# A Hierarchical Spatial Finlay-Wilkinson Model for Analysis of Multi-Environment Field Trials

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# The Genomes to Fields (G2F) Initiative



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### Multi-Environment Field Trial Analysis for G2F Data

- Initially, we focus on a subset of 24 **environments**.
- We have yield recorded for 10,971 field plots with known spatial locations.
- A total of 1,105 hybrids are planted in approximately 10 plots on average.
- Hybrid **genotypes at**  $\sim$ **1M genomic locations** are available.
- To characterize environments, weather stations provide time-indexed measurements for 10 weather variables, and several soil variables are available.

### Finlay-Wilkinson (FW) Model

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$$y_{ijk} = \mu + g_i + h_j + b_i h_j + e_{ijk},$$

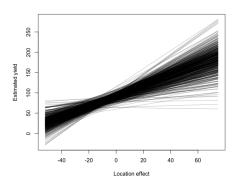
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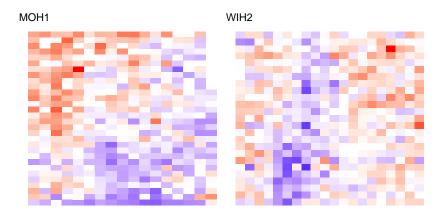
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### Residuals of FW Model for Two Fields

**Problem:** the residuals are **highly spatially correlated**.



### Hierarchical Spatial Finlay-Wilkinson (SFW) Model

• Data model:

$$[y_{ijk}|\mu, \mathbf{g}, \mathbf{b}, \mathbf{h}, \phi] \stackrel{indep}{\sim} \mathcal{N}(\mu + g_i + h_j + b_i h_j + \phi_{ijk}, \sigma_e^2),$$

Prior distributions for genotype, slope, and field effects:

$$\begin{split} [\mathbf{g}] &\sim \mathcal{N}(\mathbf{0}, \mathbf{A}\sigma_g^2); \quad [\mathbf{b}] \sim \mathcal{N}(\mathbf{0}, \mathbf{A}\sigma_b^2); \\ [\mathbf{h}|\gamma] &\sim \mathrm{N}(\gamma_1 \mathbf{Z}_1 + \dots + \gamma_l \mathbf{Z}_l + \dots + \gamma_L \mathbf{Z}_L, \mathbf{I}\sigma_h^2). \end{split}$$

- **A** is the kinship matrix describing the correlation structure of **g** and **b** (R package/software: rrBLUP, Tassel 5).
- $\mathbf{Z}_{I}$  is the *I*th standardized environmental covariate.



- A popular model for fertility adjustment in agricultural field trials is the first order intrinsic autoregression (Besag and Higdon, 1999; Dutta and Mondal, 2015).
- First order Intrinsic Autoregressive prior:

$$[\psi_j| heta_j,\sigma_j^2]\propto |\sigma_j^{-2}\mathbf{W}_j|_+^{1/2}\exp\left(-rac{1}{2}\sigma_j^{-2}\psi_j\mathbf{W}_j\psi_j
ight)$$

where

$$\psi_j \mathbf{W}_j \psi_j = \theta_j \sum \sum (\psi_{u,v} - \psi_{u-1,v})^2 + \bar{\theta}_j \sum \sum (\psi_{u,v} - \psi_{u,v-1})^2$$

ullet The distribution of  $\psi_j$  is **invariant** to the addition of arbitrary constant.

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#### Recall:

• 
$$[y_{ijk}|\mu, \mathbf{g}, \mathbf{b}, \mathbf{h}, \phi] \stackrel{indep}{\sim} \mathcal{N}(\mu + g_i + h_j + b_i h_j + \phi_{ijk}, \sigma_e^2),$$

#### Recall:

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#### Problem:

- The intrinsic spatial prior has an indeterminate overall level.
- The overall levels of spatial effects are confounded with the location effects.
- Estimation of **b** is biased.
- Hierarchical structure of h is not applicable.

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#### Solution:

• A hard constraint: set the average of the spatial effects to zero.



# Projected Intrinsic Autoregression (PIAR) Prior

• The Gaussian projected intrinsic autoregression (PIAR) on the  $r_j \times c_j$  regular array is then defined as:

$$\phi_j = \mathsf{B}_j arphi_j, \qquad arphi_j \sim \mathcal{N}(\mathbf{0}, \mathsf{D}_j^{-1}),$$

- $\mathbf{B}_j$  is an  $r_j c_j \times (r_j c_j 1)$  matrix.
- $\mathbf{D}_j$  is an  $(r_jc_j-1)\times(r_jc_j-1)$  diagonal matrix.
- ullet We can show:  $arphi_j = \mathbf{B}_j^{\mathrm{T}} \phi_j$ .

### Matrix Free Computation

- The covariance matrix of the Gaussian PIAR is a dense singular matrix.
- The computation load for generating  $\phi_j$  from PIAR using knowledge of multivariate statistics is  $\mathcal{O}((r_i c_i)^3/3)$ .
- Assume small number of missing plots (denote  $r_j c_j N_j$  as the number of missing plots).
- Thus matrix-vector multiplications with  $\mathbf{B}_j$  and  $\mathbf{B}_j^{\mathrm{T}}$  can also be performed using these discrete cosine transformations (DCT).
- The computation load of our proposed algorithm is  $\mathcal{O}(r_jc_j + (r_jc_j N_j)r_jc_j \log r_jc_j + (r_jc_j N_j)^3/3)$ .



### Prediction

- Implement posterior predictive distributions.
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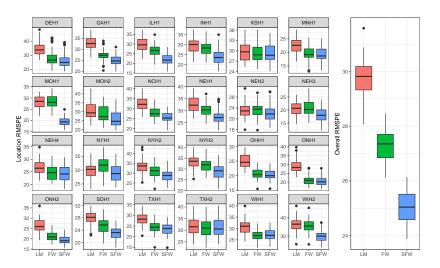
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#### Predict in new environments:

 By learning how environment effects depend on the weather and soil variables.

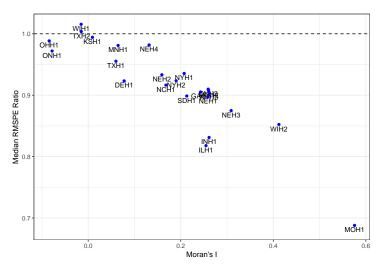
### Model Evaluation via Within-Field Prediction

Reduced error for yield prediction.



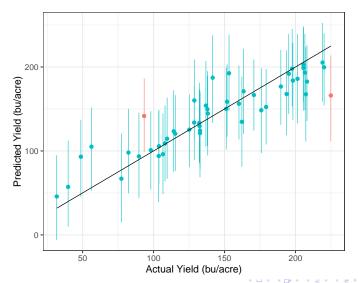
### Model Evaluation via Within-Field Prediction

Level of spatial correlation vs performance of SFW model.



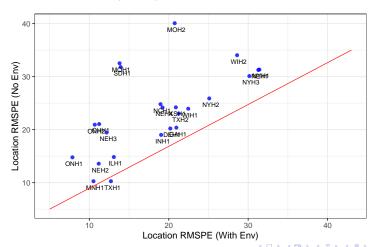
### Prediction Intervals

50 plot yield prediction intervals (95% credible level).



### Predict in New Environments

Location-wise RMSPEs computed using temperature and rainfall data (x-axis), versus the location-wise RMSPEs computed not using any environment information (y-axis).



### Selected References

- Besag, J. and Higdon, D. (1999). Bayesian analysis of agricultural field experiments. <u>Journal of the Royal Statistical Society: Series B</u> (Statistical Methodology), 61(4):691–746.
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Any opinions, findings, conclusions, or recommendations expressed in this publication are those of the authors and do not necessarily reflect the views of the U.S. Department of Agriculture.

# Thank You!

# Construction of $\mathbf{B}_j$ and $\mathbf{D}_j$

• Then the spectral decomposition of  $W_j$  is given by:

$$(\mathbf{N}_{r_j} \otimes \mathbf{N}_{c_j}) \mathbf{W}_j (\mathbf{N}_{r_j}^{\mathrm{T}} \otimes \mathbf{N}_{c_j}^{\mathrm{T}}) = \theta_j \mathbf{\Lambda}_{r_j} \otimes \mathbf{I}_{c_j} + \bar{\theta}_j \mathbf{I}_{r_j} \otimes \mathbf{\Lambda}_{c_j}.$$

- $\Lambda_k$  denote the  $k \times k$  diagonal matrix whose uth diagonal entry is  $4\sin^2\{\pi(u-1)/(2k)\}$ .
- $\mathbf{N}_k$  denotes the  $k \times k$  orthogonal matrix whose (u, v)th entry is  $1/\sqrt{k}$  if u = 1,  $\forall v$ , and  $(2/k)^{1/2} \cos\{\pi(u-1)(v-1/2)/k\}$  otherwise.
- $\mathbf{B}_{j}^{\mathrm{T}}$  denotes the  $(r_{j}c_{j}-1)\times r_{j}c_{j}$  matrix consisting of last  $r_{j}c_{j}-1$  rows of  $\mathbf{N}_{r_{j}}\otimes\mathbf{N}_{c_{j}}$ .
- $\mathbf{D}_j$  denotes the diagonal matrix consisting of the nonzero elements of  $\theta_j \mathbf{\Lambda}_{r_i} \otimes \mathbf{I}_{c_i} + \bar{\theta}_j \mathbf{I}_{r_i} \otimes \mathbf{\Lambda}_{c_i}$ .



# Assessing Uncertainty about FW Regression Lines

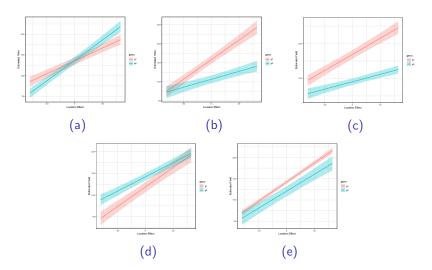


Figure: Estimated Yield vs Location Effect for pairs of genotypes