A2A2 and PP status. Combining 417 BPI with 112 Daughter Fertility and 2957 GTPI, +2.14 PTAT and +1.73 UDC. He represents a rare combination of conformation, functionality and modern polled convenience.

When it comes to looking to the future our expanded Polled Feature demonstrates that going polled no longer means compromising on performance. Bulls that bring type to the table, like **MDOUGLAS-PP**, add chest and muzzle width without excess stature, while sires such as **BEACON-PP** meet the industry's everyday requirements for high fertility and longer teat length. Adding to this group is **LANGLEY-PP**, an Australian-bred homozygous polled son of **MDOUGLAS-PP** who combines balanced production (+29 Fat, +16 Protein) with improved udders (+109 Udder Depth, +105 Fore Attachment) and strong fertility (+108 Daughter Fertility). Others in the line-up bring extra milk, strength, and overall balance. This team means breeders can finally have their homozygous polled without any compromises.

Our Jersey offering is just as exciting, Attaview **INCHSTA-P** offers a balanced package with a 324 BPI, and strong mammary system and type composites, making him a versatile option for production with functional type. Peak **DOUBS-PP**, a Genex based A2A2 Homozygous Polled Jersey, delivers +665 lbs Milk, +70 CFP, +1.00 PTAT and positive teat length, offering international genetics with commercial appeal. For Brown Swiss breeders, Hilltop Acres **TRACTION** combines strength, production, and udder quality, adding +1171 lbs Milk, +3.4 Productive Life, and strong udder attachment to a modern, balanced type profile.

Whether your goal is genetic progress, herd health, or future-proofing your breeding program, TLG remains committed to genetics that are focused on you, your herd, and your future.



Mat Dennis
TLG Product Specialist



Disclaimer:

The content of this Catalogue is provided for information only. It is published with care and attention to accuracy, nevertheless Genetics Australia Holdings Pty Ltd and all affiliates (GAH) accepts no liability if, for any reason the information is inaccurate or out of date whether negligent or otherwise.

Limitation of Liability:

GAH warrants that it has clear title to the products sold and that all semen processed in its laboratory is processed under rigid quality controls. Since GAH does not control conditions under which its products are used it does not give, and its agents and employees are forbidden to give, any warranty expressed or implied as to the conception rate or productiveness or the results which may be obtained by the use of any product sold or in connection with any techniques recommended. GAH makes no other warranty of any kind whatsoever expressed or implied which extends beyond the description of its products. Where so permitted the liability of GAH for breach of a condition or warranty implied by statute is limited at GAH's option to the replacement or repair of the goods or the supply of equivalent goods or the cost of replacing or repairing the goods or of acquiring equivalent goods. So far as the law permits GAH shall not be liable in any way whatsoever for any indirect or consequential loss or loss of profit.



PLUS20 - it's our commitment that the conventional semen carrying this logo contains over 20 Million sperm cells per straw and has over 20 years of development.

Sexed (sorted) semen:

Artificial Insemination products (straws) that GAH offers for sale as sexed (sorted) semen using Sexing Technologies proprietary technology, shall only be used for single-use artificial insemination and not for in-vitro-fertilisation or embryo transfer.

Prices:

All prices listed within this publication are listed exclusive of Goods and Services Tax (GST), so GST will be added at the time of purchase.

Source data:

Data referencing Australian Breeding Values (ABVs') information unless otherwise stated is sourced from DataGene August 2025 breeding value release. Data referencing United States of America Breeding Values are provided through the Council on Dairy Cattle Breeding (CDCB) this information unless otherwise stated is sourced from CDCB August 2025 breeding value release.

Genetic Codes and Haplotype Abbreviations:

For a full breakdown of the Genetic Codes and Haplotypes referenced in this directory please visit the website www.genaust.com.au



Available also in Sexed Ultraplus Semen

APOLLO-pp

Vector Fra Apollo-PP (Logic-PP / Allday / Hotspot-P)

NEW

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGAPOLLO NAAB: 724H002040 HB: HOCAN000014911360 DOB: 28/11/2023

SEXED \$70.00
UltraPlus

RRP \$38.00

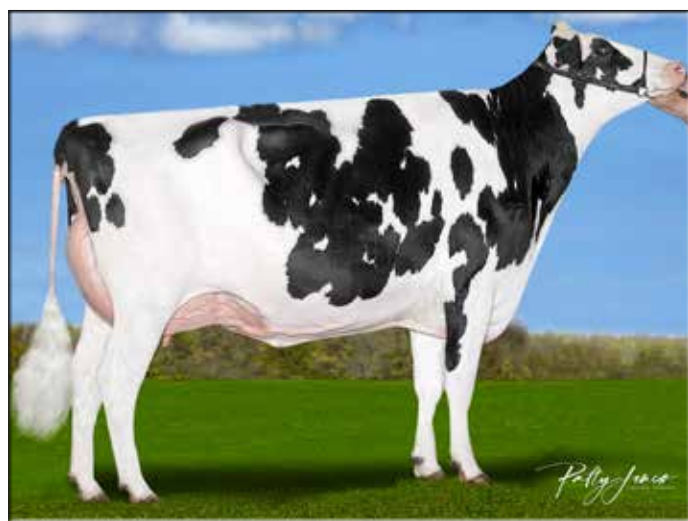
USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
2957	163	50	0.16	21	0.06	79	273
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0	0	-0.3	1.3	1.4	2.92	2.14	1.73

CONFORMATION 0 Dtrs 0 Herds				USDA/HA Genomic Evaluation				August 2025
PTAT: 2.14	UDC: 1.73	FLC: 1.21						GTPI 2957
Stature	Short						Tall	1.60
Strength	Frail						Strong	0.46
Body Depth	Shallow						Deep	0.98
Dairy Form	Tight Rib						Open Rib	1.66
Rump Angle	High						Sloped	0.91
Rump width	Narrow						Wide	1.29
RL SV	Posty						Sickle	-0.06
RL RV	Hock-In						Straight	1.25
Foot Angle	Low						Steep	0.93
F & L Score	Low						High	1.55
Fore Attachment	Loose						Strong	2.11
Rear Udder Height	Low						High	2.66
Rear Udder Width	Narrow						Wide	2.57
Udder Cleft	Weak						Strong	0.91
Udder Depth	Deep						Shallow	1.52
Fore Teat Placement	Wide						Close	0.25
Rear Teat Placement	Wide						Close	0.39
Teat Length	Short						Long	0.68

- "One of the most complete sires I have ever seen" Dave Eastman, Validity Sires
- Positive CCR, DPR, A2A2,PP & Teat Length
- 417 BPI, 112 Daughter Fertility
- +2.14 PTAT +1.73 UDC 2957 GTPI
- +14 Conformation, 3924 LPI



Vector Fra Apollo-PP



Dam of Apollo: Fraholme Allday Arielle-P VG86-2YRS

BPI 417 65 %R

HWI 420 59 %R

SI 622 62 %R

PRODUCTION ABV (g) 08/25

Milk	-15	77% R
Fat	44 kg	0.65%
Protein	11 kg	0.22%
ASI	169	77 % R

WORKABILITIES

Milking Speed	103
Temperament	101
Likability	104

Conformation ABV(g)

Overall Type	103
Mammary	108
Rump	105
Dairy Strength	97
Feet and Legs	98

HEALTH ABV (g) 08/25

Cell Count	163	66
Survival	108	56
Feed Saved	-121	41
Dtr Fertility	112	58
Calving Ease	102	68
Gest Length	-1	61
Heat Tol	100	48

CONFORMATION ABV (g)	0 Dtrs	0 Herds	61%Rel	August 2025
OVERALL TYPE	103		MAMMARY SYSTEM	108
Stature	103		Pin Width	105
Bone Quality	108		Pin Set	104
Angularity	97		Udder Texture	99
Muzzle Width	96		Udder Depth	106
Body Depth	98		Fore Attachment	105
Chest Width	96		Rear Att Height	110
Loin Strength	105		Rear Att Width	104
Foot Angle	99		Centre Ligament	103
Rear Set	102		Teat Placement (Front)	102
Rear Leg Rear View	100		Teat Placement (Rear)	101
			Teat Length	97

EXCITEMENT

Peak Excitement (Exquisite / Pursuit / Emerald)

GENEX

A2 STATUS: A1/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXEXCITEMENT NAAB: 001H016675 HB: H0840M003252198074 DOB: 20/10/2022

SEXED \$58.00 RRP \$28.00

UltraPlus

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3313	130	78	0.27	42	0.14	81	805

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
1.7	0.4	4.8	1	-0.7	2.75	1.14	0.26



Peak Excitement

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
435	364	756	253	-210	47	0.81%	20	0.49%	77

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability	
106	101	175	102	102	103	
C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
103	-5	-108	109	91	102	102

○ Short Gestation -5 Days and 103 Calving Ease

○ International sire of sons

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.14 UDC: 0.26 FLC: 1.01 GTPI 3313

Stature	Short					Tall	0.75
Strength	Frail					Strong	0.47
Body Depth	Shallow					Deep	0.47
Dairy Form	Tight Rib					Open Rib	0.75
Rump Angle	High					Sloped	0.29
Rump width	Narrow					Wide	0.87
RL SV	Posty					Sickle	0.06
RL RV	Hock-In					Straight	1.26
Foot Angle	Low					Steep	0.92
F & L Score	Low					High	1.06
Fore Attachment	Loose					Strong	0.70
Rear Udder Height	Low					High	0.59
Rear Udder Width	Narrow					Wide	1.21
Udder Cleft	Weak					Strong	-0.96
Udder Depth	Deep					Shallow	0.13
Fore Teat Placement	Wide					Close	0.00
Rear Teat Placement	Wide					Close	-0.51
Teat Length	Short					Long	0.45

-1.5 -1.0 -0.5 0 0.5 1.0 1.5

ELVIN

Summit View Elvin (Excitement / Porsche / Challenger)

NEW

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGELVIN NAAB: 187H005857 HB: AUS2288808 DOB: 19/09/2024

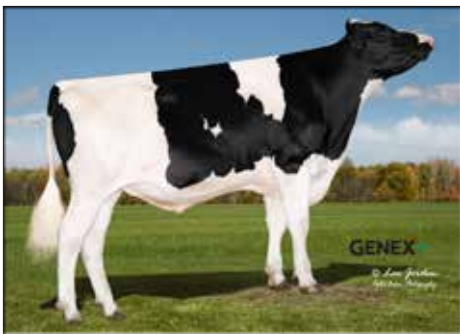
SEXED \$55.00 RRP \$26.00

UltraPlus

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3219	55	73	0.27	30	0.11	79	630

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
1.9	0.4	3.6	1.5	0.2	2.76	1.62	0.85



Sire: Peak Excitement

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
361	329	567	159	-286	45	0.83%	6	0.26%	76

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability	
104	99	163	102	103	103	
C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
103	-1	-27	109	99	101	105

○ Exciting NEW Australian standing Excitement son

○ 3219 GTPI, Positive CCR, DPR, A2A2

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.62 UDC: 0.85 FLC: 0.84 GTPI 3219

Stature	Short					Tall	1.28
Strength	Frail					Strong	1.06
Body Depth	Shallow					Deep	1.11
Dairy Form	Tight Rib					Open Rib	1.11
Rump Angle	High					Sloped	0.30
Rump width	Narrow					Wide	1.33
RL SV	Posty					Sickle	-0.21
RL RV	Hock-In					Straight	1.11
Foot Angle	Low					Steep	1.15
F & L Score	Low					High	1.02
Fore Attachment	Loose					Strong	0.99
Rear Udder Height	Low					High	1.22
Rear Udder Width	Narrow					Wide	1.80
Udder Cleft	Weak					Strong	0.55
Udder Depth	Deep					Shallow	0.52
Fore Teat Placement	Wide					Close	0.93
Rear Teat Placement	Wide					Close	0.71
Teat Length	Short					Long	-0.17

-1.5 -1.0 -0.5 0 0.5 1.0 1.5

ENTERPRISE

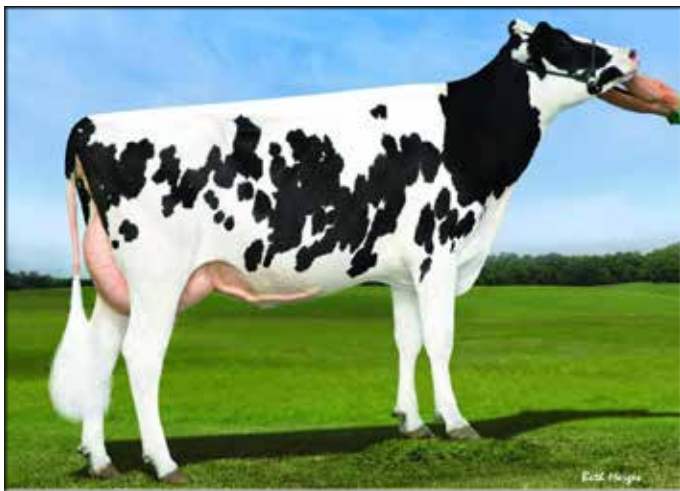
A2 STATUS: A1/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXENTERPRISE NAAB: 001H017453 HB: H0840003283239666 DOB: 17/05/2024

RRP \$32.00

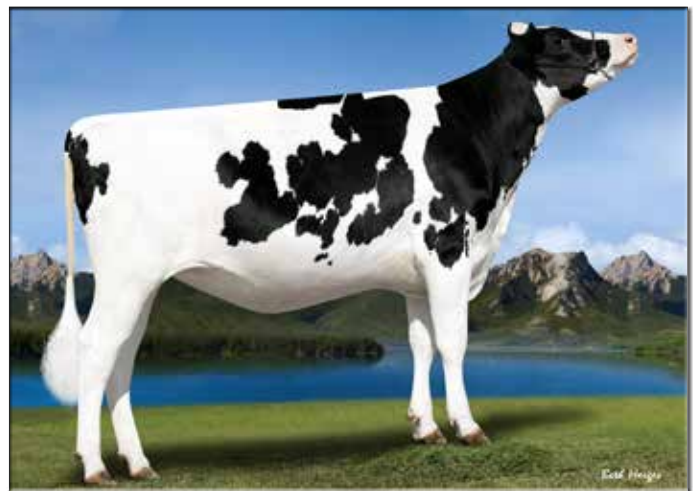
USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3286	330	73	0.22	41	0.11	78	706
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
1.2	0.3	4.9	1.4	-1.6	2.71	0.93	1.08

- New Release Excitement son
- 553 BPI, 3286 GTPI
- Moderate stature with strength and width
- Delivering milk with components
- High demand sire order early to avoid disappointment

CONFORMATION		0 Dtrs	0 Herds	USDA/HA Genomic Evaluation			August 2025	
PTAT: 0.93		UDC: 1.08		FLC: 0.33		GTPI 3286		
Stature	Short						Tall	0.94
Strength	Frail						Strong	0.88
Body Depth	Shallow						Deep	0.27
Dairy Form	Tight Rib						Open Rib	-0.44
Rump Angle	High						Sloped	1.36
Rump width	Narrow						Wide	1.24
RL SV	Posty						Sickle	-0.94
RL RV	Hock-In						Straight	0.75
Foot Angle	Low						Steep	1.14
F & L Score	Low						High	0.42
Fore Attachment	Loose						Strong	1.90
Rear Udder Height	Low						High	1.03
Rear Udder Width	Narrow						Wide	1.52
Udder Cleft	Weak						Strong	-0.24
Udder Depth	Deep						Shallow	1.61
Fore Teat Placement	Wide						Close	0.21
Rear Teat Placement	Wide						Close	-0.20
Teat Length	Short						Long	0.19
		-1.5	-1.0	-0.5	0	0.5	1.0	1.5



7th Dam: Moonry Myesha 9071



5th Dam: Pine Tree Mandolin

BPI 553 64 %R

HWI 527 58 %R

SI 963 61 %R

PRODUCTION ABV (g) 08/25

Milk	88	76% R
Fat	49 kg	0.65%
Protein	24 kg	0.42%
ASI	259	76 % R

WORKABILITIES

Milking Speed	102
Temperament	103
Likability	104

Conformation ABV(g)

Overall Type	102
Mammary	104
Rump	102
Dairy Strength	99
Feet and Legs	102

HEALTH ABV (g) 08/25

Cell Count	185	65
Survival	112	55
Feed Saved	-111	41
Dtr Fertility	112	55
Calving Ease	102	68
Gest Length	-6	61
Heat Tol	97	48

CONFORMATION ABV (g) 0 Dtrs 0 Herds 60%Rel August 2025

OVERALL TYPE	102	MAMMARY SYSTEM	104
Stature	97	Pin Width	104
Bone Quality	95	Pin Set	98
Angularity	94	Udder Texture	91
Muzzle Width	106	Udder Depth	104
Body Depth	98	Fore Attachment	106
Chest Width	105	Rear Att Height	105
Loin Strength	103	Rear Att Width	102
Foot Angle	102	Centre Ligament	97
Rear Set	100	Teat Placement (Front)	101
Rear Leg Rear View	100	Teat Placement (Rear)	96
		Teat Length	97



TODAY IS YOUR OPPORTUNITY **TO** BUILD THE TOMORROW YOU WANT.

At Trans Ova Genetics, we're empowering breeders to build a tomorrow that exceeds expectations. We provide the complete toolbox of services you need to create a legacy of excellence. Today's choices shape tomorrow's success. Choose Trans Ova Genetics and leave a lasting legacy for the next generation.

For more information, visit www.transova.com

TRANS 
genetics

**NOW PROVIDING IVF SERVICES IN AUSTRALIA
IN PARTNERSHIP WITH TOTAL LIVESTOCK GENETICS.**



PHONE (03) 5593 2016

LANGLEY - PP Summit View Langley PP (MDouglas -PP / Nipit-P/ Luster P)

NEW

A2 STATUS: A2/A2 Genetic Codes: TE, PP Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGLANGLEY NAAB: 187H005856 HB: AUS2281874 DOB: 27/07/2024

SEXED \$55.00 RRP \$26.00
Ultraplus CONVENTIONAL PLUS

USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
2972	703	56	0.10	37	0.05	79	392
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0.8	-0.9	0.9	1.6	1.4	3.09	1.09	1.33

CONFORMATION 0 Dtrs 0 Herds				USDA/HA Genomic Evaluation				August 2025
PTAT: 1.09	UDC: 1.33	FLC: -0.06						GTPI 2972
Stature	Short						Tall	0.40
Strength	Frail						Strong	0.07
Body Depth	Shallow						Deep	0.20
Dairy Form	Tight Rib						Open Rib	0.70
Rump Angle	High						Sloped	0.35
Rump width	Narrow						Wide	1.07
RL SV	Posty						Sickle	0.62
RL RV	Hock-In						Straight	-0.28
Foot Angle	Low						Steep	0.16
F & L Score	Low						High	0.12
Fore Attachment	Loose						Strong	1.49
Rear Udder Height	Low						High	1.67
Rear Udder Width	Narrow						Wide	1.42
Udder Cleft	Weak						Strong	0.86
Udder Depth	Deep						Shallow	1.27
Fore Teat Placement	Wide						Close	0.28
Rear Teat Placement	Wide						Close	0.79
Teat Length	Short						Long	0.69

- PP sire with +2972 GTPI, +1.33 UDC
- Wide sloped rumps
- A2A2, PP, Teat Length, CCR +0.8
- New release sire that we are excited about
- Medium stature with open rib



4th Dam: Stanton's Big Maker



Sire: Vala MDouglas-PP

BPI 343 65 %R

HWI 313 60 %R

SI 569 63 %R

PRODUCTION ABV (g) 08/25		
Milk	245	77% R
Fat	29 kg	0.27%
Protein	16 kg	0.18%
ASI	149	77 % R

WORKABILITIES	
Milking Speed	103
Temperament	102
Likability	104

Conformation ABV(g)	
Overall Type	97
Mammary	102
Rump	104
Dairy Strength	98
Feet and Legs	99

HEALTH ABV (g) 08/25		
Cell Count	160	67
Survival	109	57
Feed Saved	-165	42
Dtr Fertility	108	58
Calving Ease	102	70
Gest Length	-1	63
Heat Tol	99	48

CONFORMATION ABV (g)	0 Dtrs	0 Herds	62%Rel	August 2025
OVERALL TYPE	97		MAMMARY SYSTEM	102
Stature	102		Pin Width	103
Bone Quality	99		Pin Set	102
Angularity	97		Udder Texture	94
Muzzle Width	102		Udder Depth	109
Body Depth	99		Fore Attachment	105
Chest Width	101		Rear Att Height	106
Loin Strength	102		Rear Att Width	101
Foot Angle	98		Centre Ligament	97
Rear Set	102		Teat Placement (Front)	99
Rear Leg Rear View	98		Teat Placement (Rear)	104
			Teat Length	96

ENZO

Summit View Dropbox Enzo - ET (Dropbox / Einstein / Marius)

A2 STATUS: A1/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGENZO NAAB: 187H005738 HB: AUS2238370 DOB: 25/06/2023

SEXED \$55.00

RRP \$24.00

UltraPlus

CONVENTIONAL PLUS20

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3049	513	48	0.1	44	0.1	81	366

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0.1	-0.3	1.8	1.8	-0.1	2.66	1.32	1.18



Summit View Enzo

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
541	570	854	208	-105	35	0.57%	19	0.40%	77

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
118	103	194	100	102	103

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
103	-5	-118	110	95	101	104

First calves are pick of the calf pens

Fresh sexed semen sire

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.32 UDC: 1.18 FLC: -0.16 GTPI 3049

Stature	Short					Tall	2.08
Strength	Frail					Strong	1.41
Body Depth	Shallow					Deep	1.32
Dairy Form	Tight Rib					Open Rib	0.95
Rump Angle	High					Sloped	1.34
Rump width	Narrow					Wide	1.96
RL SV	Posty					Sickle	-0.72
RL RV	Hock-In					Straight	-0.36
Foot Angle	Low					Steep	1.11
F & L Score	Low					High	0.44
Fore Attachment	Loose					Strong	1.46
Rear Udder Height	Low					High	1.65
Rear Udder Width	Narrow					Wide	2.15
Udder Cleft	Weak					Strong	0.95
Udder Depth	Deep					Shallow	1.29
Fore Teat Placement	Wide					Close	0.98
Rear Teat Placement	Wide					Close	0.98
Teat Length	Short					Long	0.27

MAGNET

Bundalong Parfect Magnet (Parfect / Magnitude / Muscadet)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGMAGNET NAAB: 187H005716 HB: AUS2234385 DOB: 02/03/2023

RRP \$24.00

CONVENTIONAL PLUS20

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
2997	762	54	0.08	35	0.04	81	435

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0.3	-0.6	2.1	1.4	-1.2	2.79	0.58	0.42



Bundalong Parfect Magnet

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
443	416	858	232	610	44	0.26%	28	0.22%	79

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
109	103	154	102	102	105

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
102	-3	-112	110	91	99	103

+610L Milk with positive components

Wide sloped rumps and added teat length

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 0.58 UDC: 0.42 FLC: 0.34 GTPI 2997

Stature	Short					Tall	0.19
Strength	Frail					Strong	0.54
Body Depth	Shallow					Deep	0.26
Dairy Form	Tight Rib					Open Rib	-0.01
Rump Angle	High					Sloped	0.02
Rump width	Narrow					Wide	1.42
RL SV	Posty					Sickle	-0.08
RL RV	Hock-In					Straight	0.88
Foot Angle	Low					Steep	0.35
F & L Score	Low					High	0.21
Fore Attachment	Loose					Strong	0.47
Rear Udder Height	Low					High	0.85
Rear Udder Width	Narrow					Wide	0.86
Udder Cleft	Weak					Strong	0.21
Udder Depth	Deep					Shallow	-0.12
Fore Teat Placement	Wide					Close	-0.17
Rear Teat Placement	Wide					Close	0.25
Teat Length	Short					Long	0.65

VINDICATE

Peak Vindicate (AltaSamson / AltaZemini / Fortnite)

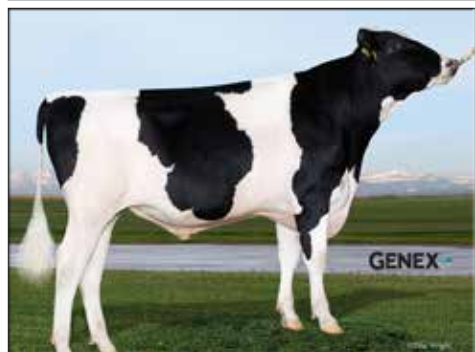
NEW GENEX

RRP \$34.00

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXVINDICATE NAAB:001H017196 HB: H0840003272456673 DOB: 23/10/2023

USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3353	1183	95	0.17	61	0.08	79	860

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0.1	-1.3	3.6	1.4	-1	2.59	0.85	0.27



Peak Vindicate

Australian Indices										August 2025
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%	
455	375	801	252	88	55	0.74%	21	0.36%	76	

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
105	100	172	101	102	103

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
102	-5	-143	111	87	99	100

○ High milk quality 2.59 SCS, 172 SCC

○ Flawless linear, high production 252 ASI

CONFORMATION	0 Dtrs	0 Herds	USDA/HA Genomic Evaluation	August 2025
PTAT: 0.85	UDC: 0.27	FLC: -0.06		GTPI 3353

Stature	Short				Tall	1.40
Strength	Frail				Strong	0.66
Body Depth	Shallow				Deep	0.65
Dairy Form	Tight Rib				Open Rib	0.98
Rump Angle	High				Sloped	0.27
Rump width	Narrow				Wide	1.06
RL SV	Posty				Sickle	-0.11
RL RV	Hock-In				Straight	-0.05
Foot Angle	Low				Steep	0.65
F & L Score	Low				High	0.29
Fore Attachment	Loose				Strong	0.65
Rear Udder Height	Low				High	0.30
Rear Udder Width	Narrow				Wide	0.99
Udder Cleft	Weak				Strong	0.16
Udder Depth	Deep				Shallow	0.63
Fore Teat Placement	Wide				Close	0.35
Rear Teat Placement	Wide				Close	0.11
Teat Length	Short				Long	0.16

CASIMIRO

Peak Casimiro (AltaOvertake / AltaZazzle / Renegade)

GENEX

SEXED \$55.00

RRP \$24.00

A2 STATUS: A2/A2 Genetic Codes: MW Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXCASIMIRO NAAB:001H016650 HB: USA3250025924 DOB: 22/09/2022

USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3338	363	60	0.17	41	0.11	80	723

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
4.4	2.7	4.5	1.5	0.8	2.72	1.12	1.39



Peak Casimiro

Australian Indices										August 2025
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%	
474	480	832	224	209	43	0.50%	22	0.32%	77	

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
116	99	161	102	100	102

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
104	0	-121	110	98	94	98

○ Global Sires of Sons with extreme fertility

○ 4.4 CCR, 2.7 DPR, 116 daughter fertility

CONFORMATION	0 Dtrs	0 Herds	USDA/HA Genomic Evaluation	August 2025
PTAT: 1.12	UDC: 1.39	FLC: 0.85		GTPI 3338

Stature	Short				Tall	0.35
Strength	Frail				Strong	-0.20
Body Depth	Shallow				Deep	-0.46
Dairy Form	Tight Rib				Open Rib	-0.03
Rump Angle	High				Sloped	0.10
Rump width	Narrow				Wide	0.82
RL SV	Posty				Sickle	-1.07
RL RV	Hock-In				Straight	0.97
Foot Angle	Low				Steep	1.15
F & L Score	Low				High	0.84
Fore Attachment	Loose				Strong	1.72
Rear Udder Height	Low				High	1.60
Rear Udder Width	Narrow				Wide	1.35
Udder Cleft	Weak				Strong	0.41
Udder Depth	Deep				Shallow	1.71
Fore Teat Placement	Wide				Close	0.24
Rear Teat Placement	Wide				Close	0.09
Teat Length	Short				Long	0.40

COMET

Peak Comet (Maverick / Wheelhouse / Lionel)

GENEX

A2 STATUS: A1/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXCOMET NAAB: 001H016959 HB: H0840M00326337326 DOB: 06/03/2023

SEXED \$55.00
UltraPlus

RRP \$24.00

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3222	867	80	0.16	36	0.03	79	688

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0.7	-0.4	3	1.6	0	2.91	1.41	1.37



Peak Comet

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
370	332	610	187	70	47	0.63%	14	0.23%	76

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability	
106	100	153	103	102	105	
C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
103	-4	-116	108	97	101	107

○ Trouble free, No holes proof

○ Backed by generations of success

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.41 UDC: 1.37 FLC: 0.4 GTPI 3222

Stature	Short					Tall	0.73
Strength	Frail					Strong	0.36
Body Depth	Shallow					Deep	0.38
Dairy Form	Tight Rib					Open Rib	1.25
Rump Angle	High					Sloped	0.54
Rump width	Narrow					Wide	1.40
RL SV	Posty					Sickle	0.65
RL RV	Hock-In					Straight	0.46
Foot Angle	Low					Steep	0.12
F & L Score	Low					High	0.59
Fore Attachment	Loose					Strong	1.24
Rear Udder Height	Low					High	1.90
Rear Udder Width	Narrow					Wide	2.32
Udder Cleft	Weak					Strong	0.30
Udder Depth	Deep					Shallow	0.90
Fore Teat Placement	Wide					Close	0.63
Rear Teat Placement	Wide					Close	0.67
Teat Length	Short					Long	0.22

LUCKYCHARM

Peak Luckycharm (Marlon x Magnifique x AltaZazzle)

GENEX

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXLUCKYCHARM NAAB: 001H016863 HB: H0840M003269404436 DOB: 13/03/2023

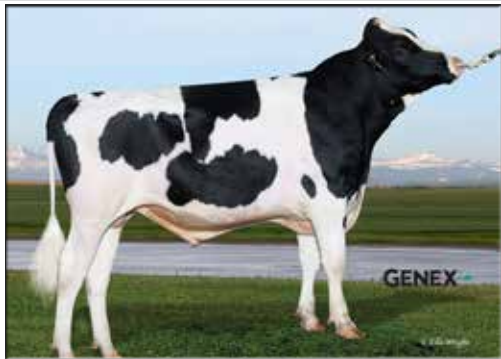
SEXED \$52.00
UltraPlus

RRP \$22.00

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3253	822	75	0.15	44	0.06	79	685

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
1.5	0.4	3.6	1.3	-1.3	2.79	1.09	1.17



Peak Luckycharm

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
440	430	806	231	303	49	0.53%	23	0.27%	76

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability	
112	99	164	100	100	102	
C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
104	-6	-68	109	94	101	99

○ Semen Fertility of +1.8 SCR

○ 3253 GTPI, high milk flow with positive components

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.09 UDC: 1.17 FLC: 0.52 GTPI 3253

Stature	Short					Tall	0.88
Strength	Frail					Strong	0.45
Body Depth	Shallow					Deep	0.36
Dairy Form	Tight Rib					Open Rib	0.41
Rump Angle	High					Sloped	1.66
Rump width	Narrow					Wide	0.85
RL SV	Posty					Sickle	-0.63
RL RV	Hock-In					Straight	0.67
Foot Angle	Low					Steep	1.06
F & L Score	Low					High	0.65
Fore Attachment	Loose					Strong	1.63
Rear Udder Height	Low					High	1.39
Rear Udder Width	Narrow					Wide	1.56
Udder Cleft	Weak					Strong	0.25
Udder Depth	Deep					Shallow	1.14
Fore Teat Placement	Wide					Close	0.99
Rear Teat Placement	Wide					Close	0.86
Teat Length	Short					Long	-0.37

CARTEL

Emu Banks Cartel (Redcarpet / Balisto / Numero Uno)

A2 STATUS: A2/A2
NASIS: TLGCARTEL

Genetic Codes: TE
HB: AUS2146036

Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
DOB: 28/03/2021

SEXED \$50.00
Ultraplus

RRP \$22.00
CONVENTIONAL PLUS20



Dam: Emu Banks Balisto Donatello EX90-1E

PRODUCTION ABV (g) 08/25

Milk	-771	86% R
Fat	1 kg	0.49%
Protein	-5 kg	0.31%
ASI	34	86 % R

WORKABILITIES

Milking Speed	102
Temperament	101
Likability	102

BPI 296 73 %R

HWI 368 67 %R

SI 336 70 %R

Conformation ABV(g)

Overall Type	110
Mammary	102
Rump	104
Dairy Strength	100
Feet and Legs	99

HEALTH ABV (g) 08/25

Cell Count	163	70
Survival	107	63
Feed Saved	-28	45
Dtr Fertility	114	65
Calving Ease	104	99
Gest Length	-4	99
Heat Tol	105	48

CONFORMATION ABV (g)	4 Dtrs	2 Herds	67%Rel	August 2025
OVERALL TYPE	110	MAMMARY SYSTEM	102	
Stature	99	Pin Width	103	
Bone Quality	106	Pin Set	97	
Angularity	98	Udder Texture	100	
Muzzle Width	97	Udder Depth	106	
Body Depth	100	Fore Attachment	106	
Chest Width	100	Rear Att Height	104	
Loin Strength	101	Rear Att Width	98	
Foot Angle	98	Centre Ligament	98	
Rear Set	102	Teat Placement (Front)	99	
Rear Leg Rear View	95	Teat Placement (Rear)	93	
		Teat Length	102	

- Now with over 100 daughters in his proof
- High fertility sire +114 with calving ease 104 and short gestation -4 days
- A2A2 and predominately black

MRRIGHT

Vala Mr Right (Almamater / Bandares / Silver)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGMRRIGHT NAAB:187H005559 HB: AUS2143902 DOB: 27/03/2021

SEXED \$55.00
Ultraplus

RRP \$24.00
CONVENTIONAL PLUS20



Vala MrRight

PRODUCTION ABV (g) 08/25

Milk	67	79% R
Fat	36 kg	0.47%
Protein	15 kg	0.26%
ASI	172	79 % R

WORKABILITIES

Milking Speed	105
Temperament	100
Likability	104

BPI 482 69 %R

HWI 519 64 %R

SI 776 66 %R

Conformation ABV(g)

Overall Type	107
Mammary	106
Rump	105
Dairy Strength	99
Feet and Legs	104

HEALTH ABV (g) 08/25

Cell Count	156	70
Survival	112	62
Feed Saved	-113	43
Dtr Fertility	117	66
Calving Ease	103	97
Gest Length	-5	98
Heat Tol	101	48

CONFORMATION ABV (g)	0 Dtrs	0 Herds	64%Rel	August 2025
OVERALL TYPE	107	MAMMARY SYSTEM	106	
Stature	101	Pin Width	101	
Bone Quality	98	Pin Set	106	
Angularity	94	Udder Texture	97	
Muzzle Width	101	Udder Depth	109	
Body Depth	98	Fore Attachment	107	
Chest Width	105	Rear Att Height	106	
Loin Strength	102	Rear Att Width	101	
Foot Angle	105	Centre Ligament	101	
Rear Set	101	Teat Placement (Front)	106	
Rear Leg Rear View	103	Teat Placement (Rear)	102	
		Teat Length	101	

- First daughters hitting dairies now
- 482 BPI 519 HWI A2A2

BEEZER T-Spruce Peak Beezer (Olympus / Taos / Lionel)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXBEEZER NAAB: 001H017148 HB: H0840M003272622397 DOB: 28/08/2023

NEW GENEX™

RRP \$25.00

USDA CDCB Genomic Index						August 2025	
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NMS
3338	1672	106	0.13	66	0.04	80	843
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0.1	-1.8	2.5	0.6	-1.1	2.96	0.88	0.62



T-Spruce Peak Beezer

Australian Indices								August 2025			
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%		
371	325	643	218	266	44	0.46%	22	0.28%	76		
Dtr Fert		LiveW		SCC		M/Speed		Temp		Likeability	
107		100		166		102		100		103	
C Ease		Gest L		Feed S		Survival		Heat Tol		TYPE MAMM	
104		-5		-138		105		89		101 104	

- Calving ease specialist 104, with a short gestation length of -5 days
- Outcross pedigree with extreme milk flow

CONFORMATION		0 Dtrs	0 Herds	USDA/HA Genomic Evaluation			August 2025	
PTAT: 0.88		UDC: 0.62	FLC: -0.1			GTPI 3338		
Stature	Short						Tall	0.84
Strength	Frail						Strong	0.53
Body Depth	Shallow						Deep	0.62
Dairy Form	Tight Rib						Open Rib	1.35
Rump Angle	High						Sloped	1.04
Rump width	Narrow						Wide	1.38
RL SV	Posty						Sickle	0.35
RL RV	Hock-In						Straight	0.00
Foot Angle	Low						Steep	-0.28
F & L Score	Low						High	0.14
Fore Attachment	Loose						Strong	0.68
Rear Udder Height	Low						High	1.02
Rear Udder Width	Narrow						Wide	1.80
Udder Cleft	Weak						Strong	0.29
Udder Depth	Deep						Shallow	-0.17
Fore Teat Placement	Wide						Close	0.47
Rear Teat Placement	Wide						Close	0.60
Teat Length	Short						Long	0.45
		-1.5	-1.0	-0.5	0	0.5	1.0	1.5

LORAX

Peak Lorax (Lockstep / Upside / AltaZazzle)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXLORAX NAAB: 001H017248 HB: H0840M003275362776 DOB: 11/03/2023



RRP \$26.00

USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3331	913	93	0.20	44	0.05	79	980
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
2.0	0.0	5.4	1.1	-0.5	2.66	0.1	0.24



Peak Lorax

Australian Indices								August 2025	
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
458	477	766	188	289	46	0.48%	17	0.18%	76
Dtr Fert		LiveW		SCC		M/Speed		Temp	
115		99		182		101		100	
C Ease		Gest L		Feed S		Survival		Heat Tol	
103		-3		-105		110		102	
								TYPE	
								MAMM	
								96	
								100	

- +2 CCR, +5.4 Productive Life with low cell count
- Combining High TPI, NM\$ and High Milk Flow

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation						August 2025	
PTAT: 0.10 UDC: 0.24 FLC: -0.16						GTP! 3331	
Stature	Short					Tall	-0.22
Strength	Frail					Strong	-0.84
Body Depth	Shallow					Deep	-0.73
Dairy Form	Tight Rib					Open Rib	0.28
Rump Angle	High					Sloped	0.68
Rump width	Narrow					Wide	-0.22
RL SV	Posty					Sickle	-0.31
RL RV	Hock-In					Straight	-0.47
Foot Angle	Low					Steep	-0.08
F & L Score	Low					High	-0.12
Fore Attachment	Loose					Strong	0.39
Rear Udder Height	Low					High	0.23
Rear Udder Width	Narrow					Wide	0.36
Udder Cleft	Weak					Strong	-0.27
Udder Depth	Deep					Shallow	0.24
Fore Teat Placement	Wide					Close	-0.59
Rear Teat Placement	Wide					Close	-0.41
Teat Length	Short					Long	0.08

OVERHAUL-P Stantons Overhaul-P (Revamp-P /Delta Lambda / Bighit-P)

NEW

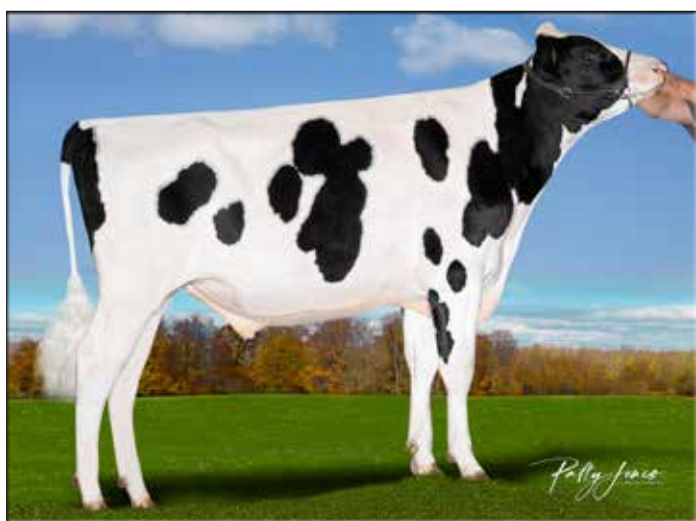
A2 STATUS: A2/A2 Genetic Codes: TE, P Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGOVERHAUL NAAB: 724H002037 HB: HOCAN000014691536 DOB: 18/10/2023

SEXED \$65.00 RRP \$35.00
Ultraplus[®] CONVENTIONAL PLUS20

USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NMS
3003	471	32	0.05	26	0.04	79	420
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
2.2	1.5	3.1	1.1	0.8	2.96	1.25	1.61

- Dam is #1 LPI cow in Canada
- #1 P or PP sire in Canada
- +1.5 DPR +3003 GTPI +1.61 UDC
- +4034 LPI, +11 Conformation, +11 Feet&Legs
- One not to be missed

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation										August 2025
PTAT: 1.25 UDC: 1.61 FLC: 0.81										GTPI 3003
Stature	Short								Tall	0.34
Strength	Frail								Strong	0.40
Body Depth	Shallow								Deep	0.28
Dairy Form	Tight Rib								Open Rib	0.04
Rump Angle	High								Sloped	0.06
Rump width	Narrow								Wide	0.46
RL SV	Posty								Sickle	0.46
RL RV	Hock-In								Straight	0.73
Foot Angle	Low								Steep	1.12
F & L Score	Low								High	0.83
Fore Attachment	Loose								Strong	2.47
Rear Udder Height	Low								High	1.70
Rear Udder Width	Narrow								Wide	1.13
Udder Cleft	Weak								Strong	0.56
Udder Depth	Deep								Shallow	2.02
Fore Teat Placement	Wide								Close	0.23
Rear Teat Placement	Wide								Close	0.35
Teat Length	Short								Long	-0.43



Stantons Overhaul-P



Dam: Stantons Lambda Margaret VG 2yo

BPI 405 64 %R

HWI 403 59 %R

SI 742 62 %R

PRODUCTION ABV (g) 08/25

Milk	353	76% R
Fat	37 kg	0.31%
Protein	19 kg	0.18%
ASI	177	76 % R

WORKABILITIES

Milking Speed	101
Temperament	101
Likability	105

Conformation ABV(g)

Overall Type	101
Mammary	105
Rump	99
Dairy Strength	100
Feet and Legs	99

HEALTH ABV (g) 08/25

Cell Count	152	66
Survival	110	56
Feed Saved	-76	42
Dtr Fertility	110	56
Calving Ease	103	66
Gest Length	-3	61
Heat Tol	102	48

CONFORMATION ABV (g) 0 Dtrs 0 Herds 61%Rel August 2025

OVERALL TYPE	101	MAMMARY SYSTEM	105
Stature	104	Pin Width	100
Bone Quality	102	Pin Set	96
Angularity	98	Udder Texture	99
Muzzle Width	100	Udder Depth	107
Body Depth	100	Fore Attachment	106
Chest Width	102	Rear Att Height	104
Loin Strength	98	Rear Att Width	101
Foot Angle	99	Centre Ligament	103
Rear Set	104	Teat Placement (Front)	102
Rear Leg Rear View	99	Teat Placement (Rear)	107
		Teat Length	97

Have you booked your **FRESH SEXED SEMEN** dates for your Spring 2025 joining programs?



Now is the ideal time to confirm your fresh sexed semen booking, so contact TLG today

Consider the advantages

- *Success stories continue to be told*
- *We manage the logistics so it is delivered on time*
- *Easily managed synchro programs*
- *Wide selection of quality Australian standing bulls*
- *Consistently delivering improved fertility*

For the last 3 years, throughout October and November we aim to offer fresh sexed semen to our clients daily. Working closely with TLG they have made this process seamless to provide commercial, high health and low stature bulls for the Maffra market and work out the logistics for smooth delivery.

Results are consistently within a couple of % of conventional semen, in some cases the fresh sexed has come out higher. Our clients love the ease and flexibility of the fresh sexed product from TLG.

Lyndsey Flemming, Quality Breeding Services, Gippsland

We have been using fresh sexed semen since it became available in Australia. We and our clients find it easy to use and enjoy the consistency of the conception rates.

We find that the conception rates are an improvement compared to using frozen product.

Jonathan Thomas, Nth East Genetics

THE DEMAND FOR FRESH SEXED IS INCREASING *and so are the success stories...*



LIONIZE- pp Summit View Nipit Lionize - PP (Nipit-P / Luster-P / Bighit-P)

A2 STATUS: A2/A2 Genetic Codes: TE, PP Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGLIONIZE NAAB: 187H005667 HB: AUS 2199626 DOB: 30/05/2022

SEXED \$55.00 RRP \$24.00

USDA CDCB Genomic Index						August 2025	
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
2787	63	45	0.16	28	0.1	82	108
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0.3	-0.3	-1.6	1.9	1	3.16	1.52	0.66



Summit View Nipit Lionize-PP

Australian Indices								August 2025	
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
343	287	609	207	183	40	0.46%	21	0.30%	78
Dtr Fert		LiveW	SCC	M/Speed	Temp	Likeability			
107		102	146	102	101	103			
C Ease		Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM		
101		-1	-158	107	96	101	100		

- High ranking PP sire with +1.52 for Type
- Combining Polled, Milk, Type & Udders

CONFORMATION					0 Dtrs	0 Herds	USDA/HA Genomic Evaluation		August 2025		
PTAT: 1.52		UDC: 0.66		FLC: 0.17		GTPI 2787					
Stature	Short							Tall	2.38		
Strength	Frail							Strong	1.34		
Body Depth	Shallow							Deep	1.69		
Dairy Form	Tight Rib							Open Rib	1.67		
Rump Angle	High							Sloped	0.68		
Rump width	Narrow							Wide	1.69		
RL SV	Posty							Sickle	0.99		
RL RV	Hock-In							Straight	0.54		
Foot Angle	Low							Steep	0.67		
F & L Score	Low							High	0.73		
Fore Attachment	Loose							Strong	1.15		
Rear Udder Height	Low							High	0.96		
Rear Udder Width	Narrow							Wide	1.24		
Udder Cleft	Weak							Strong	1.33		
Udder Depth	Deep							Shallow	1.47		
Fore Teat Placement	Wide							Close	0.49		
Rear Teat Placement	Wide							Close	0.70		
Teat Length	Short							Long	0.92		
					-1.5	-1.0	-0.5	0	0.5	1.0	1.5

LEX-pp

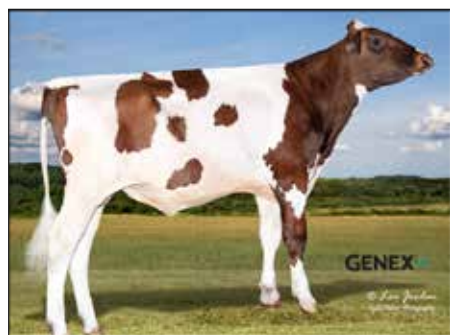
LEX-pp Siemens Lex PP Red (Lazer PP / McDonald P / Luster P)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXLEX NAAB: 001H017352 HB: H0840M003267429178 DOB: 09/08/2023

GENEX 

SEXED \$55.00 RRP \$26.00

USDA CDCB Genomic Index						August 2025	
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
2745	-137	29	0.13	-3	0.01	80	131
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
-1.5	-0.7	0.9	1.7	-0.5	2.94	2.22	1.4



Siemers Lex PP RED

Australian Indices								August 2025	
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
245	238	305	60	43	29	0.39%	1	-0.01%	76
Dtr Fert		LiveW	SCC	M/Speed		Temp	Likeability		
106		103	151	103		102	104		
C Ease		Gest L	Feed S	Survival		Heat Tol	TYPE	MAMM	
102		-4	-158	108		102	103	107	

- Semen fertility of +1.1 SCR
- +2.22 PTAT polled red sire

CONFORMATION			0 Dtrs	0 Herds	USDA/HA Genomic Evaluation		August 2025	
PTAT: 2.22	UDC: 1.40	FLC: 1.38					GTPI 2745	
Stature	Short						Tall	2.51
Strength	Frail						Strong	0.20
Body Depth	Shallow						Deep	0.99
Dairy Form	Tight Rib						Open Rib	1.96
Rump Angle	High						Sloped	-0.70
Rump width	Narrow						Wide	1.70
RL SV	Posty						Sickle	0.56
RL RV	Hock-In						Straight	1.85
Foot Angle	Low						Steep	2.15
F & L Score	Low						High	1.78
Fore Attachment	Loose						Strong	2.36
Rear Udder Height	Low						High	1.85
Rear Udder Width	Narrow						Wide	1.34
Udder Cleft	Weak						Strong	1.37
Udder Depth	Deep						Shallow	2.23
Fore Teat Placement	Wide						Close	1.80
Rear Teat Placement	Wide						Close	1.76
Teat Length	Short						Long	0.29

POLLED SPRING SPECIAL

Buy 100 doses or more of any combination
of conventional semen below for \$18.00 a dose

Buy 50 doses or more of any sexed semen below and
save 10% OFF RRP

LEX-PP-RED

\$26.00 | \$55.00 (SEXED)

BREED DEFINING COW FAMILY THAT CONTINUES TO DELIVER.

MDOUGLAS-PP

\$22.00 | \$55.00 (SEXED)

ADDS CHEST AND MUZZLE TO A MODERATE STATURE RATING.

BEACON-PP

\$20.00 | \$50.00 (SEXED)

DAUGHTER FERTILITY SPECIALIST, WITH ADDED TEAT LENGTH.

GONE-PP

\$26.00

OUTCROSS SIRE WITH LOADS OF MILK.

MDOUGLAS-pp Vala Fomo MDouglas - PP (Fomo / Bighit / Montana)

A2 STATUS: A2/A2 Genetic Codes: TE, PP Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGMDOUGLAS NAAB: 187H005637 HB: AUS2191761 DOB: 04/02/2022

SEXED \$55.00 RRP \$22.00

Ultraplus™

CONVENTIONAL PLUS20

USDA CDCB Genomic Index

August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
2972	-114	52	0.22	17	0.08	82	415

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
1	0	2	1.7	0.2	2.98	0.88	1.86



Vala Fomo MDouglas - PP

Australian Indices

August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
412	405	666	163	51	35	0.48%	14	0.24%	77

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability	
111	100	161	104	100	103	
C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
101	-2	-111	109	104	98	103

PP sire with +2972 GTPI and 412 BPI, +1.83 UDC

International sire of sons

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation

August 2025

PTAT: 0.88 UDC: 1.86 FLC: 0.49 GTPI 2972

Stature	Short						Tall	-0.34
Strength	Frail						Strong	0.09
Body Depth	Shallow						Deep	-0.33
Dairy Form	Tight Rib						Open Rib	-0.76
Rump Angle	High						Sloped	-0.09
Rump width	Narrow						Wide	1.06
RL SV	Posty						Sickle	0.10
RL RV	Hock-In						Straight	0.28
Foot Angle	Low						Steep	0.42
F & L Score	Low						High	0.41
Fore Attachment	Loose						Strong	2.28
Rear Udder Height	Low						High	1.94
Rear Udder Width	Narrow						Wide	1.39
Udder Cleft	Weak						Strong	0.41
Udder Depth	Deep						Shallow	2.16
Fore Teat Placement	Wide						Close	0.26
Rear Teat Placement	Wide						Close	0.44
Teat Length	Short						Long	-0.76

-1.5 -1.0 -0.5 0 0.5 1.0 1.5

BEACON - pp

Vala Magicball Beacon - PP (Magicball / Jeronimo / Bandares)

A2 STATUS: A2/A2 Genetic Codes: TE, PP Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGBEACON NAAB: 187H005636 HB: AUS2192579 DOB: 02/04/2022

SEXED \$50.00

RRP \$20.00

UltraPlus

CONVENTIONAL PLUS26



Vala Magicball Beacon-PP

PRODUCTION ABV (g) 08/25		
Milk	-651	78% R
Fat	32 kg	0.86%
Protein	5 kg	0.42%
ASI	152	78 % R

WORKABILITIES	
Milking Speed	101
Temperament	102
Likability	102

BPI 412 68 %R

HWI 445 63 %R

SI 616 65 %R

Conformation ABV(g)	
Overall Type	100
Mammary	99
Rump	103
Dairy Strength	94
Feet and Legs	102

HEALTH ABV (g) 08/25		
Cell Count	175	69
Survival	109	60
Feed Saved	15	44
Dtr Fertility	114	66
Calving Ease	102	87
Gest Length	-3	87
Heat Tol	102	48

CONFORMATION ABV (g)	0 Dtrs	0 Herds	64%Rel	August 2025
OVERALL TYPE	100	MAMMARY SYSTEM		99
Stature	103	Pin Width		96
Bone Quality	100	Pin Set		101
Angularity	94	Udder Texture		94
Muzzle Width	98	Udder Depth		106
Body Depth	94	Fore Attachment		102
Chest Width	96	Rear Att Height		105
Loin Strength	104	Rear Att Width		96
Foot Angle	99	Centre Ligament		95
Rear Set	96	Teat Placement (Front)		98
Rear Leg Rear View	100	Teat Placement (Rear)		96
		Teat Length		103

- Homozygous Polled sire that widens and lengthens rear teats
- High Fertility sire with calving ease of 102 and short gestation -3

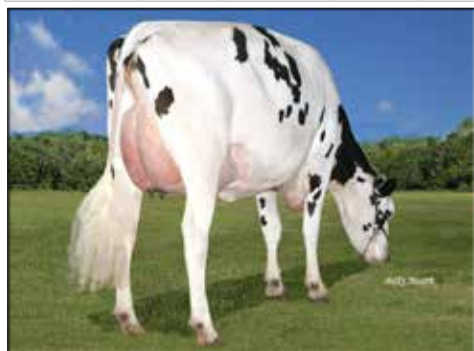
GONE - pp

Coldsprings All-Gone-PP (Allday-P / Basic P / Denver)

A2 STATUS: A2/A2 Genetic Codes: TE, PP Haplotypes: HH1F,HH2F,HH3C,HH4F,HH5F,HH6F
NASIS: GONEPP NAAB: 724H002025 HB: H08400003210109717 DOB: 18/06/2021

RRP \$22.00

USDA CDCB Genomic Index								August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$	
2718	198	18	0.04	3	-0.01	82	85	
CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC	
0.4	0.2	-0.2	1.3	1.4	2.84	1.44	1.54	



Cold Springs Basic 114477

Australian Indices										August 2025
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%	
307	329	482	86	411	29	0.17%	9	-0.05%	78	
Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability					
108	101	172	100	98	100					
C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM				
103	1	-61	106	105	106	108				

- Imagine the udders on this PP sire
- Milk, calving ease, type and production in one

CONFORMATION	0 Dtrs	0 Herds	USDA/HA Genomic Evaluation	August 2025
PTAT: 1.44	UDC: 1.54	FLC: 0.74	GTPI 2718	
Stature	Short			Tall 1.78
Strength	Frail			Strong 0.73
Body Depth	Shallow			Deep 0.71
Dairy Form	Tight Rib			Open Rib 0.62
Rump Angle	High			Sloped -0.32
Rump width	Narrow			Wide 1.20
RL SV	Posty			Sickle 0.26
RL RV	Hock-In			Straight 0.76
Foot Angle	Low			Steep 0.83
F & L Score	Low			High 1.16
Fore Attachment	Loose			Strong 1.73
Rear Udder Height	Low			High 2.32
Rear Udder Width	Narrow			Wide 1.82
Udder Cleft	Weak			Strong 1.16
Udder Depth	Deep			Shallow 1.91
Fore Teat Placement	Wide			Close 0.64
Rear Teat Placement	Wide			Close 1.14
Teat Length	Short			Long 0.33

RIZBOY

Emu Banks Rizboy (Manhattan / Montana / Supershot)

A2 STATUS: A1/A2 Genetic Codes: MW Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGRIZBOY NAAB: 187H005641 HB: AUS2181357 DOB: 30/11/2021

SEXED \$50.00 RRP \$22.00
Ultraplus

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
2933	-94	39	0.16	23	0.1	82	275

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
2.4	1.6	2.1	1.6	-0.8	2.8	1.1	0.71



Dam: Westcoast Montana Riza 4700

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
519	558	772	153	-365	31	0.67%	9	0.35%	77

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability	
116	102	185	103	103	104	
C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
101	-2	-89	113	96	108	104

○ Australian standing sire combining high TPI and BPI

○ +2933 GTPI, 519 BPI, 1.1 PTAT, +2.4 CCR

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.1 UDC: 0.71 FLC: -0.21 GTPI 2933

Stature	Short	Tall	1.53
Strength	Frail	Strong	1.10
Body Depth	Shallow	Deep	0.76
Dairy Form	Tight Rib	Open Rib	00.0
Rump Angle	High	Sloped	1.02
Rump width	Narrow	Wide	1.47
RL SV	Posty	Sickle	-0.70
RL RV	Hock-In	Straight	-0.24
Foot Angle	Low	Steep	0.77
F & L Score	Low	High	0.21
Fore Attachment	Loose	Strong	1.70
Rear Udder Height	Low	High	0.82
Rear Udder Width	Narrow	Wide	1.05
Udder Cleft	Weak	Strong	-0.11
Udder Depth	Deep	Shallow	1.69
Fore Teat Placement	Wide	Close	-0.31
Rear Teat Placement	Wide	Close	-0.64
Teat Length	Short	Long	0.44

POCALYPSE

Peak Bullpocalypse (AltaGadzook x No Excuse x Redrock)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXPOCALYPSE NAAB: 001H016649 HB: H084000003247843699 DOB: 24/05/2022

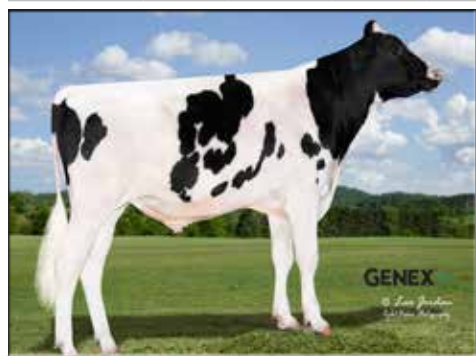
SEXED \$48.00 RRP \$20.00
Ultraplus

GENEX

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3176	716	66	0.13	43	0.07	80	625

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
+1.3	-0.3	2.6	1.5	-1.0	2.77	1.03	1.03



Peak Bullpocalypse

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
475	423	848	240	189	57	0.70	20	0.29	77

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability	
107	100	180	101	101	103	
C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
103	-4	-91	111	97	94	99

○ 3176 GTPI with sloped rumps and teat length

○ +1.3 CCR, Short gestation, 475 BPI, A2A2

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.03 UDC: 1.03 FLC: -0.6 GTPI 3176

Stature	Short	Tall	1.30
Strength	Frail	Strong	0.43
Body Depth	Shallow	Deep	0.54
Dairy Form	Tight Rib	Open Rib	0.93
Rump Angle	High	Sloped	0.40
Rump width	Narrow	Wide	1.32
RL SV	Posty	Sickle	0.24
RL RV	Hock-In	Straight	-0.60
Foot Angle	Low	Steep	0.37
F & L Score	Low	High	-0.23
Fore Attachment	Loose	Strong	1.67
Rear Udder Height	Low	High	1.25
Rear Udder Width	Narrow	Wide	1.41
Udder Cleft	Weak	Strong	0.60
Udder Depth	Deep	Shallow	1.27
Fore Teat Placement	Wide	Close	0.41
Rear Teat Placement	Wide	Close	0.37
Teat Length	Short	Long	0.16

LEITH-P

Vala Nacash Leith - P (Nacash / Polar P / Duke)

A2 STATUS: A2/A2 Genetic Codes: TE, HP Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGLEITH NAAB: 187H005560 HB: AUS2146004 DOB: 05/03/2021

RRP \$18.00

CONVENTIONAL PLUS20

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NMS
2622	-148	15	0.08	6	0.04	84	27

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
0.5	0.8	0.5	1.9	0.8	2.89	0.72	0.43



Vala Nacash Leith-P

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
279	307	398	70	-22	12	0.19%	6	0.13%	93

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
112	100	164	102	101	103

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
101	0	-155	109	102	101	100

Combining BPI, TPI, fertility, A2A2 and polled

Predominately black with added teat length

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 0.72 UDC: 0.43 FLC: -0.03 GTPI 2622

Stature	Short					Tall	0.94
Strength	Frail					Strong	0.65
Body Depth	Shallow					Deep	0.40
Dairy Form	Tight Rib					Open Rib	-0.43
Rump Angle	High					Sloped	0.84
Rump width	Narrow					Wide	0.92
RL SV	Posty					Sickle	0.89
RL RV	Hock-In					Straight	-0.27
Foot Angle	Low					Steep	0.17
F & L Score	Low					High	0.32
Fore Attachment	Loose					Strong	1.39
Rear Udder Height	Low					High	0.14
Rear Udder Width	Narrow					Wide	0.17
Udder Cleft	Weak					Strong	0.09
Udder Depth	Deep					Shallow	1.36
Fore Teat Placement	Wide					Close	0.64
Rear Teat Placement	Wide					Close	0.30
Teat Length	Short					Long	0.93

SHERIFF

Vala Bolti Sheriff (Bolti / Almamater / King Royal)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGSHERIFF NAAB: 187H005712 HB: AUS2227506 DOB: 12/03/2023

SEXED \$48.00

RRP \$20.00

Ultraplus

CONVENTIONAL PLUS20

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NMS
2743	-428	1	0.07	1	0.06	78	149

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
3	2.1	2.5	1.3	0.6	2.78	1.03	1.46



Vala Bolti Sheriff

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
505	550	792	139	-223	21	0.45%	11	0.33%	78

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
116	102	180	105	103	105

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
103	0	-69	113	100	102	105

Among the leading bulls available for longevity

+1.46 UDC and +105 Mammary with +2.1 DPR

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.03 UDC: 1.46 FLC: -0.06 GTPI 2743

Stature	Short					Tall	0.94
Strength	Frail					Strong	0.28
Body Depth	Shallow					Deep	-0.09
Dairy Form	Tight Rib					Open Rib	-0.23
Rump Angle	High					Sloped	1.96
Rump width	Narrow					Wide	0.69
RL SV	Posty					Sickle	-0.75
RL RV	Hock-In					Straight	-0.14
Foot Angle	Low					Steep	0.92
F & L Score	Low					High	0.19
Fore Attachment	Loose					Strong	1.74
Rear Udder Height	Low					High	1.40
Rear Udder Width	Narrow					Wide	1.17
Udder Cleft	Weak					Strong	1.25
Udder Depth	Deep					Shallow	2.22
Fore Teat Placement	Wide					Close	1.21
Rear Teat Placement	Wide					Close	1.13
Teat Length	Short					Long	-0.30

ZIMMER

Peak Zimmer (Jalapeno / Incredible / Pursuit)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXZIMMER NAAB: 001H016646 HB: H0840M003247843570 DOB: 25/04/2022

GENEX

SEXED \$48.00

RRP \$18.00

Ultraplus

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NMS
3080	1109	49	0.01	49	0.05	81	697

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
-0.8	-1.5	3	1.1	-1.3	2.74	0.77	0.7



Peak Zimmer

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
375	366	705	160	282	32	0.29%	17	0.18%	77

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
106	98	165	101	102	104

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
104	-3	-24	110	104	101	103

○ Outcross pedigree with nice linear profile

○ +1109 Milk, +98 CFP, +3080 GTPI

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 0.77 UDC: 0.7 FLC: -0.67 GTPI 3080

Stature	Short						Tall	0.87
Strength	Frail						Strong	-0.28
Body Depth	Shallow						Deep	0.54
Dairy Form	Tight Rib						Open Rib	2.74
Rump Angle	High						Sloped	1.50
Rump width	Narrow						Wide	1.55
RL SV	Posty						Sickle	0.75
RL RV	Hock-In						Straight	-1.01
Foot Angle	Low						Steep	-0.06
F & L Score	Low						High	-0.27
Fore Attachment	Loose						Strong	0.47
Rear Udder Height	Low						High	1.22
Rear Udder Width	Narrow						Wide	1.98
Udder Cleft	Weak						Strong	0.64
Udder Depth	Deep						Shallow	-0.06
Fore Teat Placement	Wide						Close	0.50
Rear Teat Placement	Wide						Close	0.59
Teat Length	Short						Long	0.81

MAUNO

Peak Mauno (Zemini / Pursuit / Medley)

A2 STATUS: A2/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: GXMAUNO NAAB: 001H016397 HB: H0840M003247843563 DOB: 24/04/2022

GENEX

RRP \$22.00

USDA CDCB Genomic Index August 2025

GTPI	MILK	FAT	%F	PROT	%P	Rel%	NMS
3274	1435	67	0.03	51	0.01	82	834

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
1	-0.2	4.5	1	-2.4	2.7	0.61	0.71



Peak Mauno

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
395	359	745	197	567	41	0.24%	23	0.15%	77

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
107	100	168	101	100	103

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
105	-6	-113	109	100	101	102

○ +3274 GTPI and 1% Calving Ease

○ Combining Milk, Udders & Fertility, A2A2

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 0.61 UDC: 0.71 FLC: 0.64 GTPI 3274

Stature	Short						Tall	0.22
Strength	Frail						Strong	-0.04
Body Depth	Shallow						Deep	-0.29
Dairy Form	Tight Rib						Open Rib	0.20
Rump Angle	High						Sloped	0.64
Rump width	Narrow						Wide	0.75
RL SV	Posty						Sickle	-0.57
RL RV	Hock-In						Straight	0.68
Foot Angle	Low						Steep	0.60
F & L Score	Low						High	0.64
Fore Attachment	Loose						Strong	0.78
Rear Udder Height	Low						High	0.75
Rear Udder Width	Narrow						Wide	1.18
Udder Cleft	Weak						Strong	-0.18
Udder Depth	Deep						Shallow	0.51
Fore Teat Placement	Wide						Close	0.77
Rear Teat Placement	Wide						Close	0.51
Teat Length	Short						Long	-0.14

Coming Soon

We have some exciting sires soon to be available in the
COMING MONTHS



View Fort

HUCKLEBERRY

(Pace x Doc)

- *Pace brother to Yarrawood*
- *Huckleberry is a very impressive individual*
- *+2.51 PTAT +1.71 UDC +1.47 F&L*
- *+211 lbs Milk 1.7% Calving ease*

RRP \$32.00

Peak

EXPEDIA

(Excitement x Zappy)

- *Extreme TPI sire 3473 GTPI*
- *+1.73 PTAT with ideal linear profile*
- *1205 lbs milk 174kg CFP*
- *Sloped rumps and adds teat length*

RRP \$45.00 SEXED \$70.00

H-Bridge Vec

PAPRIKA-PP-RED

(Eveready-PP x Augustus-P-Red)

- *+2.08 UDC +1.51 PTAT*
- *Outcross sire line*
- *+0.7 DPR +1.0 CCR*
- *Combining Type, TPI, Polled and Fertility*

P.O.A

Peak

LEXION

(Axford x Merle)

- *3463 GTPI +968 NM\$*
- *+0.1 DPR +1.8 CCR*
- *+839 Milk +158 CFP*
- *557 BPI 275 ASI*

SEXED \$85.00

ELEVATION

Summit View RocknRoll Elevation (RocknRoll / Einstein / Marius)

A2 STATUS: A1/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGELEVATION NAAB: 187H05710 HB: AUS2229105 DOB: 04/05/2023

RRP \$18.00
CONVENTIONAL PLUS20

USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
3084	547	62	0.15	33	0.06	80	496

CCR	DPR	PL	C/EASE	GEST L	SCS	PTAT	UDC
1.7	1	3	1.6	-1.4	2.79	0.51	0.77



Summit View Rocknroll Elevation

Australian Indices										August 2025
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%	
500	511	870	213	284	46	0.49%	21	0.25%	78	

Dtr Fert	LiveW	SCC	M/Speed	Temp	Likeability
114	101	179	102	101	104

C Ease	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
102	-6	-71	111	97	102	103

○ Fertility and milk quality improver

○ 3084 GTPI, 500 BPI, 511 HWI

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation										August 2025
PTAT: 0.51 UDC: 0.77 FLC: -0.18										GTPI 3084
Stature	Short								Tall	0.08
Strength	Frail								Strong	1.29
Body Depth	Shallow								Deep	0.28
Dairy Form	Tight Rib								Open Rib	-1.31
Rump Angle	High								Sloped	-0.32
Rump width	Narrow								Wide	0.66
RL SV	Posty								Sickle	-0.78
RL RV	Hock-In								Straight	-0.71
Foot Angle	Low								Steep	0.67
F & L Score	Low								High	-0.02
Fore Attachment	Loose								Strong	1.16
Rear Udder Height	Low								High	0.10
Rear Udder Width	Narrow								Wide	0.73
Udder Cleft	Weak								Strong	0.94
Udder Depth	Deep								Shallow	0.91
Fore Teat Placement	Wide								Close	1.04
Rear Teat Placement	Wide								Close	1.40
Teat Length	Short								Long	-0.25

TURBO

Wilara Turbo (Endgame / Delta Lambda / Doorman)

A2 STATUS: A1/A2 Genetic Codes: TE Haplotypes: HH1F,HH2F,HH3F,HH4F,HH5F,HH6F
NASIS: TLGTURBO HB: AUS2133034 DOB: 16/10/2020

SEXED \$48.00 RRP \$18.00
UltraPlus CONVENTIONAL PLUS20



Same family as Gorbro Atwood Tiffany EX91-1E

PRODUCTION ABV (g) 08/25		
Milk	137	77% R
Fat	24 kg	0.27%
Protein	21 kg	0.33%
ASI	183	77% R

WORKABILITIES	
Milking Speed	103
Temperament	103
Likability	105

BPI 446 68 %R

HWI 428 64 %R

SI 771 65 %R

Conformation ABV(g)	
Overall Type	106
Mammary	105
Rump	108
Dairy Strength	99
Feet and Legs	107

HEALTH ABV (g) 08/25	
Cell Count	171 68
Survival	110 63
Feed Saved	-129 44
Dtr Fertility	110 66
Calving Ease	101 93
Gest Length	-1 94
Heat Tol	98 48

CONFORMATION ABV (g) 0 Dtrs 0 Herds 66%Rel	August 2025	
OVERALL TYPE	106	MAMMARY SYSTEM 105
Stature	104	Pin Width 107
Bone Quality	103	Pin Set 109
Angularity	95	Udder Texture 100
Muzzle Width	99	Udder Depth 104
Body Depth	99	Fore Attachment 101
Chest Width	103	Rear Att Height 111
Loin Strength	102	Rear Att Width 104
Foot Angle	107	Centre Ligament 101
Rear Set	95	Teat Placement (Front) 97
Rear Leg Rear View	105	Teat Placement (Rear) 101
		Teat Length 96



Wilara Turbo

○ Solid Type and production from famous Tiffanys

○ High daughter fertility 110

Are you looking for **VALUE OPTIONS?**

Holstein Value Sires

TLGLENN - P RDC	\$14.00 - A2A2	GXBACKSPIN	\$16.00 - A2A2
TLGBEAMER - P	\$14.00 - A2A2	ESCHER - PP	\$16.00 - A1A2
TLGBRINX - P	\$14.00 - A1A2	TLGLEITH - P	\$16.00 - A2A2
TLGLARK	\$14.00 - A2A2	TLGWIM	\$16.00 - A2A2
TLGEPIC	\$16.00 - A2A2	TLGEMMANUEL	\$16.00 - A2A2
EMMETT - P	\$16.00 - A2A2	TLGCHILLI - PP	\$16.00 - A2A2

Jersey Value Sires

TLGVALBONE	\$12.00 - A1A2
TLGDEPOLE - P	\$14.00 - A1A2
TLGQANTAS	\$14.00 - A2A2
GXGRAYSON - PP	\$16.00 - A2A2

**TALK TO US ABOUT VOLUME
DISCOUNTS ON THIS WIDE
RANGE OF VALUE SIRES...**

Talk to us today for all your
**GENOMIC
TESTING**
REQUIREMENTS

All major testing services available including:

- BPI, TPI, JPI & LPI Bases
- A2 & Full Milk Proteins Testing
- Recumbency and Haplotypes
- Polled, JH1, BVD Testing

Backed with assistance for sampling on farm,
submission support and genetic decisions.



One sample, one simple
comprehensive solution.



Our Genomic Services team are available to
assist you with all your enquires
by phoning (03) 5593 2016 or
genomics@genaust.com.au

TOTAL
LIVESTOCK GENETICS

GENEX
HerdMonitor®
Powered by  **nedap**



Transform your future.

The GENEX HerdMonitor™ cow monitoring system will revolutionize your approach to cow management, by bringing you real-time solutions today for a successful tomorrow. This system features 24/7 monitoring, with identification, rumination, heat detection and cow positioning options, all designed to take your herd management to the next level.



Unmatched monitoring features and management tools in a highly advanced, yet easy to use system.



Integration & Connection

Easily connect HerdMonitor to your farm automation systems and dairy management programs thanks to advanced integration applications.



Heat Detection

Highly accurate heat detection with advice for the optimal insemination moment and reproduction insights.



Health Monitoring

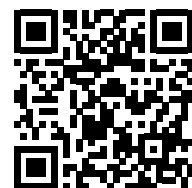
Individual health management for early detection of health issues and intensive monitoring of transition cows and post-treatment recovery.



Herd Performance Trends

Chart the behavior patterns of groups and your entire herd. Get alerts when possible risks affect a group's performance. Evaluate the impact of management decisions on herd performance to make improvements.

Contact your Genetics Australia or Total Livestock Genetics Representative today. For further information scan the QR code or visit genaust.com.au/herdmonitor.



TRACTION

Hilltop Acres Traction (Telsa / Kingsley / Doboy)

GENEX

NEW

A2 STATUS: A1/A2 Haplotypes: BH2F

NASIS: GXTRACTION NAAB: 001BS00720 HB: BSS840M003273133744 DOB: 21/03/2024

RRP \$26.00



Hilltop Acres Traction

- Exciting New Brown Swiss Sire
- Moderate frames with great udders
- +0.90 PTAT with wide sloped rumps
- High milk flow with positive components

USDA CDCB Genomic Index							August 2025
GTPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
151	1138	57	0.04	48	0.04	64	555
DPR	PL	SCS	GEST L	HCR	CCR	PTAT	JUI
-0.8	3.4	2.81	0	0.0	-2.0	0.90	0.59

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation										August 2025
PTAT: 0.90 UDC: 0.59										GTPI 151
Stature	Short								Tall	-0.90
Strength	Frail								Strong	+0.90
Dairy Form	Tight Rib								Open Rib	+1.00
Body Depth	Shallow								Deep	+0.30
Rump Angle	High								Sloped	+0.60
Rump Width	Narrow								Wide	+0.70
RL SV	Posty								Sickle	-0.40
Foot Angle	Hock-In								Straight	+0.10
Fore Attachment	Loose								Strong	+1.60
Rear Udder Height	Low								High	+0.70
Rear Udder Width	Narrow								Wide	+0.70
Udder Cleft	Weak								Strong	+0.10
Udder Depth	Deep								Shallow	-0.20
Fore Teat Placement	Wide								Close	+0.10
Teat Length	Short								Long	+0.20



GDam: Hilltop Acres DB Tabby EX91



GDam: Hilltop Acres DB Tabby EX91



Dam: Hilltop Acres K Tammie EX92 EX94 MS



Dam: Hilltop Acres K Tammie EX92 EX94 MS

DINGO

Windy Ways CCC Dingo (Roulette / Galaxies EX95 / Headline EX90)

A2 STATUS: A2/A2 Haplotypes: JH1F
NASIS: TLGDINGO HB: AUS814027 DOB: 02/03/2023

SEXED \$55.00 RRP \$24.00
Ultraplus[®] CONVENTIONAL PLUS20

PRODUCTION ABV 08/25		
Milk	427	79% R
Fat	15 kg	-0.16%
Protein	14 kg	-0.01%
ASI	91	70% R

WORKABILITIES	
Milking Speed	101
Temperament	104
Likability	105

BPI	242	70%R
HWI	196	58%R
SI	487	63%R

Conformation ABV(g)	
Overall Type	106
Mammary	110

HEALTH ABV (g) 08/25		
Cell Count	131	70%
Survival	107	56%
Feed Saved	-39	34%
Dtr Fertility	99	55%
Heat Tol	99	39%

CONFORMATION ABV (g)	0 Dtrs	0 Herds	63% Rel	August 2025
OVERALL TYPE	106	MAMMARY SYSTEM	110	
Stature	105	Pin Width	101	
Bone Quality	107	Pin Set	100	
Angularity	104	Udder Texture	105	
Muzzle Width	103	Udder Depth	104	
Body Depth	100	Fore Attachment	104	
Chest Width	97	Rear Att Height	104	
Loin Strength	100	Rear Att Width	104	
Body Length	101	Centre Ligament	114	
Rump Length	99	Teat Placement (Front)	105	
Foot Angle	95	Teat Placement (Rear)	109	
Rear Set	98	Teat Length	99	
Rear Leg Rear View	99			

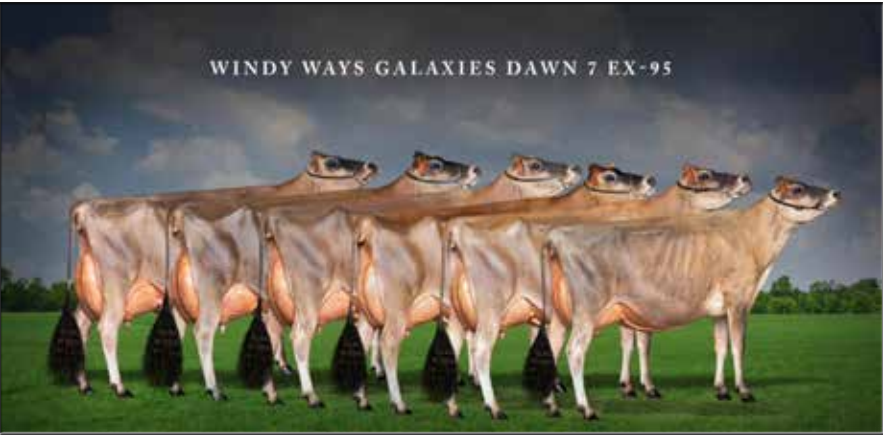
- Imagine the potential - a Roulette son from the famous Galaxies Dawn EX95
- Dingo is an impressive individual who is in high demand worldwide.



Dam: Windy Ways Galaxies Dawn EX95



Windy Ways CCC Dingo



Windy Ways CCC Dingo

SONGSIDE

Miami Songside-ET (Woodside / Vanahlem EX92 / Tbone VG85 2yrs)

A2 STATUS: A2/A2 Haplotypes: JH1F
NASIS: TLGSONGSIDE HB: AUS822422 DOB: 03/09/2022

SEXED \$55.00 RRP \$22.00
Ultraplus CONVENTIONAL PLUS20

PRODUCTION ABV 08/25		
Milk	336	78% R
Fat	19 kg	0.02%
Protein	21 kg	0.21%
ASI	155	70% R

WORKABILITIES	
Milking Speed	104
Temperament	107
Likability	107

BPI 293 70%R

HWI 209 59%R

SI 629 63%R

Conformation ABV(g)	
Overall Type	109
Mammary	112

HEALTH ABV (g) 08/25		
Cell Count	107	68%
Survival	108	59%
Feed Saved	-28	34%
Dtr Fertility	96	58%
Heat Tol	93	39%

CONFORMATION ABV (g)	0 Dtrs	0 Herds	65% Rel	August 2025
OVERALL TYPE	109	MAMMARY SYSTEM		112
Stature	102	Pin Width		102
Bone Quality	104	Pin Set		103
Angularity	106	Udder Texture		110
Muzzle Width	101	Udder Depth		102
Body Depth	102	Fore Attachment		108
Chest Width	100	Rear Att Height		105
Loin Strength	102	Rear Att Width		106
Body Length	100	Centre Ligament		107
Rump Length	102	Teat Placement (Front)		109
Foot Angle	98	Teat Placement (Rear)		108
Rear Set	102	Teat Length		102
Rear Leg Rear View	103			

- Adds Teat length to world class udders
- +109 Type, +112 Mammary backed by the strong and deep "Song" cow family



Miami Songside



Dam: Miami Vanahlem Song 4608 EX92



Miami Songside

DOUBS-pp

Peak Doubs-PP (AltaLuken / Joiner / Gallantry)

NEW

A2 STATUS: A2/A2 Genetic Codes: PP Haplotypes: JH1F JNSF
HB: JE840003272457390 NASIS: GXDOUBS DOB: 15/08/2024

RRP \$24.00



Peak Doubs-PP

Australian Indices August 2025									
BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
328	267	704	171	241	28	0.29%	20	0.25%	72
Dtr Fert	Mast	SCC	M/Speed	Temp	Likeability				
101	104	129	101	103	105				
Liveweight	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM			
101	0	-14	111	96	99	102			

USDA CDCB Genomic Index August 2025							
GJPI	MILK	FAT	%F	PROT	%P	Rel%	NM\$
127	665	40	0.03	30	0.02	73	379
DPR	PL	SCS	GEST L	HCR	CCR	PTAT	JUI
-0.7	0.1	3.00	0	2.2	-0.2	1.00	15.4

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025									
PTAT: 1.00 JUI: 15.4					GJPI 127				
Stature	Short				Tall	-0.10			
Strength	Frail				Strong	+0.30			
Dairy Form	Tight Rib				Open Rib	+0.70			
Rump Angle	High				Sloped	+0.20			
Rump Width	Narrow				Wide	-0.10			
RL SV	Posty				Sickle	-0.80			
Foot Angle	Hock-In				Straight	+0.50			
Fore Attachment	Loose				Strong	+0.30			
Rear Udder Height	Low				High	+0.90			
Rear Udder Width	Narrow				Wide	+0.70			
Udder Cleft	Weak				Strong	+0.80			
Udder Depth	Deep				Shallow	-0.10			
Fore Teat Placement	Wide				Close	+0.50			
Rear Teat Placement	Wide				Close	+0.30			
Teat Length	Short				Long	+0.30			

-1.5 -1.0 -0.5 0 0.5 1.0 1.5

Exciting New Sire, PP, A2A2, Fertility, Teat length

A game changer for Australian Jersey breeders

INCHSTA-P

Attaview Inchsta-P (Starlord / Oliver-P / Brax)

NEW

A2 STATUS: A2/A2 Haplotypes: JH1F
NASIS: TLGINCHSTA HB: AUS834812 DOB: 10/10/2024

SEXED \$55.00 RRP \$24.00
Ultraplus[®] CONVENTIONAL PLUS20



Half sister to Dam: Attaview Bontino Inch EX92

BPI 324 68%R

HWI 329 56%R

SI 587 62%R

Conformation ABV(g)

Overall Type 106

Mammary 109

CONFORMATION ABV (g) 0 Dtrs 0 Herds 62% Rel August 2025			
OVERALL TYPE	106	MAMMARY SYSTEM	109
Stature	103	Pin Width	103
Bone Quality	102	Pin Set	99
Angularity	102	Udder Texture	106
Muzzle Width	102	Udder Depth	104
Body Depth	99	Fore Attachment	105
Chest Width	101	Rear Att Height	107
Loin Strength	104	Rear Att Width	105
Body Length	102	Centre Ligament	104
Rump Length	101	Teat Placement (Front)	104
Foot Angle	107	Teat Placement (Rear)	104
Rear Set	99	Teat Length	101
Rear Leg Rear View	103		

PRODUCTION ABV 08/25		
Milk	267	78% R
Fat	24 kg	0.18%
Protein	12 kg	0.06%
ASI	108	68% R

WORKABILITIES	
Milking Speed	102
Temperament	105
Likability	107

HEALTH ABV (g) 08/25		
Cell Count	119	67%
Survival	110	55%
Feed Saved	-26	33%
Dtr Fertility	106	55%
Heat Tol	102	39%

First bull in AI from Attaview Jerseys from the consistent Inch family

Heterozygous polled sire with high daughter fertility 106

IRYMPLE

Loxleigh Roulette Irymple (Roulette / Valentino EX90 / Navarian EX91)

A2 STATUS: A2/A2 Haplotypes: JH1F
NASIS: TLGIRYMPLE HB: AUS808583 DOB: 21/09/2022

RRP \$22.00

CONVENTIONAL PLUS20



Same family as Loxleigh Badger Iris 4 - EX94

PRODUCTION ABV 08/25		
Milk	527	79% R
Fat	26 kg	-0.04%
Protein	21 kg	0.06%
ASI	152	70% R

WORKABILITIES	
Milking Speed	102
Temperament	106
Likability	107

BPI 283 70%R

HWI 208 58%R

SI 595 64%R

Conformation ABV(g)

Overall Type 108

Mammary 109

HEALTH ABV (g) 08/25

Cell Count	111	70%
Survival	109	57%
Feed Saved	-77	33%
Dtr Fertility	99	56%
Heat Tol	97	39%

CONFORMATION ABV (g)	0 Dtrs	0 Herds	63% Rel	August 2025
OVERALL TYPE	108	MAMMARY SYSTEM		109
Stature	107	Pin Width		107
Bone Quality	106	Pin Set		102
Angularity	104	Udder Texture		106
Muzzle Width	107	Udder Depth		100
Body Depth	102	Fore Attachment		100
Chest Width	102	Rear Att Height		103
Loin Strength	101	Rear Att Width		107
Body Length	103	Centre Ligament		111
Rump Length	101	Teat Placement (Front)		107
Foot Angle	103	Teat Placement (Rear)		111
Rear Set	99	Teat Length		98
Rear Leg Rear View	97			

- Roulette son from the famous "Iris" family
- +108 Type, 109 Mammary, Wide sloped rumps



Loxleigh Routette Irymple

LEMONPEEL - P

Wallacedale Lemonpeel -P (Polled Gold / Lemonhead EX90 / Marvarie)

A2 STATUS: A2/A2 Haplotypes: JH1F
NASIS: TLGLEMONPEEL HB: AUS792933 DOB: 12/03/2021

SEXED \$48.00 RRP \$18.00

Ultraplus

CONVENTIONAL PLUS20



GDAM Wallacedale Marvs Rosella VG88

PRODUCTION ABV 08/25		
Milk	179	75% R
Fat	13 kg	0.06%
Protein	7 kg	0.01%
ASI	55	68% R

WORKABILITIES	
Milking Speed	104
Temperament	107
Likability	107

BPI 215 68%R

HWI 187 58%R

SI 386 62%R

Conformation ABV(g)

Overall Type 106

Mammary 106

HEALTH ABV (g) 08/25

Cell Count	121	66%
Survival	109	60%
Feed Saved	-38	34%
Dtr Fertility	100	61%
Heat Tol	100	39%

CONFORMATION ABV (g)	0 Dtrs	0 Herds	64% Rel	August 2025
OVERALL TYPE	106	MAMMARY SYSTEM		106
Stature	104	Pin Width		106
Bone Quality	103	Pin Set		105
Angularity	102	Udder Texture		102
Muzzle Width	104	Udder Depth		103
Body Depth	98	Fore Attachment		103
Chest Width	102	Rear Att Height		107
Loin Strength	105	Rear Att Width		105
Body Length	101	Centre Ligament		98
Rump Length	102	Teat Placement (Front)		103
Foot Angle	104	Teat Placement (Rear)		97
Rear Set	99	Teat Length		101
Rear Leg Rear View	101			

- Early daughters calving in are well liked
- +106 Type and +106 Mammary with +107 Likability

BIGTOP

Bushlea Bigtop (Craze / Valentino EX92 / Nathan EX94)

A2 STATUS: A2/A2 Haplotypes: JH1F
NASIS: TLGBIGTOP NAAB: 187JE05595 HB: AUS798071 DOB: 11/05/2021

SEXED \$50.00

RRP \$18.00

Ultraplus

CONVENTIONAL PLUS20



Bushlea Bigtop

Australian Indices August 2025

BPI	HWI	SI	ASI	Milk	Fat	Fat%	Prot	Prot%	Rel%
262	198	518	97	312	23	0.12%	11	0.01%	79

Dtr Fert	Mast	SCC	M/Speed	Temp	Likeability
97	103	127	103	106	107

Liveweight	Gest L	Feed S	Survival	Heat Tol	TYPE	MAMM
103	-5	-39	110	98	106	111

USDA CDCB Genomic Index August 2025

GJPI	MILK	FAT	%F	PROT	%P	Rel%	NMS
76	-164	7	0.08	-5	0.01	77	72
DPR	PL	SCS	GEST L	HCR	CCR	PTAT	JUI
0.4	1.3	2.99	-2	-1.0	0.9	1.20	19.1

CONFORMATION 0 Dtrs 0 Herds USDA/HA Genomic Evaluation August 2025

PTAT: 1.20 JUI: 19.1 GJPI 76

Stature	Short						Tall	+1.10
Strength	Frail						Strong	+0.20
Dairy Form	Tight Rib						Open Rib	+1.10
Rump Angle	High						Sloped	+0.20
Rump Width	Narrow						Wide	+0.40
RL SV	Posty						Sickle	-0.20
Foot Angle	Hock-In						Straight	+0.60
Fore Attachment	Loose						Strong	+0.50
Rear Udder Height	Low						High	+1.90
Rear Udder Width	Narrow						Wide	+1.30
Udder Cleft	Weak						Strong	+1.20
Udder Depth	Deep						Shallow	+0.90
Fore Teat Placement	Wide						Close	+0.60
Rear Teat Placement	Wide						Close	+0.90
Teat Length	Short						Long	+1.30

- Incredible breed leading type +1.20 PTAT
- Remains an extreme udder sire at +19.1 JUI

EXQUISITE - P

Wallacedale Exquisite-PO (Goldband P / Lemonhead EX92 / Tbone EX90)

A2 STATUS: A2/A2 Genetic Codes: HP Haplotypes: JH1F
HB: AUS814935 NASIS: TLGEXQUISITE DOB: 21/02/2023

SEXED \$52.00

RRP \$20.00

Ultraplus

CONVENTIONAL PLUS20



Dam: Wallacedale Lemon Melaine - EX92

PRODUCTION ABV 08/25

Milk	326	78% R
Fat	21 kg	0.06%
Protein	17 kg	0.13%
ASI	133	70% R

WORKABILITIES

Milking Speed	104
Temperament	105
Likability	104

BPI 233 70%R

HWI 185 58%R

SI 522 63%R

Conformation ABV(g)

Overall Type	102
Mammary	101

HEALTH ABV (g) 08/25

Cell Count	110	69%
Survival	107	57%
Feed Saved	-11	34%
Dtr Fertility	100	56%
Heat Tol	97	39%

CONFORMATION ABV (g) 0 Dtrs 0 Herds 64% Rel August 2025

OVERALL TYPE	102	MAMMARY SYSTEM	101
Stature	100	Pin Width	109
Bone Quality	100	Pin Set	103
Angularity	103	Udder Texture	104
Muzzle Width	100	Udder Depth	100
Body Depth	100	Fore Attachment	99
Chest Width	102	Rear Att Height	100
Loin Strength	108	Rear Att Width	102
Body Length	98	Centre Ligament	99
Rump Length	99	Teat Placement (Front)	102
Foot Angle	104	Teat Placement (Rear)	101
Rear Set	100	Teat Length	106
Rear Leg Rear View	101		

- Teat Length and Daughter Fertility improver
- 286 BPI, A2A2 and Heterozygous polled jersey



The highest levels of industry accreditation ensure quality outcomes.

Delivering world class products and services since 1989.

At the forefront of animal genetics and reproduction technology.

Offering a quality range of dairy sires for your next joining period.

PHOTO BY BRAD CULLEN

TLG offer a wide range of professional services:

- ⊙ Semen Collection
- ⊙ Sexed Semen Collections
- ⊙ Storage & Despatch
- ⊙ ET and IVF Technology
- ⊙ Import & Export
- ⊙ Genomic & A2 Testing



TLG Product Specialist: Mat Dennis
Phone: 0467 027 178 Email: mat.dennis@genaust.com.au
4969 Princes Hwy, Camperdown Victoria 3260
Phone: 61 3 5593 2016 Email: sales@tlg.com.au www.tlg.com.au