


Preliminary results on de-regressed proof in single-step GBLUP

Yutaka Masuda
University of Georgia

Zengting Liu, VIT, Germany
Peter Sullivan, Lactanet, Canada

“Genomic-free” de-regression



Option 3: ssDRP

- Deregressed ssGEBV Proofs
 - Methods developed, but **no proof of concept** for MACE
 - Do ssDRP include daughter phenotype but NOT genotype?
 - Does de-regressing genotypes re-introduce **GPS bias**?
 - To be applied by each country (**need genotype access**)
 - **New programming** by each country, or by Interbull for all

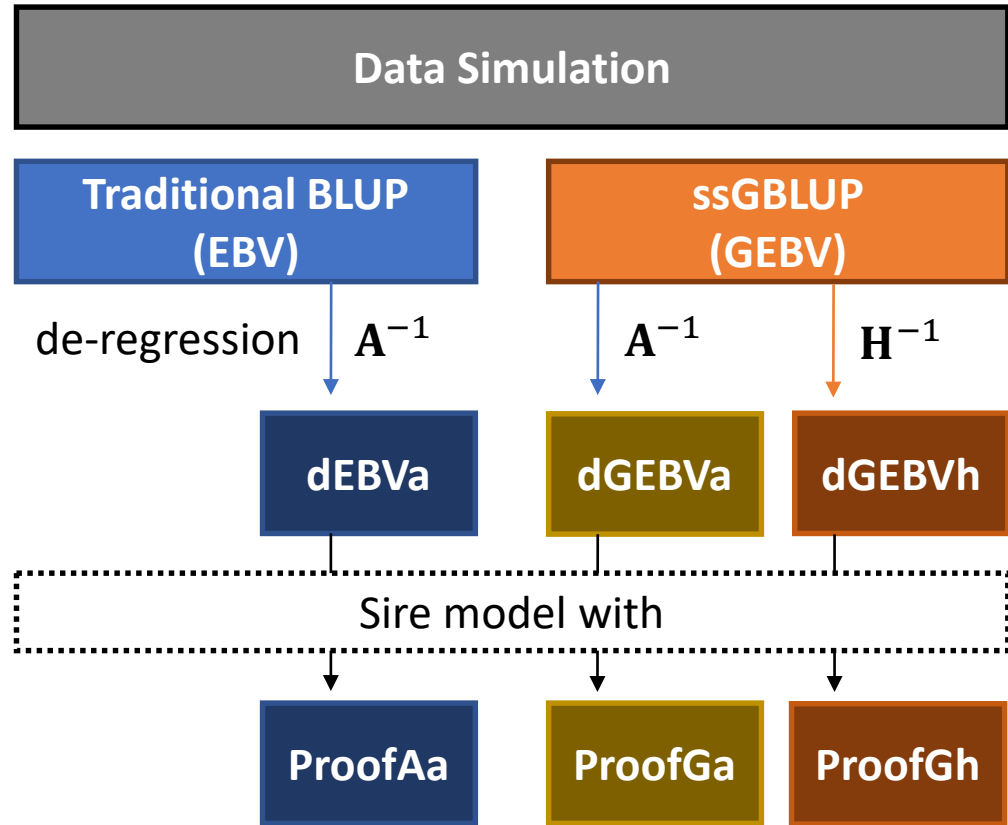
Genomic-free EBV for MACE *(Interbull Webinar, Feb 11, 2021)*

- What if using as

in each country

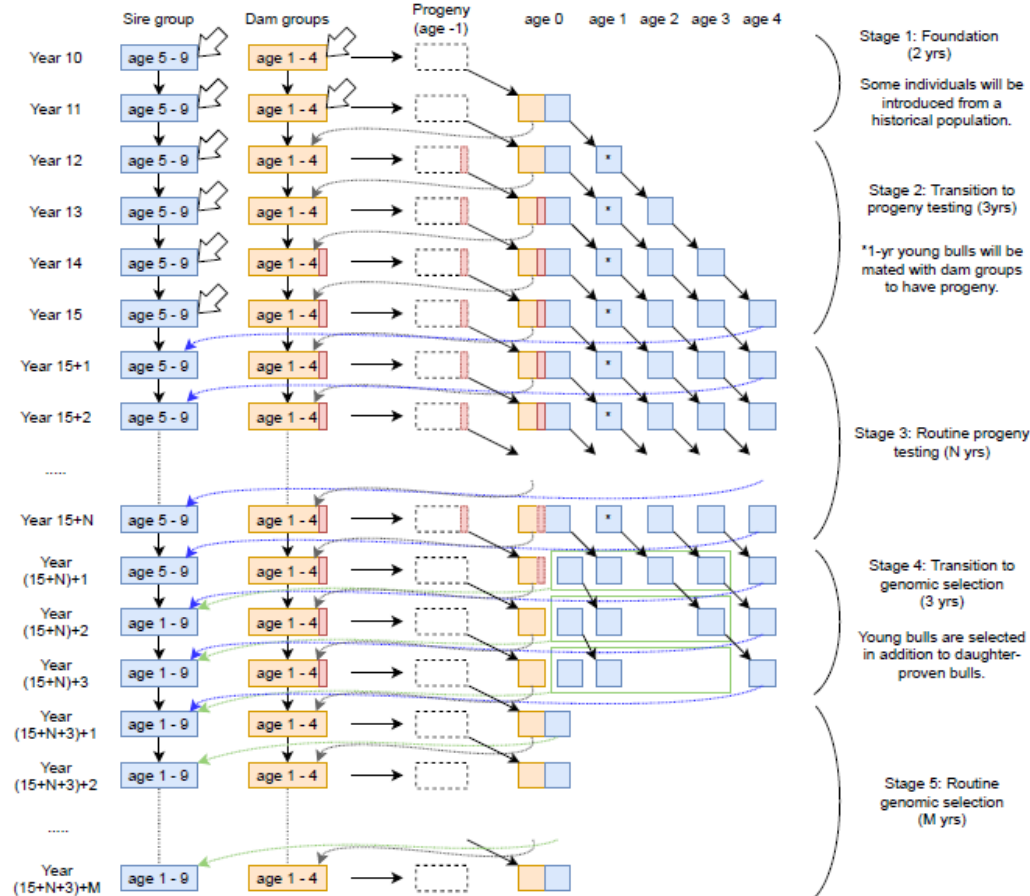
- Zengting has suggested but never published it in public.
 - What kind of animals should we use for de-regression?
 - What is the computing cost?
 - How does

Objectives



- Examine if
 - De-regressed proofs with reflect GEBV well.
 - De-regressed proofs account for the pre-selection bias in a sire model with
- Target: proven bulls
 - Genetic trend of EBV/GEBV
 - Genetic trend of re-evaluated proofs with sire model

Simulation



- Mimicking a dairy-cattle breeding program
 - Yrs 10-11: foundation
 - Yrs 12-15: transition to **progeny testing** (20 selected bulls out of 200 candidates; 40 active bulls)
 - Yrs 16-30: **progeny testing**
 - Yrs 31-33: transition to **genomic selection** (25 selected young-bulls out of 500 candidates; 50 active)
 - Yrs 34-42:

Data and model

Data	N	Description
Phenotypes	284,783	<ul style="list-style-type: none">• One phenotype per cow• PT bull with at least 50 daughters• Selected bulls with 100 daughters on average per year (Gamma dist.)
Pedigree	1,541,288	<ul style="list-style-type: none">• No missing parents• Dam pedigree traced back within a herd
Genotypes	6,900	<ul style="list-style-type: none">• Bulls born after generation 20• 1989 proven bulls• 4911 young bulls

- Genetic evaluation

- and
- Progeny testing: animal model BLUP:
- Genomic selection: TBV + noise with reliability = REL_G (equivalent to DE=15) + REL_PA
- Simplified EDC based on Interbull Method (Fikse and Banos, 2001)

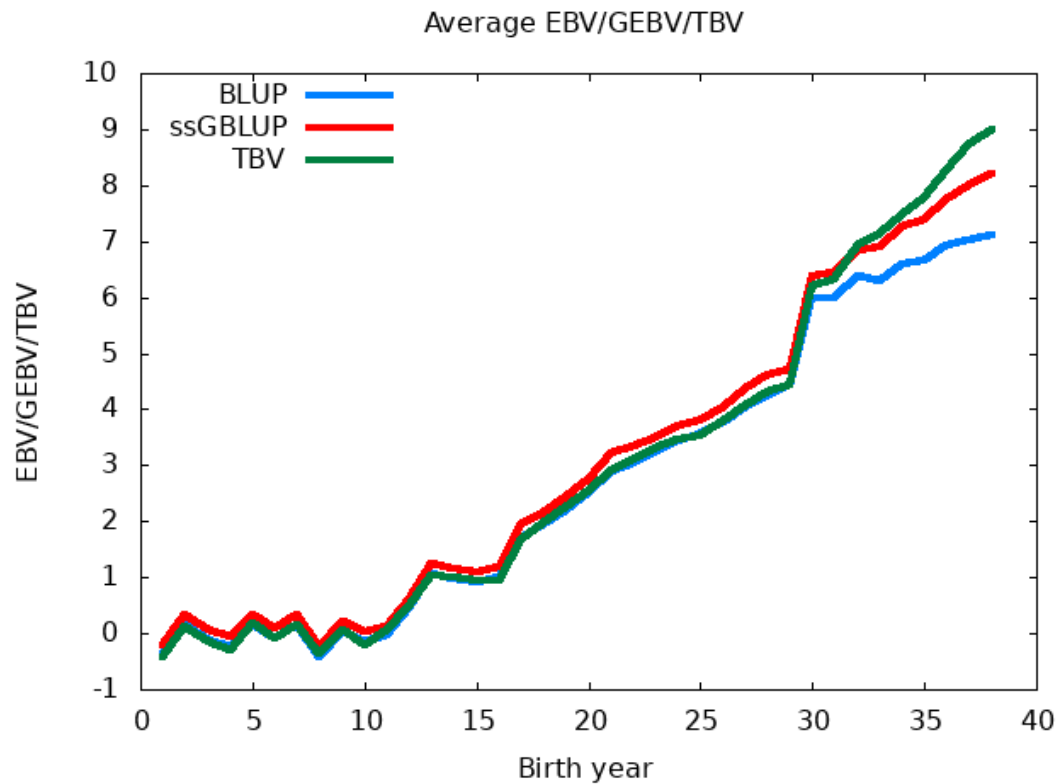
De-regression methods

- Method 1: based on
 - Based on Jairath et al. (1998) but no UPG
 - : sire EBV, : sires' ancestor EBV,
: diagonal matrix of EDC,
: de-regressed proof
- Method 2: based on
 - Same as Method 1 except for replacing with
 - With all animals i.e., genotyped (both proven and young bulls) and non-genotyped animals
 - Thanks to Zengting's

Re-evaluation of de-regressed proof

- Mimicking “MACE” with de-regressed proofs
 - (= fixed birth-year group effect)
It is close to the MACE model.
- Single-trait sire model with MGS pedigree
- Confirm if
 - reproduces the original GEBV.
 - Pre-selection bias in disappears.
- Thanks to Peter’s suggestion

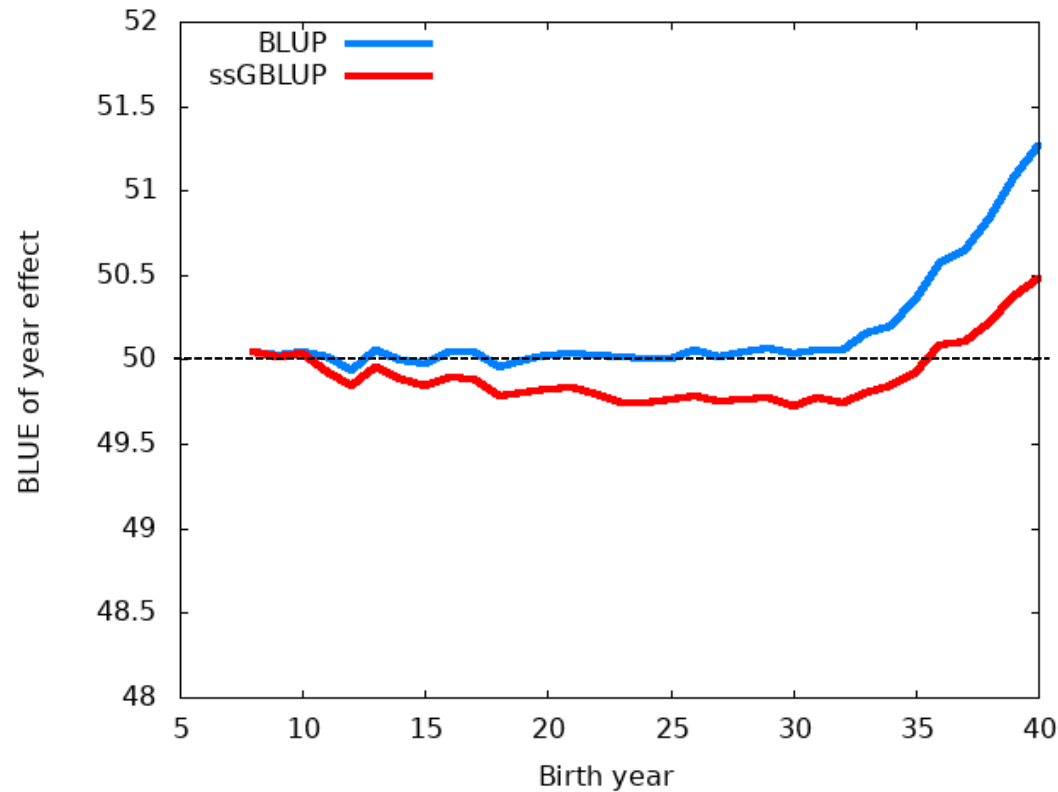
Genetic trend of TBV/EBV/GEBV



For bulls with daughters

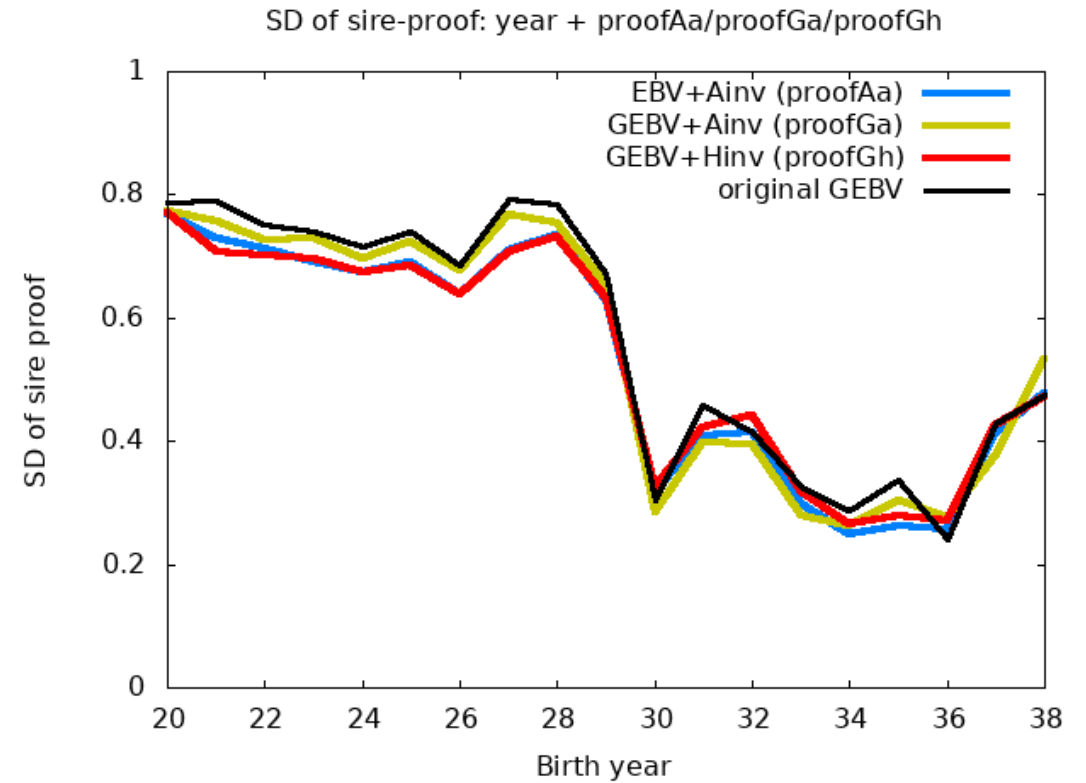
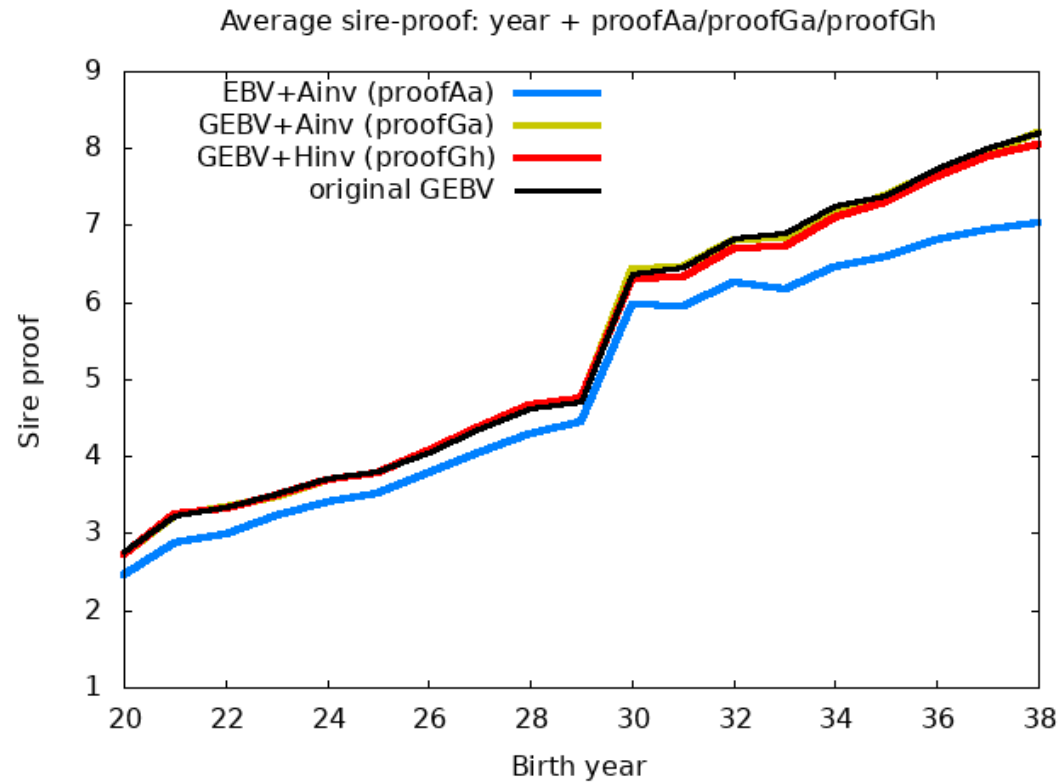
- Clear change of genetic trends after year 30
 - **EBV**: highly biased
 - **GEBV**: less but still biased
- Missing information in G
 - Selection based on pseudo GEBV (TBV + noise)
 - Only proven bulls (and young genotypes) in G
 - Small size of data

Where the bias goes?

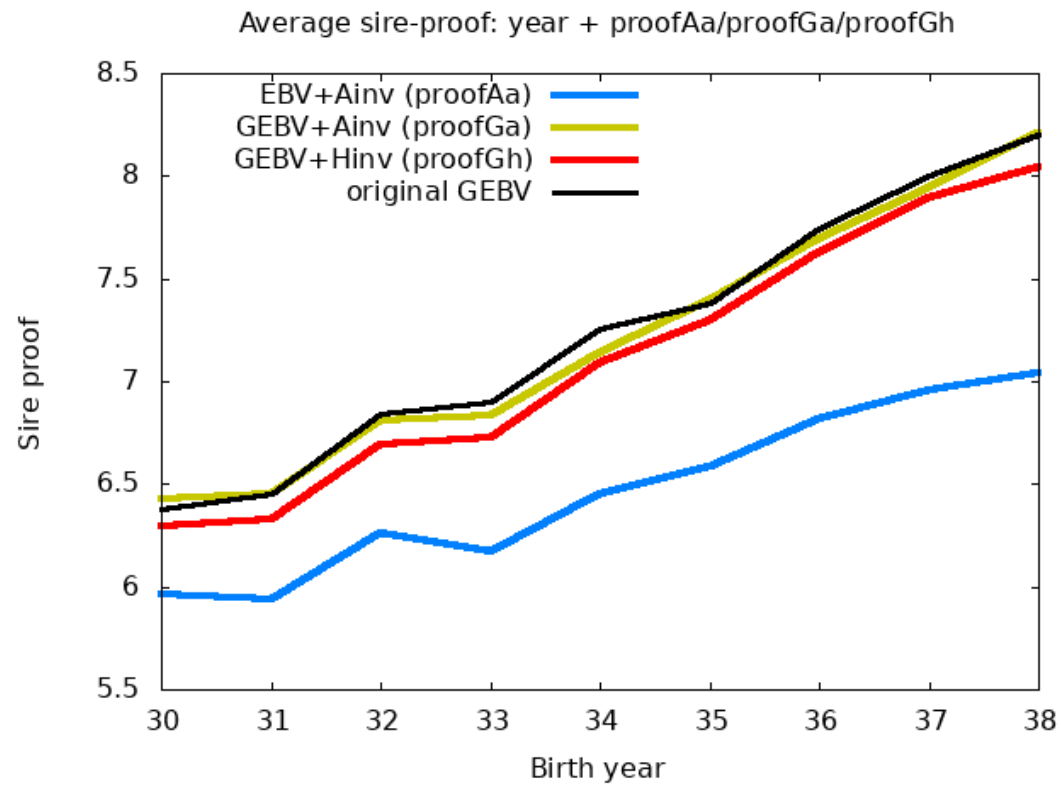


- Pre-selection bias merged to the year effect
 - Pointed out by Esa.
 - In practice, it will be confounding with herd-year or the other contemporary effects.
 - And possibly, it could be merged to UPG effects.

Sire-model proof with year effect ()



Years ≥ 30 ()



proof	Ga	Gh	TBV	EBV	GEBV
Aa	0.99	>0.99	0.97	0.90	0.97
Ga		>0.99	0.97	0.84	0.95
Gh			0.98	0.86	0.95
TBV				0.83	0.94
EBV					0.96

Summary

- De-regressed proofs from ssGBLUP can work.
 - Deregressed with or .
 - Able to reproduce GEBV by MACE-style sire model.
 - No pre-selection bias in reproduced sire proofs.
- ... with simulated data
 - Correct validation method?
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- Very preliminary results: many concerns
 - Real data?
 - Genotyped daughters
 - Multiple-country data
 - Missing pedigree
 - Foreign (external) information
 - General de-regressed method: single-step GBLUP/SNPBLUP