

Effects of weighting schemes and base adjustment on US genomic validation

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Key findings from (Mota et al. 2023, 2024)

- Large breeds and highly heritable traits are most stable
- Small breeds and low-heritability traits are often unstable
- Failures are driven by a few candidate bulls rather than model issues
- TMACE provided more consistent validation



Why revisit genomic validation now?



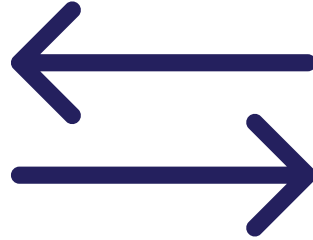
GEBV are widely used to support early selection and marketing of young bulls.

- International semen marketing requires standardized and transparent genomic validation
- Increasing model complexity and trait expansion
- Limitations of traditional 4-year reference validation
- Persistent challenges for small breeds and low-heritability traits

Why compare validation scenarios?



1 GEBVtest outcomes influence international acceptance



2 Small methodological differences may alter regression statistics



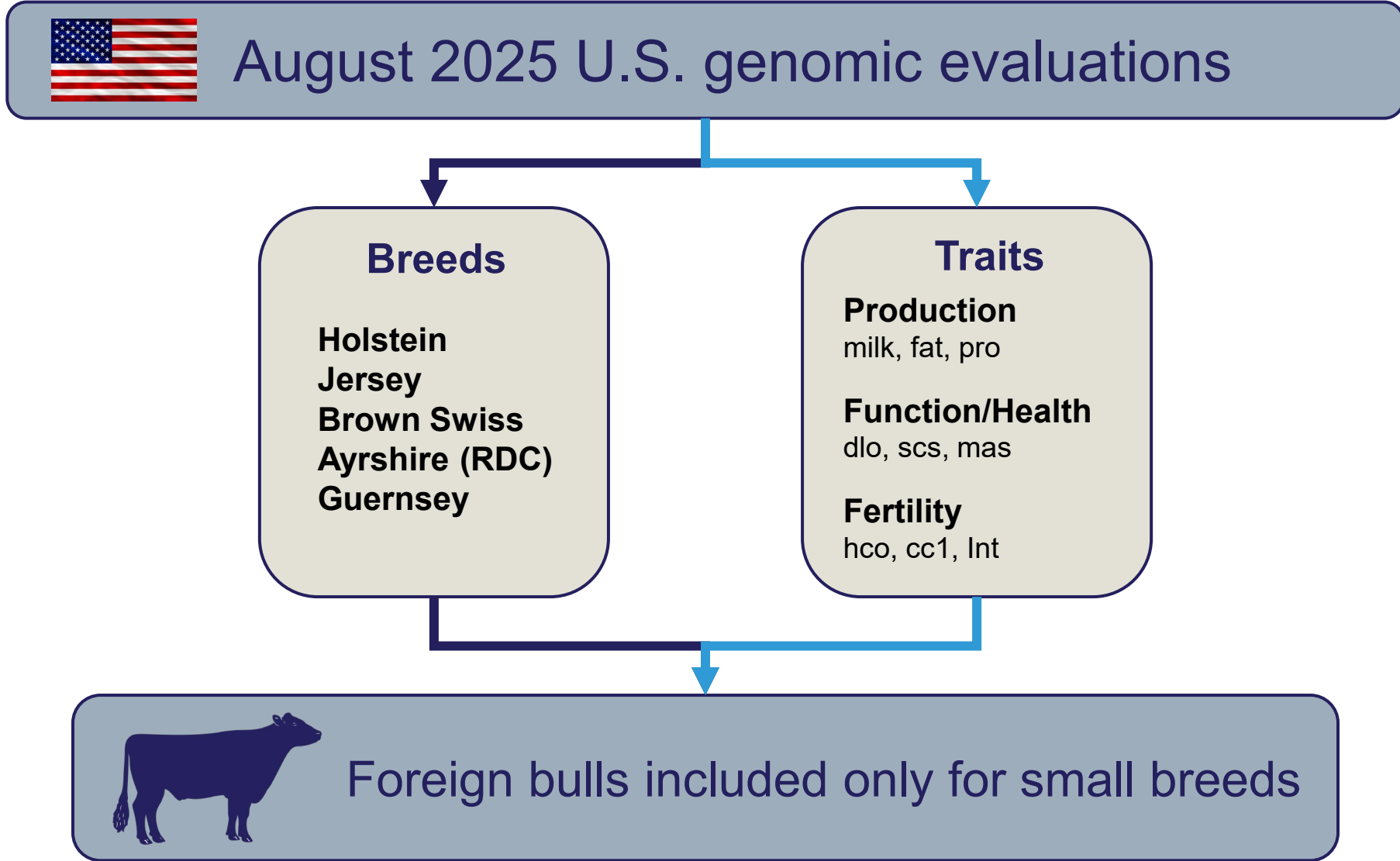
3 Different scenarios may lead to different PASS/FAIL outcomes

Understanding scenario sensitivity is essential for robust genomic validation.

Validation Scenarios

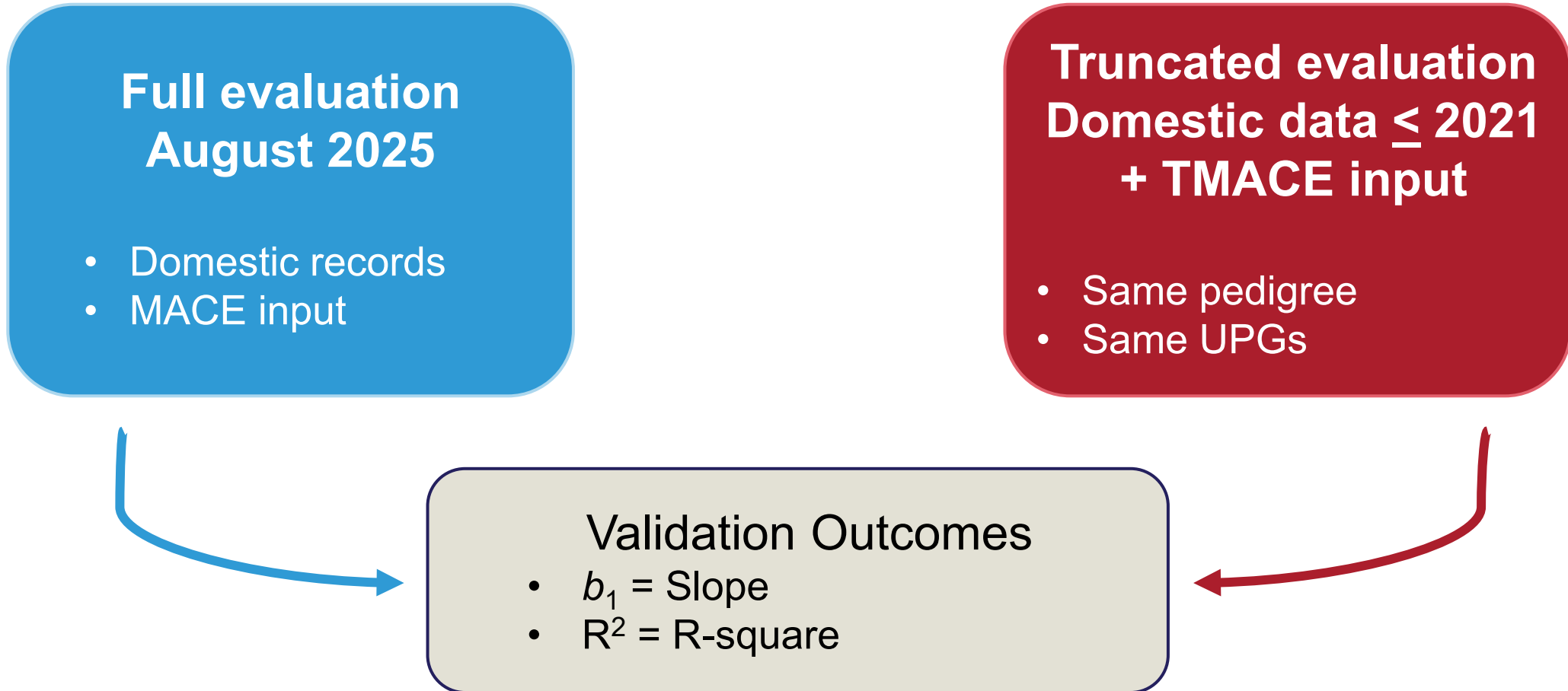
Scenario	Target	Regression	Base Adjustment	Weight
1	dGEBV	ITB	No	Yes
2	dGEBV	ITB	Yes	Yes
3	dGEBV	LR	No	No
4	dGEBV	LR	Yes	No

Data and Breeds



Validation Framework

TMACE-based genomic validation



Overall results

- All five breeds remained stable across scenarios
- Smaller breeds were mainly constrained by the number of validation bulls
- Some exceptions were observed depending on breed and trait



Validation group definition across scenarios

Tested bulls are defined by the regression method:

ITB (weighted) and LR (unweighted) use different sets of test bulls.

Scenario	Regression Method	Base Adjustment	Slope (b_1)	Validation Result
1	ITB	No	1.174	Pass
2	ITB	Yes	1.225	Fail
3	LR	No	1.1465	Pass
4	LR	Yes	1.1968	Pass

LR methods include additional candidate bulls, so the candidate group size is higher.

Key Observations

The number of tested bulls **are not the sole driver** of validation failures.

Validation outcomes are driven by **interactions among key methodological choices**, not a single factor.

What the Differences Tell Us

Validation behavior depends on the **interaction** among key methodological choices:

- Regression method (ITB vs. LR)
- Base adjustment (Yes vs. No)
- Validation group definition

Take Home Messages



Validation outcomes depend on configuration



Base adjustment and validation-group definition interact



Interpretation of GEBVtest requires caution



CDCB is transitioning to single-step evaluation with direct GEBV output, which may **improve scale consistency** and **validation performance**

Acknowledgements and disclaimers

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- Participating dairy producers for data supply
- DHI organizations and DRPCs for processing and relaying data to CDCB
- Purebred breed associations for providing pedigree data

The mention of trade names or commercial products is solely for the purpose of providing specific information and does not imply recommendation or endorsement by CDCB.

Thank you!

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