



# Two approaches to account for G x E interactions for production traits in South African Holstein



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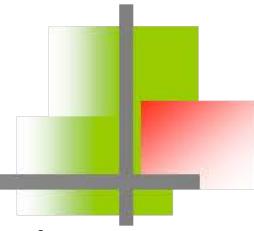
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# GxE in South Africa

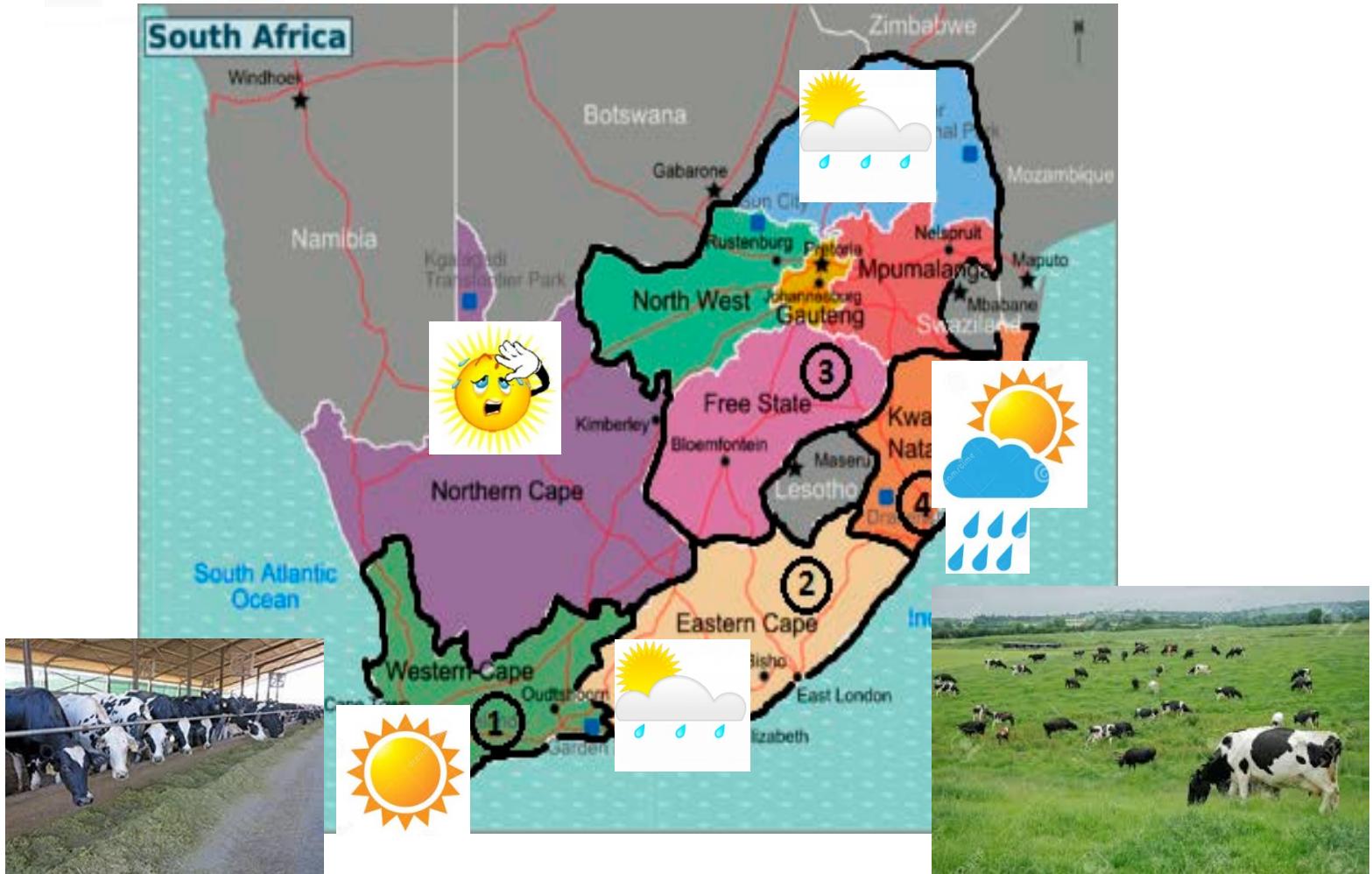


- Part of the GENOSOUTH project (INRA « metaprogramme »)
- Data on :
  - 4 traits: Milk, fat, protein, age at first calving
  - 378,782 first lactations between 1982 and 2012
  - 10 climatic variables at herd level

(averaged over 50 years, *closest weather station from the herd*):

*Average rainfall, temperature (min, max), relative humidity (min, max), solar radiation, evapotranspiration, summer temperature (max), summer solar radiation (max), summer relative humidity (max)*

# Environmental conditions in South Africa



# Contrasted performances

Region	Number of animals	Milk yield (kg)	Fat yield (kg)	Protein yield (kg)	Age at First Calving (mo)
(1) Western Cape	91 024	<b>7 729</b>	282	249	27
(2) Eastern Cape	56 251	<b>5 955</b>	214	191	29
(3) Free State, Gauteng, North West and Limpopo	135 218	<b>6 758</b>	246	215	29
(4) KwaZulu Natal	90 415	<b>5 767</b>	207	183	29

# Models (1)

## 1. Univariate model:

$$y_{305d} = \sum \text{fixed effects} + a + e$$

*Herd-year + calving season + age at first calving*

The diagram illustrates a univariate model. It shows a red oval labeled  $A \sigma_a^2$  connected by a red arrow to the term 'a' in the equation. A blue oval labeled  $I \sigma_e^2$  is connected by a blue arrow to the term 'e'. A black brace underlines the terms 'a' and 'e' in the equation, grouping them together.

## 2. with heterogeneous residual variance

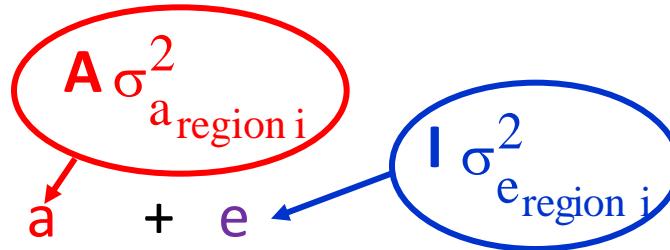
$$y_{305d} = \sum \text{fixed effects} + a + e$$

The diagram illustrates a model with heterogeneous residual variance. It shows a red oval labeled  $A \sigma_a^2$  connected by a red arrow to the term 'a' in the equation. A blue oval labeled  $I \sigma_e^2_{\text{region } i}$  is connected by a blue arrow to the term 'e'. A black brace underlines the terms 'a' and 'e' in the equation, grouping them together.

# Models (2)

## 3. Multivariate models (« MACE type ») : 1 region = 1 trait

$$y_{305d} = \sum \text{fixed effects} +$$



(in fact,  $\text{var}(a) = G_0 \otimes A$ )

- a.  $G_0 = \text{rank } 4$  (10 parameters)
- b.  $G_0 = \text{rank } 3$  (9 parameters)
- c.  $G_0 = \text{rank } 2$  (7 parameters)
- d.  $G_0 = \text{rank } 1$  (4 parameters)

More parsimonious  
models using the  
Wombat software  
(K. Meyer)

# Models (3)

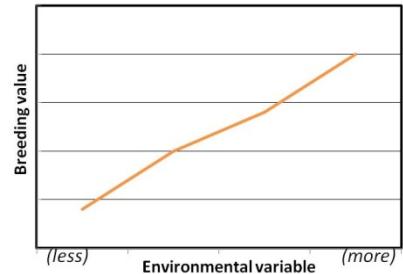
## 4. reaction norm models:

$$y_{305d} = \sum \text{fixed effects} + a_1 + c_k a_2 + \dots + e$$

↑  
Climatic variable (standardized)

$\text{var}(a) = G_{rn} \otimes A$

$I \sigma^2_{e_{\text{region } i}}$



$G_{rn}$  is of rank 2

## 5. more complex reaction norm models:

$$y_{305d} = \sum \text{fixed effects} + a_1 + c_k a_2 + \dots + c_p a_n + \dots + e$$

$G_{rn}$  can be of rank n or < n (for example: 2)

# Results : uni/multi variate models

- **Genetic parameters ( $h^2$ ,  $\rho_a$ ) in two situations**
  - 1 country = 1 trait

Analysis	Milk yield	Age at first calving
Univariate	<b>0.25</b>	<b>0.18</b>

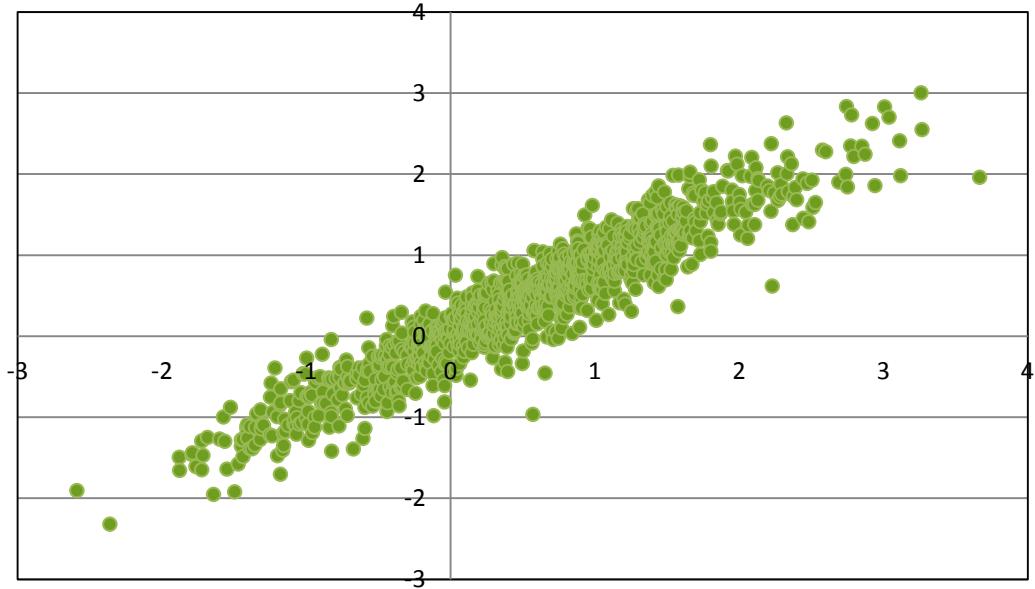
- 1 region = 1 trait → 4 traits

	(1)	0.23			0.22				
Multi-	(2)	0.75	0.36		0.62	0.18			
variante	(3)	0.87	0.83	0.25	0.74	0.82	0.18		
	(4)	0.82	0.76	0.86	0.27	0.71	0.64	0.82	0.12

# Implications = reranking !

EBV for protein yield

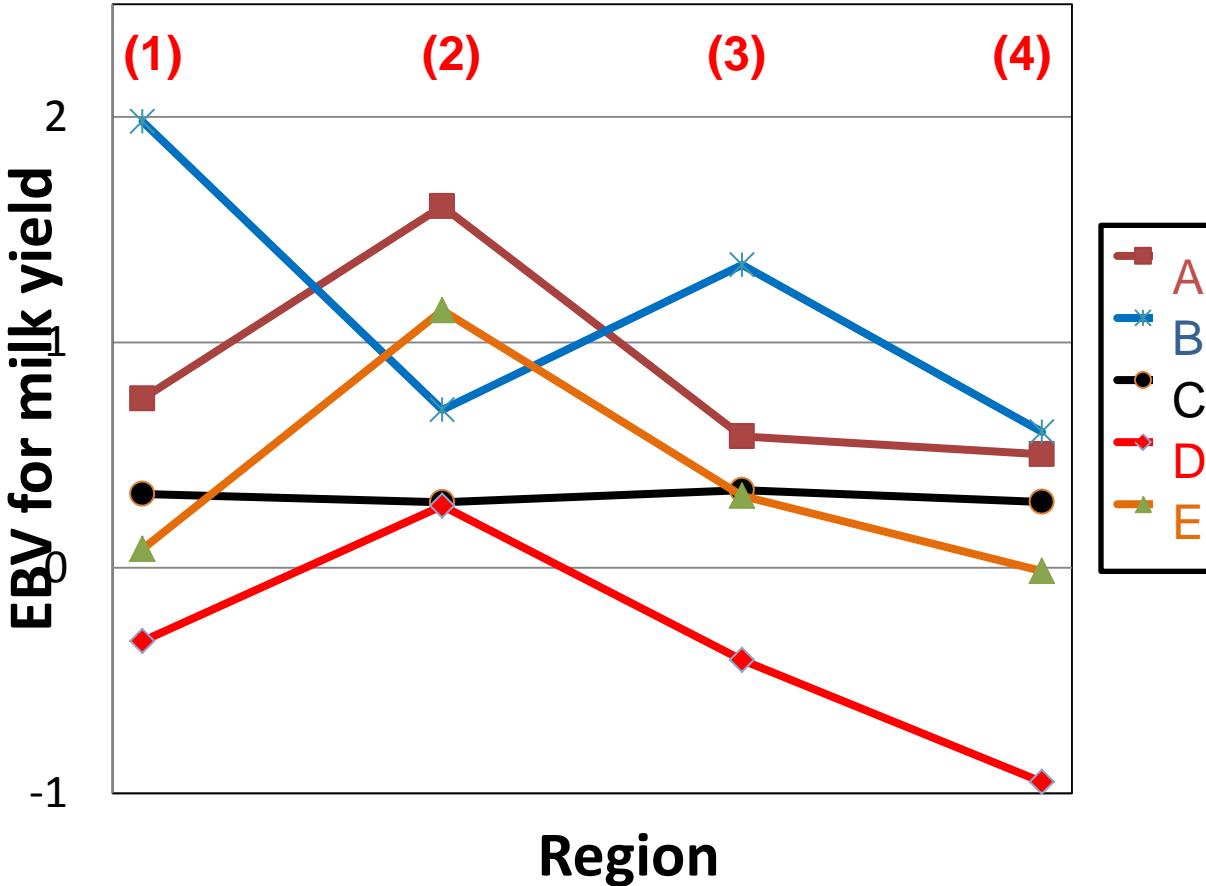
In KwaZulu Natal



EBV at national level

- The best bulls in a given region are not necessarily the best ones at national level ...

# Implications = reranking !



- Farmers are better off if they can pick the best bulls for their own region ...

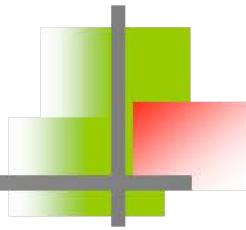
# Results : reduced rank models

## ■ Genetic parameters ( $h^2$ , $\rho_a$ ) for milk yield

region	(1)	0.23	Rank 4			0.23	Rank 3	
	(2)	0.75	0.36			0.78	0.37	
	(3)	0.87	0.83	0.25		0.85	0.85	0.25
	(4)	0.82	0.76	0.86	0.27	0.95	0.75	0.95
(+0.029)						(+0.013)		

region	(1)	0.24	Rank 2			0.23	Rank 1	
	(2)	0.77	0.37			1.00	0.36	
	(3)	0.91	0.96	0.25		1.00	1.00	0.25
	(4)	1.00	0.82	0.95	0.27	1.00	1.00	1.0
(+0.003)								

# Results : Model comparison



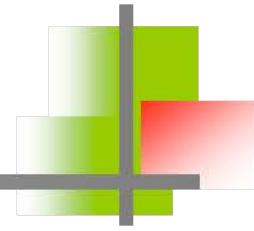
Model	BIC - BICmin
Univariate	<b>10410</b>
<b>Univariate + residual variance by region</b>	<b>1236</b>
Multivariate by region, rank 1	<b>194</b>
Multivariate by region, rank 2	<b>67</b>
Multivariate by region, rank 3	<b>39</b>
<b>Multivariate by region, rank 4</b>	<b>0</b>

# A reaction norm model using climatic variables

- Fraction (%= of the total genetic variability) explained by the (best) climatic variables

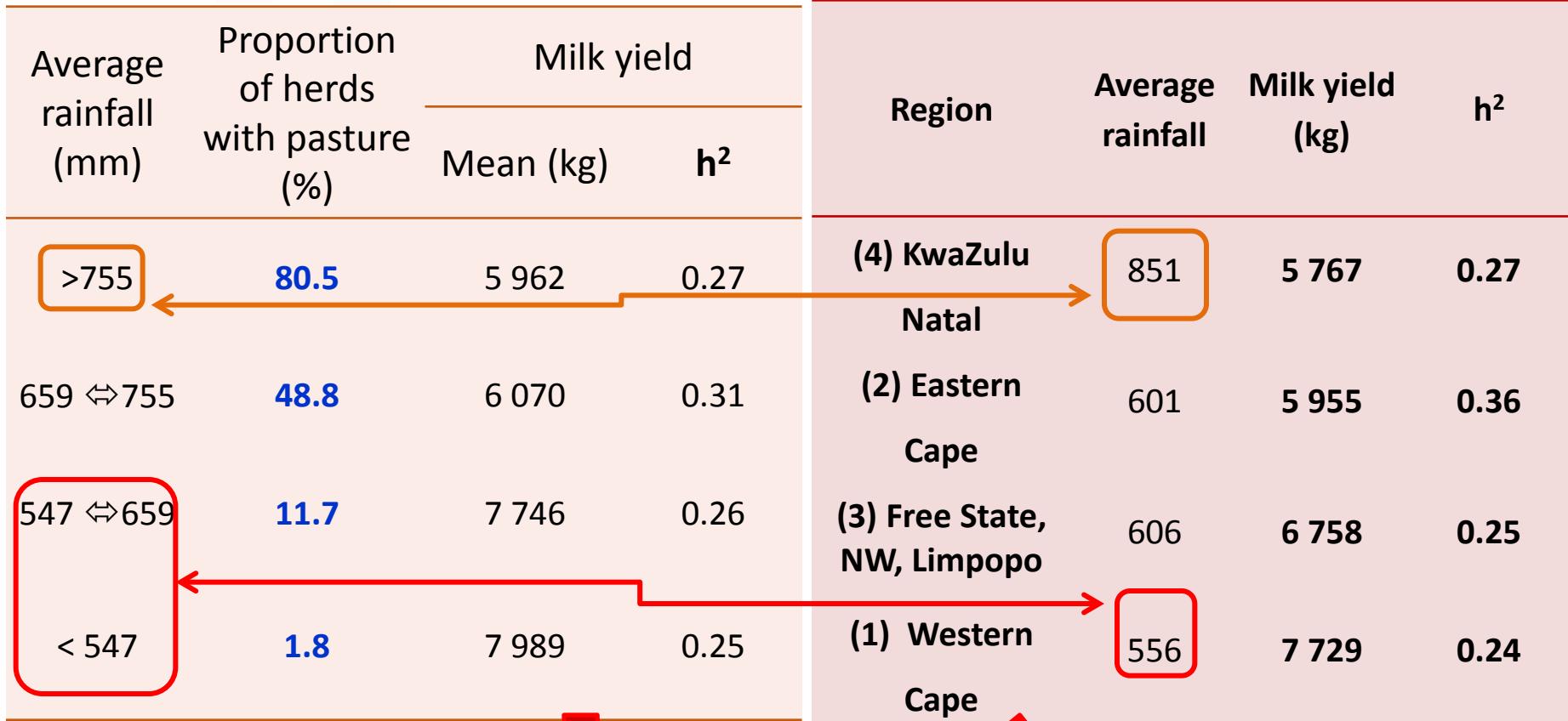
Climatic variable	Milk yield	Fat yield	Protein yield	Age at first calving
Average rainfall	9.9	9.5	10.5	9.0
Max temperature	2.6	2.8	3.4	7.3
Max Summer Temperature	1.0	1.3	0.8	26.1
Min Relative Humidity	1.0	1.0	0.9	23.3

# Results : Model comparison



Model	BIC - BICmin
Univariate	11 190
Univariate + residual variance by region	2 016
<b>Multivariate by region, rank 4</b>	<b>780</b>
<b>Reaction norm on average rainfall, rank 2</b>	<b>745</b>
<b>Reaction norm on average rainfall and max temperature, rank 3</b>	<b>0</b>

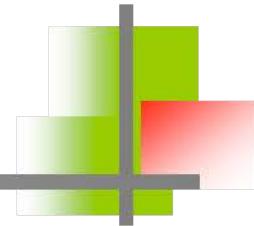
# Grouping herds in 4 groups according to average rainfall



# Conclusion

- within South Africa : as much G x E interaction as between Europe and New-Zealand ! probably related to large **climatic (rainfall) / feeding system (pasture vs TMR) differences**
- a reaction norm model based on average rainfall gives results as good as when considering 1 trait per region (« MACE type »), - possibly because rainfall « imposes » a particular feeding system
- G x E should be considered at national level in the future





***THANK YOU  
FOR YOUR ATTENTION!***