

# UNDERSTANDING U.S. SIRE PROOFS

## GENETIC CODES

- BD** Bulldog\*
- BL** Bovine Leukocyte Adhesion Deficiency (BLAD)\*
- TL** Tested free of BLAD
- BY** Brachyspina\*
- TY** Tested free of Brachyspina
- CD** Cholesterol Deficiency\*
- TC** Tested free of Cholesterol Deficiency
- CV** Complex Vertebral Malformation (CVM)\*
- TV** Tested free of CVM
- DP** Deficiency of Uridine Monophosphate Synthase (DUMPS)\*
- TD** Tested free of DUMPS
- MF** Mulefoot\*
- TM** Tested free of Mulefoot
- PO** Observed Polled\*\*
- PC** Tested Heterozygous Polled\*\*
- PP** Tested Homozygous Polled\*\*
- TP** Tested free of the Polled Condition (horned)
- RC** Carrier for red hair color
- B/R** Black/Red hair colour\*
- TR** Tested free to red hair colour
- DR1** Tested Heterozygous for Dominant Red\*
- DR2** Tested Homozygous for Dominant Red\*

**Net Merit (NM)**  
Additional net profit the offspring will provide over her lifetime.  
Expressed in \$US.

**Predicted Transmitting Ability - Type (PTAT)**  
Average: 0  
Higher is better.

**Total Performance Index (TPI)**  
Combines genetic proofs for production, type, longevity and fertility into a single value.  
Higher is better.  
GTPi = Genomic TPI value

**Daughters and Herds**  
with production data  
Reliability accuracy (increases with more daughters)

**HAPLOTYPES AFFECTING FERTILITY**  
Researchers have identified 5 haplotypes (HH1-HH5) that affect cattle fertility mainly through early embryo losses. If an animal inherits a copy of this haplotype from each parent, that embryo will not survive to become a calf, so fertility is affected.

**HAPLOTYPE FOR CHOLESTEROL DEFICIENCY (HCD)**  
A haplotype has been identified that causes calves to have no cholesterol in the homozygous state. Calves who produce no cholesterol will die at several months of age. The economic impact of this haplotype is quite large, due to the costs of raising the calf for months before it dies.

**Origin of Production proof**  
CDCB = Domestic proof;  
HA = Domestic Proof  
MACE = Interbull MACE Proof  
The letter "G" indicates the inclusion of genomic information

format: proof month/year

**Productive Life (PL)**  
Measurement of longevity, including yield information.  
Higher is better.

**Somatic Cell Score (SCS)**  
An indicator trait for mastitis resistance based on the direct measure of somatic cells in milk samples. Lower is better.

**Daughter Pregnancy Rate (DPR)**  
The percentage of non-pregnant cows that become pregnant during each 21-day period. A bull with a DPR of 1 indicates that his daughters have 1% higher pregnancy rate than a bull with a DPR of 0.  
Higher is better.

**Based on daughter classification.**  
Average: 0  
Higher is better.

\*For these traits, 0 is ideal. These traits include **Rump Angle, Rear Legs Side View, Fore Teat Placement, Rear Teat Placement and Fore Teat Length.**

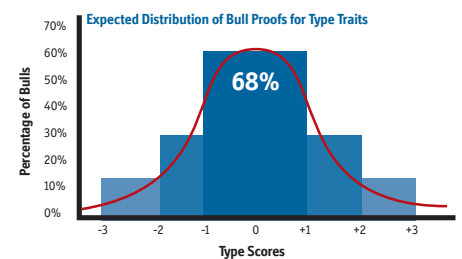
<b>GTPi 2698</b>		EX-90-CAN TR TL TY TV		HH1F HH2F HH3F HH4F HH5F HH6F HCDF	
Reg. #:	HO84013130920421	aAa:	342156	DMS:	345,135
Born:	08/02/2015	Kappa Casein:	BE	Beta Casein:	A1A2
<b>PRODUCTION</b>	53 Herds	169 Dtrs	94% Rel	MACE-G / 12-19	
Milk lbs	278	Fat lbs	106	Fat %	+0.35
Protein lbs	28	Protein %	+0.07		
NM\$	878	CM\$	915	FM\$	801
GMS	843	DWPS	1021		
Daughter Average (ME) Milk <b>28,167 lbs</b> Fat <b>1,166 lbs</b> Protein <b>910 lbs</b>					
<b>HEALTH &amp; REPRODUCTION</b>					
Productive Life	5.9	Sire Calving Ease	7.5%	99% Rel	
Somatic Cell Score	2.75	Daughter Calving Ease	5.2%	73% Rel	
Daughter Pregnancy Rate	2.4	Sire Stillbirth	7.2%		
Livability	3.0	Daughter Stillbirth	5.3%		
Fertility Index	2.5	Feed Efficiency	186		
<b>CONFORMATION</b>	32 Herds	70 Dtrs	91% Rel	MACE / 12-19	
PTAT	2.24	Body Composite	-0.06		
Udder Composite	2.89	Dairy Composite	1.21		
Feet & Legs Composite	0.79				
Stature			Tall	+1.16	
Strength			Strong	+0.27	
Body Depth			Deep	+0.52	
Dairy Form			Open Rib	+1.42	
Rump Angle *			High	-1.53	
Rump Width			Wide	+0.60	
Rear Legs Side View *			Posty	-0.17	
Rear Legs Rear View			Straight	+0.23	
Foot Angle			Steep	+1.88	
F & L Score			High	+1.03	
Fore Attachment			Strong	+3.02	
Rear Udder Height			High	+4.12	
Rear Udder Width			Wide	+3.79	
Udder Cleft			Strong	+1.40	
Udder Depth			Shallow	+2.27	
Fore Teat Placement *			Close	+0.63	
Rear Teat Placement *			Close	+0.56	
Fore Teat Length *			Long	+0.34	

**ANIMAL MODEL** – The animal model represents proven methodology for calculating Predicted Transmitting Abilities (PTAs). Evaluations are based on the animal and its relationship to other animals being evaluated. Information from the animal itself, its ancestors, and its progeny are incorporated, with all known relationships among the animals considered.

**PRODUCTION TRAITS** – Production traits are expressed in predicted transmitting ability, which is an estimate of the genetic superiority or inferiority that an animal will transmit to the offspring.

**TYPE TRAITS** – Type Traits are expressed as Standardized Transmitting Abilities (STAs). This allows you to easily compare different traits of the same bull and see which traits have the most extreme values. The graph below illustrates normal distribution of bull proofs for conformation traits.

**STANDARD DEVIATIONS** – Simply put, the standard deviation tells us how far a bull is from the mean (average) value of all the bulls in the breed. The graph below shows the normal distribution of bull proofs for type traits with 68% of bulls ranging from -1 to +1, 95% from -2 to +2, 99% from -3 to +3, etc.). For most bulls their scores will be close to the average of all the bulls, while fewer bulls will score at the high end or at the low end. Very few bulls reach scores as high as 3.2 as compared with over 99% of bulls that fall in the range of +3 to -3.



For more information on the American Proving System visit:  
Council of Dairy Cattle Breeding (CDCB) at <https://www.uscdcb.com>