

BROWNSCBSWISS

PEDIGREE

Name and Number: BLESSING GAMECHANGER GLORY

000000068203916

Date of Birth: 09/09/2019 Sex: Female

Tattoo: 920

Genetic Conditions:

Not Classified

Parent Average PPR: +2 PTAT: +0.6

PA: -117m -10f +3p -29NMS

Breeder:
BLESSING FARMS
Current Owner:
BLESSING FARMS
FT WAYNE IN



SCHERMA BLOOMING BIVER *TM

000120094290070

MARC SCHERTENLEIB 09/22/2012

Genetic Conditions: PT DT MT WT BH2T

Not Classified SUPERIOR

PPR: +154 92%R PTAT: +1.0 96%R (12/19)

PTA: +664m +41f +43p +352NMS 97%R (MACE)

PTA PL: +1.4 SCS: +2.96 DPR: -0.7 SCE: +6.1

PTA LIV: +1.1

5247 dau. av. 25349 4.2 1070 3.5 877

3527 class. dau. av. FS:84.0 UDC:0.60 FLC:0.66

VIKING VALLEY BROOKINGS GIFT

000000068147720

BRATLAND, GERALD & SONS 09/28/2012

Genetic Conditions: PT DT MT WT BH2T

07/00 E91 E95 E92 E91 V85 E92 (10/19)

ST SR DF RA TW RR RS FA UA UH UW UC UD TP TL

9 9 7 4 7 6 3 7 9 8 8 9 5 7 6

PPR: +100 70%R PTAT: +0.7 76%R

PTA: +402m +33f +26p +255NMS 90%\$ (GEN)

DEV: +961m +98f +62p (12/19)

Protein Plus

02/00 305d 3X 24580 4.8 1186 3.8 927 DHIR

02/00 365d 3X 28620 4.8 1384 3.8 1079 DHIR

03/02 305d 3X 29460 4.8 1400 3.7 1095 DHIR

03/02 365d 3X 34510 4.7 1635 3.7 1292 DHIR

04/08 305d 3X 31940 5.2 1655 3.6 1137 DHIR

04/08 365d 3X 36830 5.1 1871 3.6 1314 DHIR

06/03 305d 3X 31270 4.7 1474 3.3 1018 DHIR

06/03 365d 3X 37730 5.3 1984 3.3 1255 DHIR

Lifetime: 1777d 160850m 8082f 5849p

BLESSING TEX BRAIDEN ET *TM

000000000198435

BLESSING FARMS & MARK MORT 12/04/2005

Genetic Conditions: *TM PT DT WT BH2T

E92

PPR: -83 96%R PTAT: +0.5 98%R (12/19)

PTA: -499m -38f -23p -191NMS 99%R (GEN)

PTA PL: +2.2 SCS: +2.86 DPR: -0.1 SCE: +7.2

PTA LIV: +2.1

875 dau. av. 22280 3.9 878 3.3 730

577 class. dau. av. FS:84.7 UDC:0.18 FLC:-0.36

TOP ACRES PILOT GRITZ ET

000000000944303

SLIKER, WAYNE E. & KENNY 09/05/2005

Genetic Conditions:

07/04 2E-E91 E90 E92 E93 E90 E90 (01/13)

ST SR DF RA TW RR RS FA UA UH UW UC UD TP TL

7 5 7 4 8 8 5 5 4 7 6 6 4 4 4

PPR: -140 76%R PTAT: +0.1 81%R

PTA: -973m -50f -35p -408NMS 0%\$

DEV: -1873m -84f -59p (12/19)

02/07 305d 3X 23920 3.8 919 3.2 769 DHIR

02/07 365d 3X 27820 3.9 1098 3.3 914 DHIR

03/08 305d 3X 18490 4.1 750 3.6 674 DHIR

03/08 314d 3X 18770 4.1 762 3.6 685 DHIR

04/08 305d 3X 23660 3.8 904 3.4 812 DHIR

04/08 365d 3X 27350 3.8 1051 3.4 942 DHIR

06/09 305d 3X 20980 4.3 894 3.4 711 DHIR

06/09 365d 3X 24580 4.3 1045 3.5 852 DHIR

Lifetime: 1750d 112830m 4553f 3939p

HILLTOP ACRES B GAMECHANGER

000000068174506

MASHEK, DENNIS 12/17/2015

Genetic Conditions: PT DT MT WT BH2T

Not Classified

PPR: +151 60%R PTAT: +1.0 66%R (12/19)

PTA: +613m +41f +42p +331NMS 65%R (GEN)

PTA PL: +1.2 SCS: +2.95 DPR: -0.8 SCE:

PTA LIV: +0.6

0 dau. av.

FOREST LAWN BRAIDEN GIGI ET

000000068155747

FOREST LAWN FARM LLC 12/04/2013

Genetic Conditions:

AGE FS SS DQ RP MO MS

05/08 E90 E93 E95 E91 V87 V87 (08/19)

ST SR DF RA TW RR RS FA UA UH UW UC UD TP TL

9 9 9 5 7 7 4 4 6 7 8 8 4 4 3

PPR: -147 54%R PTAT: +0.1 59%R

PTA: -847m -60f -37p -388NMS 0%\$

DEV: -1987m -134f -84p (12/19)

02/01 305d 3X 19680 3.5 685 3.2 620 DHIR

02/01 365d 3X 22790 3.6 812 3.2 729 DHIR

03/03 305d 2X 17540 3.5 615 3.3 584 DHIR

03/03 365d 2X 19540 3.5 690 3.3 654 DHIR

04/05 305d 2X 18660 3.4 635 3.2 595 DHIR

04/05 365d 2X 21450 3.5 759 3.3 698 DHIR

PEDIGREE INFORMATION

Official Brown Swiss Three-Generation Performance Pedigrees are computer generated from information permanently stored on BSCBA files. Information contained in a pedigree includes:

ANIMAL IDENTIFICATION

1. Registration name and number of the animal, its parents, and grandparents (underlined areas).
2. For the animal itself, date of birth, sex, and tattoo, Breeder and Current Owner.

MALES - PERFORMANCE INFORMATION

1. Name & Registration Number followed by Breeder and Birthdate.
2. Abnormality test indicators and Haplotype test indicators.
3. Current classification rating for Final Score and Sire recognition of 'Superior' or 'Qualified' if eligible.
4. Progressive Performance Ranking (PPR) value and its associated reliability (%R).
5. Predicted Transmitting Ability for Type (PTAT) for Final Score and its reliability (%R).
6. Date of all listed genetic evaluations.
7. Predicted Transmitting Ability (PTA) for production traits of milk, fat, and protein yields, Lifetime Net Merit dollars (NMS) and its reliability (%R). MACE indicates Interbull evaluation, GEN indicates Genomic evaluation, otherwise USDA evaluation only.
8. Predicted Transmitting Ability (PTA) for Productive Life (PL), Somatic Cell Score (SCS), Daughter Pregnancy Rate (DPR), Sire Calving Ease (SCE) and Livability(LIV).
9. Number of daughters included in the production evaluation and their 305d, 2x, ME average for milk, %fat, fat, % protein, and protein.
10. Number of classified daughters including their average Final Score (FS), Udder Composite (UDC) rating, and Foot & Leg Composite (FLC).
11. Show Awards

FEMALE - PERFORMANCE INFORMATION

1. Name & Registration Number followed by Breeder and Birthdate.
2. Abnormality test indicators and Haplotype test indicators.
3. Age at most recent classification and the Final Score including any multiple E designation, plus the scores of the five breakdown traits of Substance and Strength(SS), Dairy Quality(DQ), Rump(RP), Mobility(MO) and Mammary System(MS) for cows scored after 6-1-2007 or for cows scored prior to 6-1-2007 the five breakdown traits of Frame(FM), Dairy Character(DC), Body Capacity(BC), Feet & Legs(FL), and Mammary System(MS), followed by the date of classification (month, year). The numeric score associated with each breakdown is that of the most recent classification.

E	=	Excellent (90-95 points)	G	=	Good (75-79 points)
V	=	Very Good (85-89 points)	F	=	Fair (65-74 points)
+	=	Good Plus (80-84 points)	P	=	Poor (below 65 points)

4. Cow recognition of 'Elite' or 'Certified' if eligible.
5. Linear descriptive type appraisal scores on 15 traits based on a scoring system of 1 to 50 prior to 5/15/03 and a 1 to 9 scale thereafter.

ST - Stature	RR - Rear Legs (rear view)	UW - Udder Width
SR - Strength	RS - Rear Legs (side view)	UC - Udder Cleft
DF - Dairy Form	FA - Foot Angle	UD - Udder Depth
RA - Rump Angle	UA - Fore Udder Attachment	TP - Teat Placement
TW - Thurl Width	UH - Udder Height	TL - Teat Length

6. Progressive Performance Ranking (PPR) and Predicted Transmitting Ability for Type (PTAT).
7. Recognition of 'Superior Brood Cow' if applicable.
8. Predicted Transmitting Ability (PTA) for production traits of milk, fat, and protein yields, Lifetime Net Merit dollars (NMS), and percentile rank based on Lifetime Net Merit (%NM) labeled (%\$). GEN indicates Genomic evaluation, otherwise USDA evaluation.
9. Management group deviations for milk, fat, and protein yields and date of genetic evaluation run.
10. Recognition of Protein Plus if eligible.
11. Official production records.

a. Age at calving in years and months.	f. Fat in pounds.
b. Number of days milked (d).	g. Percent protein.*
c. Times milked per day (2x or 3x).	g. Protein in pounds.*
d. Milk in pounds.	h. Type of record (DHIR or DHIA).
e. Percent fat.	* (True Protein as of 5/1/2000)

12. Lifetime totals for number of days, pounds of milk, pounds of fat, and pounds of protein if over 100,000 pounds milk, 4,000 pounds fat, or 3,500 pounds protein.
13. Show and Production Awards

GENETIC CONDITIONS

SUFFIXES to Name:

- (W); (A); (M); (D) - Known Carrier for Weaver, Spiderleg, SMA, or SDM determined by identification of affected offspring or gene test.
(CW); (CA); (CM); (CD) - Confirmed affected animal for Weaver, Spiderleg, SMA, or SDM.
(W*); (A*); (M*); (D*) - Tested Carrier at a confidence level of 90% or above for Weaver, Spiderleg, SMA, or SDM.
*TW; *TA; *TM; *TD - Tested Non-carrier at a confidence level of 90% or above for Weaver, Spiderleg, SMA, or SDM.
NP, P, or PP - Known Naturally Polled (prior), Polled (current), or Homozygous Polled.

TEST INDICATORS:

- WT, MT, DT, PT, BH1T, BH2T - Indicates Genomic Tested Non-carrier of Weaver, SMA, SDM or Polled genes, or BH1, BH2 haplotypes.
WC, MC, DC, PC, BH1C, BH2C - Indicates Genomic Tested Carrier of Weaver, SMA, SDM or Polled genes, or BH1, BH2 haplotypes.

NOTE: On some database reports referencing CDCB data, the following haplotype indicators are used:
BHWC or BHWT (Weaver carrier or tested free); BHMC or BHMT (SMA carrier or tested free);
BHDC or BHDT (SDM carrier or tested free); BHPC or BHPT (Polled carrier or tested free)

GENERAL

1. Young animals without performance information show an estimate of genetic merit given as a parent average (PA).
Parent Average = 1/2 (Sire + Dam) for all genetic traits.
2. Date pedigree was printed is listed in lower left-hand corner.
3. Order number is listed in the lower right-hand corner.

Order Number: B00076663