

Deflated preconditioned conjugate gradient method for solving single-step single nucleotide polymorphism BLUP

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Single-step models

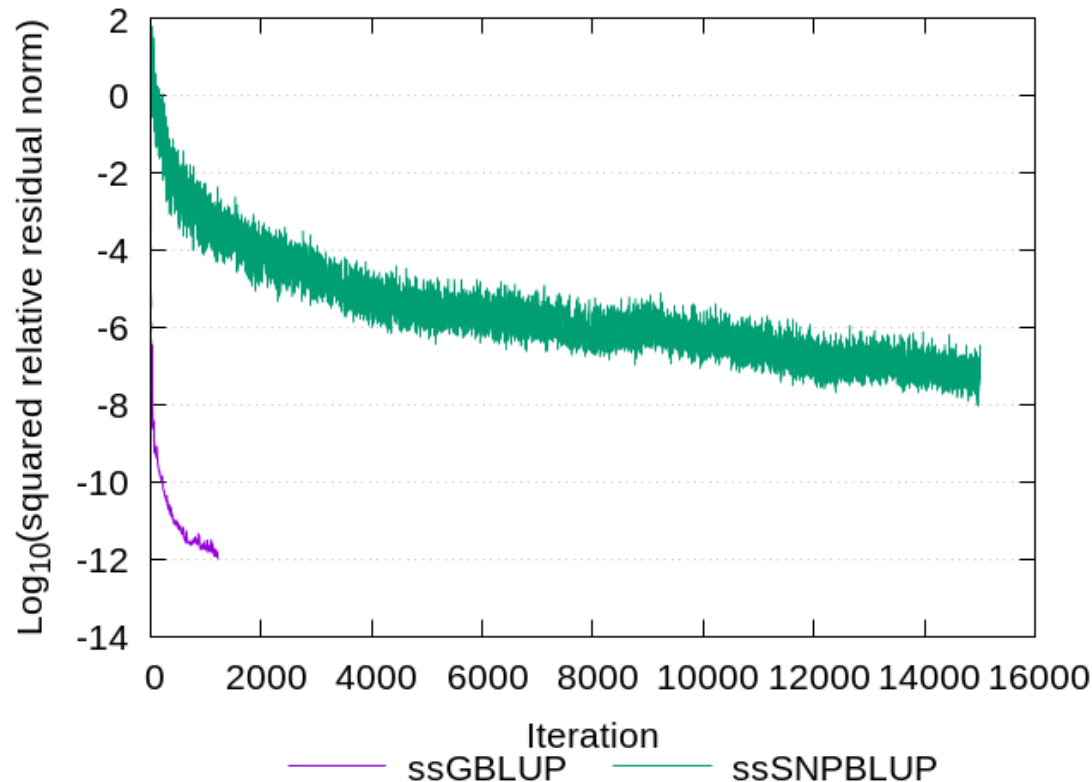
- Prediction of genomic breeding values
 - Genotyped and non-genotyped animals
- Single-step **GBLUP**
 - Animal-based model
 - Limited?
- Single-step **SNPBLUP**
 - SNP-based model
 - Several equivalent formulations
 - No limitation?!

Single-step SNPBLUP

System of equations has the form $Cx = b$

→ Iterative solver: Preconditioned Conjugate Gradient

→ Convergence issues often reported...



Aim

1. Comparison of **properties of coefficient matrices** of ssGBLUP and ssSNPBLUP

→ **Comprehension of convergence patterns of PCG**

2. Implementation of a **Deflated PCG method** for solving **efficiently** ssSNPBLUP

Equivalent single-step models

ssGBLUP:
$$\mathbf{y} = \mathbf{Xb} + \begin{bmatrix} \mathbf{W}_n & \mathbf{0} \\ \mathbf{0} & \mathbf{W}_g \end{bmatrix} \begin{bmatrix} \mathbf{u}_n \\ \mathbf{u}_g \end{bmatrix} + \mathbf{e}$$

ssSNPBLUP:
$$\mathbf{y} = \mathbf{Xb} + \begin{bmatrix} \mathbf{W}_n & \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \mathbf{W}_g & \mathbf{W}_g \mathbf{Z} \end{bmatrix} \begin{bmatrix} \mathbf{u}_n \\ \mathbf{a}_g \\ \mathbf{g} \end{bmatrix} + \mathbf{e}$$

$$\mathbf{u}_g = \mathbf{a}_g + \mathbf{Zg}$$

↑
Addition of
SNP effects

b : fixed effects

u_n, u_g : aggregate GEBVs for (non-)genotyped animals

a_g : residual polygenic effects for genotyped animals

g : SNP effects

e : residuals

Z : SNP genotypes

Conjugate Gradient (CG)

- Successive approximations to obtain a more accurate solution of \mathbf{x} by solving

$$\mathbf{C}\mathbf{x}=\mathbf{b}$$

- Convergence

- Function of the effective condition number of \mathbf{C}

$$\kappa(\mathbf{C}) = \frac{\text{largest eigenvalue of } \mathbf{C}}{\text{(non-zero) smallest eigenvalue of } \mathbf{C}}$$

- Smaller condition number \rightarrow faster convergence

Preconditioned CG (PCG)

- Improvement of the condition number from $\kappa(\mathbf{C})$ to $\kappa(\mathbf{M}^{-1}\mathbf{C})$ by introducing a preconditioner \mathbf{M}

$$\mathbf{M}^{-1}\mathbf{C}\mathbf{x} = \mathbf{M}^{-1}\mathbf{b}$$

- In animal breeding
 - PCG often implemented
 - Usually: $\mathbf{M} = \text{diag}(\mathbf{C})$ (or a variant)

Deflated PCG (DPCG)

- Improvement of the condition number from $\kappa(\mathbf{M}^{-1}\mathbf{C})$ to $\kappa(\mathbf{M}^{-1}\mathbf{P}\mathbf{C})$ by introducing a second-level preconditioner \mathbf{P}

$$\mathbf{M}^{-1}\mathbf{P}\mathbf{C}\mathbf{x} = \mathbf{M}^{-1}\mathbf{P}\mathbf{b}$$

\mathbf{P} chosen such that unfavourable eigenvalues are set to 0 (deflated)

Deflated PCG (DPCG)

- **P** = deflation matrix

$$= \mathbf{I} - \mathbf{CZ}_d(\mathbf{Z}_d'\mathbf{CZ}_d)^{-1}\mathbf{Z}_d'$$

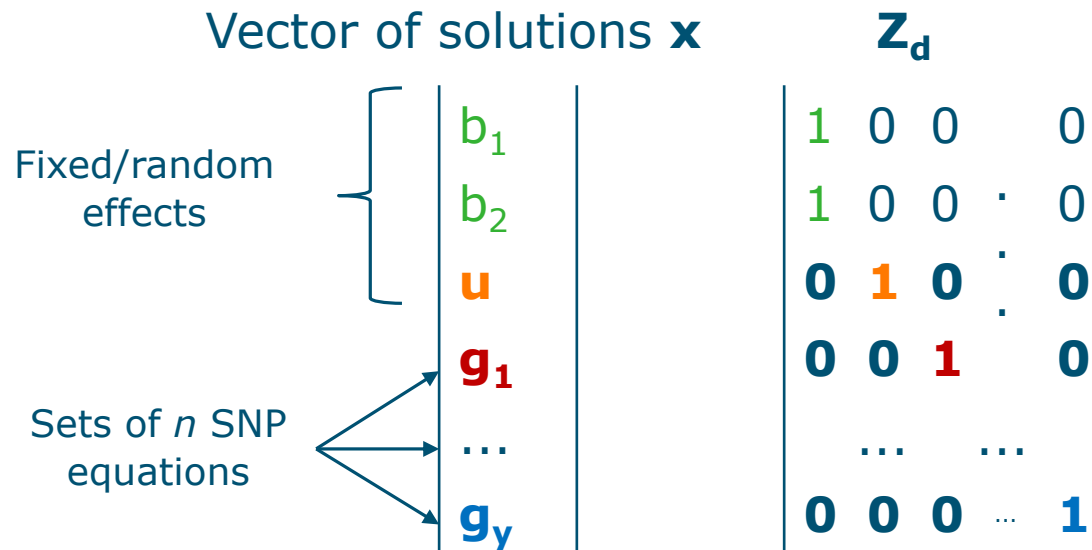
- **Z_d** contains the deflation vectors

- Approximation of the same space of the span of unfavourable eigenvectors

→ Set of deflation vectors \approx Set of unfavourable eigenvectors

Deflated PCG (DPCG)

- Setting-up of \mathbf{Z}_d following a **subdomain deflation approach**
 - **1 subdomain** per **fixed/random** effect
 - **1 subdomain** per **set of n** (1, 5, 50, or 200) **SNP equations**



Sparse $\mathbf{Z}_d \rightarrow$ efficient implementation \rightarrow small extra-costs

Data

- 61,592 Ovum pick-up sessions
- 37,021 animals
 - 4,109 phenotyped animals
 - 6,169 genotyped animals (without phenotype)
- 9,994 segregating SNPs

- Heritability = 0.35
- Residual polygenic variance = 5%

Results – Spectra and condition numbers

Model	Method	Smallest eigenvalue	Largest eigenvalue	κ	#iter.
ssG	PCG	$1.1 \cdot 10^{-4}$	11.9	$1.1 \cdot 10^5$	273
ssSNP	PCG	$1.1 \cdot 10^{-4}$	181.0	$1.7 \cdot 10^6$	1497

ssSNPBLUP vs ssGBLUP

- Unchanged smallest eigenvalues
- Increased largest eigenvalues
- ➔ Larger condition number
- ➔ Increased number of iterations

Results – Spectra and condition numbers

Model	Method	Smallest eigenvalue	Largest eigenvalue	κ	#iter.
ssG	PCG	$1.1 \cdot 10^{-4}$	11.9	$1.1 \cdot 10^5$	273
ssSNP	PCG	$1.1 \cdot 10^{-4}$	181.0	$1.7 \cdot 10^6$	1497
	DPCG (200)	$1.1 \cdot 10^{-4}$	99.4	$9.3 \cdot 10^5$	1195

200 SNPs equations per subdomain

- Unchanged smallest value
- Decreased largest eigenvalue
- Better condition number after deflation

Results – Spectra and condition numbers

Model	Method	Smallest eigenvalue	Largest eigenvalue	κ	#iter.
ssG	PCG	$1.1 \cdot 10^{-4}$	11.9	$1.1 \cdot 10^5$	273
ssSNP	PCG	$1.1 \cdot 10^{-4}$	181.0	$1.7 \cdot 10^6$	1497
	DPCG (200)	$1.1 \cdot 10^{-4}$	99.4	$9.3 \cdot 10^5$	1195
	DPCG (50)	$1.1 \cdot 10^{-4}$	40.5	$3.8 \cdot 10^5$	880
	DPCG (5)	$1.1 \cdot 10^{-4}$	6.0	$5.6 \cdot 10^4$	338
	DPCG (1)	$1.1 \cdot 10^{-4}$	6.0	$5.4 \cdot 10^4$	240

1 and 5 SNPs per subdomain

- Similar (decreased) condition numbers
- #iterations similar as ssGBLUP
- **Reduction** of #iter. by up to a factor **6!**

Conclusions

- ssSNPBLUP - PCG: larger eigenvalues
 - Larger condition number
- ssSNPBLUP - Deflated PCG
 - Treats the largest unfavourable eigenvalues
 - Smaller condition number
 - Faster convergence (similar to ssGBLUP)
- Similar pattern on large and multivariate ssSNPBLUP

Thank you!

