Jan 2, 2024

AJCA Genomic Evaluation Report

100K GENOTYPE

FEMALE: BETTER WITH THYME STARLIGHT {1}-P JEUSA000175042625 JH1F JNS-TF Born: 10/20/2021 Tattoo: BWT01

Sire: UR UNKNOWN JERSEY SIRE {0} JEUSA00099PJS0104

Dam: BETTER WITH THYME MMF MOONBEAM {1}-P JEUSA000175020085 JH1F JNS-

Owner: BETTER WITH THYME FARM

PO BOX 3105

SHEPHERDSTOWN, WV 25443

Single-breed (S) Evaluation					
P-level Inbreeding Percents					
P0	Genomic: 6.3%				
BBR	Pedigree: 8.6%				
100	Genomic Future: 1.2%				

Jersey Haplotype 1 (JH1) is associated with embryo loss. Official status is listed as F for Free, C for Carrier. Breed Base Representation (BBR): CDCB policy is to report BBR values of 94 or greater for one breed as 100. BBRs below 94 are reported as calculated. BBR is reported once, unless animal is re-genotyped with a higher density chip.

Adjustments are applied by CDCB to Traditional Evaluations for yield traits (third column from right) to correct for bias in female genetic evaluations. The Adjusted Evaluations are provided for index and yield traits. The impact of genomic information can be assessed by comparing Genomic Evaluation with the Adjusted Evaluation (column labeled Genomic Impact).

INDEX							
	Genomic	Adjusted		Traditional Eval	Genomic	Traditional	
Trait	Evaluation	Evaluation	Genomic Impact	1/2024	REL %	REL %	
JPI	-100	-31	-69		62		
Net Merit (\$)	-322	-182	-140				
Cheese Merit (\$)	-327	0	-327				
Fluid Merit (\$)	-277	0	-277				
Grazing Merit (\$)	-244						

YIELD							
Trait	Genomic Evaluation	Adjusted Evaluation	Genomic Impact	Traditional Eval 1/2024	Genomic REL %	Traditional REL %	
Milk (lb)	-1205	-478	-727		66		
Fat (lb)	-57	-25	-32		66		
Fat (%)	0.01	-0.01	0.02		66		
Protein (lb)	-46	-22	-24		67		
Protein (%)	-0.01	-0.02	0.01		67		

Note: Genomic and Traditional Evaluations for Health, Fitness and Type traits are expressed on similar scales. No adjustments have been applied. A comparison of the Genomic Evaluation with the January 2024 Traditional Eval indicates the Genomic Impact.

HEALTH and FITNESS						
To be	Genomic	Traditional Eval		Genomic	Traditional	
Trait	Evaluation	1/2024	Genomic Impact	REL %	REL %	
Pregnancy rate (%)	3.0	1.5	1.5	53		
Cow conception rate (%)	2.0	1.4	0.6	53		
Heifer conception rate (%)	-1.5	-0.5	-1.0	35		
Productive life (mo)	-0.1	0.0	-0.1	59		
Livability	2.3	1.8	0.5	39		
Somatic cell score	2.98	2.96	-0.02	64		

ТҮРЕ						
Total In	Genomic	Traditional Eval		Genomic	Traditional	
Trait	Evaluation	1/2024	Genomic Impact	REL %	REL %	
Final score (PTAT)	-1.70	0.39	-2.09	70		
Stature	-2.70	0.31	-3.01			
Strength	-1.90	0.12	-2.02			
Dairy form	-2.40	0.27	-2.67			
Rump angle	-1.00	-0.22	-0.78			
Rump width	-2.00	0.21	-2.21			
Rear legs (side view)	-0.10	-0.05	-0.05			
Foot angle	-0.50	0.21	-0.71			
Fore udder attachment	-2.10	0.63	-2.73			
Rear udder height	-1.70	0.33	-2.03			
Rear udder width	-2.00	0.02	-2.02			
Udder cleft	-0.60	0.07	-0.67			
Udder depth	-0.90	0.61	-1.51	72		
Front teat placement	-1.70	0.14	-1.84	·		
Teat length	-0.40	0.11	-0.51	·		
Jersey Udder Index	-4.13	12.92	-17.05			