

# Finlay-Wilkinson model: regression on the environmental mean

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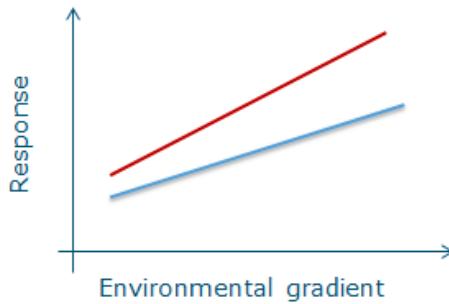
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- ❑ From ANOVA model to Finlay-Wilkinson model
- ❑ Interpretation of the model parameters
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  - Adaptability
  - Stability
- ❑ Example on Durum wheat Algeria



## Norms of reaction

- When norms of reaction are not parallel between genotypes → GxE



Issue: What to put on the X-axis?

## Finlay-Wilkinson model

### THE ANALYSIS OF ADAPTATION IN A PLANT-BREEDING PROGRAMME

By K. W. FINLAY\* and G. N. WILKINSON†

[Manuscript received January 23, 1963]

#### Summary

The adaptation of barley varieties was studied by the use of grain yields of a randomly chosen group of 277 varieties from a world collection, grown in replicated trials for several seasons at three sites in South Australia. For each variety a linear regression of yield on the mean yield of all varieties for each site and season was computed to measure variety adaptation. In these calculations the basic yields were measured on a logarithmic scale, as it was found that a high degree of linearity was thereby induced. The mean yield of all varieties for each site and season provided a quantitative grading of the environments; and from the analysis described, varieties specifically adapted to good or poor seasons and those showing general adaptability may be identified.

*Aust. J. Agric. Res.*, 1963, **14**, 742–54

Regression on the environmental mean ( $E$ )



> 1400 citations

## Finlay Wilkinson model

- Environmental variable = the *average yield across genotypes for each environment* (quality of environment).



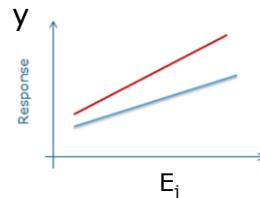
## From ANOVA to Finlay Wilkinson model

$$\underline{y}_{ij} = \mu + \textcolor{red}{G}_i + \textcolor{green}{E}_j + (\textcolor{green}{G}\textcolor{blue}{E}_{ij} + \varepsilon_{ij})$$

$$\underline{y}_{ij} = \mu + \textcolor{red}{G}_i + \textcolor{green}{E}_j + \beta_i \textcolor{green}{E}_j + \varepsilon_{ij}$$

$$\underline{y}_{ij} = \mu + \textcolor{red}{G}_i + (1 - \beta_i) \textcolor{green}{E}_j + \varepsilon_{ij}$$

$$\underline{y}_{ij} = \mu + \textcolor{red}{G}_i + \beta_i^* \textcolor{green}{E}_j + \varepsilon_{ij}$$



- The environmental main effect  $E_j$  is used as regressor.
- Uses environmental information **present in the data**:
  - $E_j$  is a measure of the environmental quality,
  - ... but no information of why.



## Finlay Wilkinson model

- ❑ Environmental variable = the *average yield across genotypes for each environment* (quality of environment).
- ❑  $y_{ij} = \mu + G_i + \beta_i^* E_j + \epsilon_{ij}$
- ❑ A popular method for estimating adaptability
  - Slope ( $\beta^*$ ) of this regression is a measure for adaptability
  - Intercept ( $\mu + G_i$ ) is a measure for general performance
- ❑ The average slope = 1
  - genotypes with slopes > 1 have **higher** than average adaptability
  - genotypes with slopes < 1 have **lower** than average adaptability

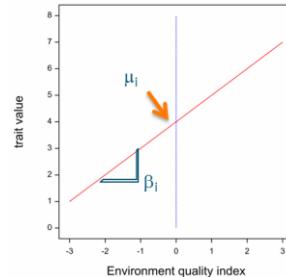
## Notation:

In the following,  $\beta$  is  $\beta^*$   
 (i.e. we drop the \* from the notation for simplicity)

## Finlay-Wilkinson model parameters

$$\underline{y}_{ij} = \mu + G_i + \beta_i E_j + \varepsilon_{ij}$$

$$\underline{y}_{ij} = \underline{\mu}_i + \beta_i E_j + \varepsilon_{ij}$$

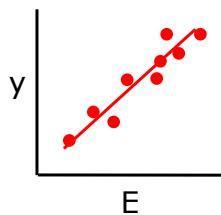


- $\mu_i$  (intercept): the expected performance of genotype  $i$  in the “average” environment.
- $\beta_i$  (slope) = the sensitivity of genotype  $i$  to the improvement in the quality of the environment (adaptability).

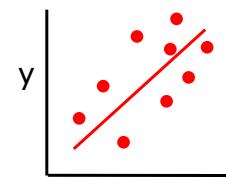


## Finlay Wilkinson model and stability

- **Eberhart and Russell stability (type 3):** deviation from predicted performance given environment
  - stable genotype = low residual variance ( $\sigma_{\varepsilon_{ij}}^2$ ) from the regression line (predicted response).



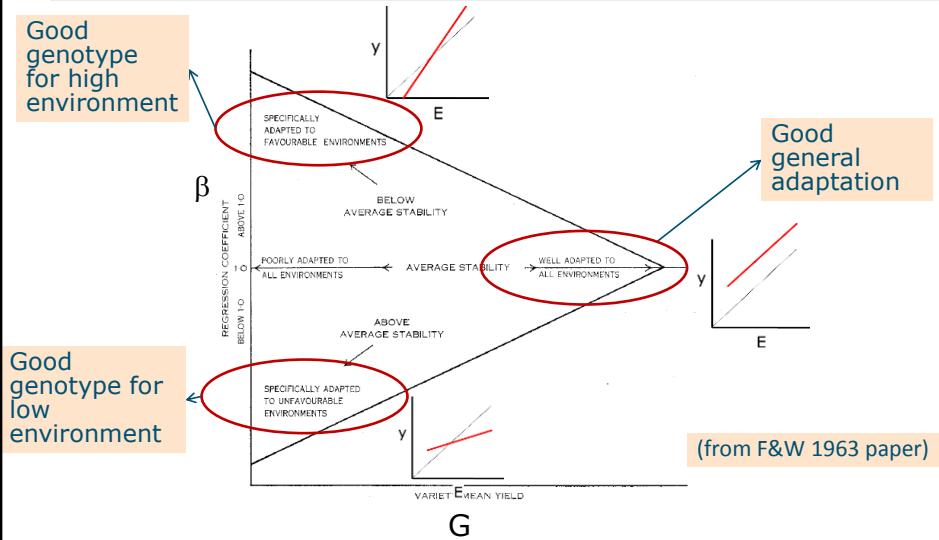
Stable,  $\sigma_{\varepsilon_{ij}}^2$  is small



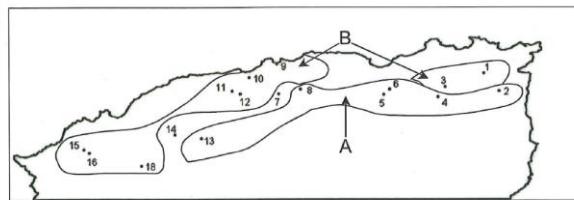
Unstable,  $\sigma_{\varepsilon_{ij}}^2$  is large



## FW parameters plotted together



## Example durum wheat Algeria



Source: Annicchiarico, 2002c.

- ❑ 24 genotypes
- ❑ 22 trials (environments), each one designed as RCBD.
  - 11 sites
  - 2 years
- ❑ Grain yield (ton/ha)

## Data set: two-way table of means

Genotypes ↓

	Environments →																							
Geno1	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	
1	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001	.001
2	ARDENT	3.641	4.462	2.485	0.715	1.894	3.217	1.087	1.662	0.961	0.646	1.424	1.819	2.139	2.267	2.286	2.305	3.14	0.965	1.649	1.509	1.648		
3	B.DUR194	3.626	3.544	2.352	1.425	2.118	1.112	1.269	1.828	1.062	0.854	1.61	2.574	1.271	2.771	2.181	2.962	2.311	3.02	1.271	1.71	1.577	1.667	
4	B.DUR195	3.626	3.544	2.352	1.425	2.118	1.112	1.269	1.828	1.062	0.854	1.61	2.574	1.271	2.771	2.181	2.962	2.311	3.02	1.271	1.71	1.577	1.667	
5	BID/WAHR	3.415	3.036	2.388	1.174	2.175	1.126	1.201	1.694	1.047	0.658	1.813	2.638	2.015	2.092	2.66	2.833	2.298	3.238	1.694	1.813	1.791	1.668	
6	BID17	3.686	1.166	2.332	1.011	1.847	1.149	1.09	1.613	0.977	0.53	1.32	1.756	1.354	2.031	2.042	2.452	2.099	3.207	1.931	1.542	0.868	1.728	
7	CHEM	3.832	1.159	2.519	1.157	2.132	1.179	1.487	1.888	1.262	0.882	1.693	2.75	1.862	2.011	2.042	2.452	2.099	3.207	1.931	1.542	0.746	1.79	
8	CHILOU	3.626	3.544	2.352	1.425	2.118	1.112	1.269	1.828	1.062	0.854	1.61	2.574	1.271	2.771	2.181	2.962	2.311	3.02	1.271	1.71	1.577	1.667	
9	EIDEY	3.524	2.68	2.387	1.056	1.931	1.117	1.27	1.85	1.201	0.715	1.86	2.368	1.621	2.13	2.299	2.318	2.205	3.189	1.569	1.434	2.023	1.681	
10	GTADUR	3.924	1.009	2.58	1.202	2.361	1.31	1.275	1.92	1.074	0.8	1.841	2.689	1.598	2.043	2.894	2.874	2.3	3.296	1.53	1.715	1.851	1.813	
11	HEIDRØD	3.703	1.009	2.58	1.202	2.361	1.31	1.275	1.92	1.074	0.8	1.841	2.689	1.598	2.043	2.894	2.874	2.3	3.296	1.53	1.715	1.851	1.813	
12	HEIDRØD	2.472	6.79	2.057	0.981	1.79	1.06	0.939	1.615	0.969	0.325	1.411	1.688	0.864	2.181	3.005	2.453	2.17	3.028	1.393	1.379	1.316	1.713	
13	INARATEB	3.103	2.7	2.071	1.029	1.92	1.07	0.683	1.349	0.705	0.726	1.411	2.031	1.667	1.982	2.353	2.828	2.212	3.11	1.629	1.38	1.96	1.566	
14	INARATEB	3.051	1.784	2.396	1.268	1.977	1.107	1.08	1.335	1.001	0.659	1.807	2.398	1.507	2.376	2.576	2.848	3.341	1.6	1.475	1.272	1.531		
15	INARATEB	3.051	1.784	2.396	1.268	1.977	1.107	1.08	1.335	1.001	0.659	1.807	2.398	1.507	2.376	2.576	2.848	3.341	1.6	1.475	1.272	1.531		
16	MEXICALI	3.103	3.2	2.104	0.801	2.082	1.031	1.264	1.655	1.137	0.713	1.909	2.056	2.063	2.301	2.53	2.68	2.3	3.262	2.171	1.783	1.389	1.668	
17	OZNARAB	2.621	4.467	2.171	0.821	1.811	1.15	1.087	1.826	0.948	0.448	1.52	1.388	1.198	1.522	2.115	2.162	2.233	2.316	0.896	1.579	1.012	1.669	
18	OZNARAB	2.621	4.467	2.171	0.821	1.811	1.15	1.087	1.826	0.948	0.448	1.52	1.388	1.198	1.522	2.115	2.162	2.233	2.316	0.896	1.579	1.012	1.669	
19	OZNARAB	3.604	1.791	2.212	1.246	2.195	1.184	1.227	1.675	1.21	0.844	1.493	2.521	1.762	1.821	2.493	2.421	2.389	3.094	1.48	1.598	1.939	1.661	
20	POLON/28	2.829	3.954	2.191	1.05	1.821	1.113	1.099	1.655	1.006	0.503	1.473	1.752	1.076	2.185	2.171	2.339	2.271	3.177	0.888	1.238	1.041	1.715	
21	POLON/28	2.592	1.083	2.098	1.49	2.004	1.117	1.001	1.758	1.087	0.524	1.45	1.707	0.917	2.191	1.88	2.509	2.178	3.038	0.905	1.431	1.77	1.433	
22	POVITRON	3.075	3.87	2.47	1.233	2.478	1.005	0.845	1.679	1.238	0.741	1.839	2.804	1.962	2.384	2.837	2.218	2.895	3.244	1.58	1.397	1.697	1.878	
23	SIMETO	3.075	3.87	2.47	1.233	2.478	1.005	1.17	1.757	1.238	0.741	1.839	2.804	1.962	2.384	2.837	2.218	2.895	3.244	1.58	1.397	1.697	1.878	
24	VITRON	3.404	3.286	2.321	0.72	1.92	1.105	1.176	1.717	1.208	0.681	1.804	2.364	1.632	2.596	2.622	2.446	2.114	3.178	1.894	1.387	1.951	1.709	
25	WAHA	3.481	1.09	2.481	0.828	2.155	1.127	1.299	1.84	0.98	0.755	1.647	2.449	1.891	2.111	2.406	2.835	2.274	3.117	1.907	1.571	2.215	1.721	

## Environmental index

- Environmental variable = Environmental main effect ( $E_j$ )
  - As a deviation from the average

$$y_{ij} = \mu + G_i + E_j + (GE_{ij} + e_{ij})$$

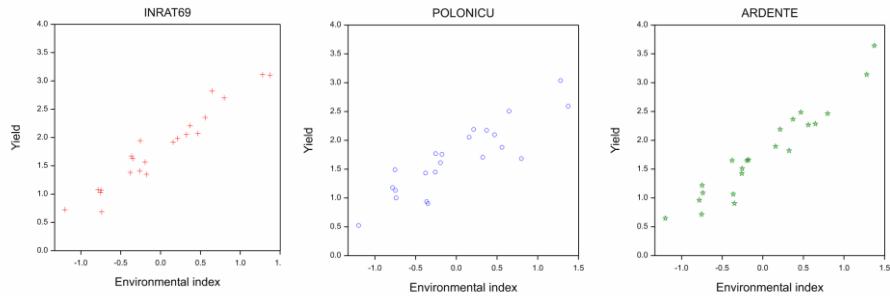
- Environments can be ranked on  $E$ 
  - $E > 0$ : better than average
  - $E < 0$ : worse than average

- $E$  = "Environmental Index"

env	E
s01_y1	1.3736
s14_y2	1.2801
s01_y2	0.7984
s12_y2	0.6474
s12_y1	0.5603
s02_y1	0.4688
s14_y1	0.3721
s10_y2	0.3238
s11_y2	0.2137
s04_y1	0.1575
s05_y2	-0.1759
s18_y2	-0.1951
s18_y1	-0.2540
s10_y1	-0.2587
s16_y1	-0.3489
s11_y1	-0.3641
s16_y2	-0.3775
s05_y1	-0.7386
s04_y2	-0.7470
s02_y2	-0.7532
s07_y1	-0.7825
s07_y2	-1.2001

WAGENINGEN UR  
For quality of life

## Yield versus environmental index (E)



- ❑ Three genotypic responses
  - Differences / similarities?



## FW model for genotype POLONICU

### Regression analysis

Response variate: yld[20]  
Fitted terms: Constant, EnvIndex

### Interpretation of the model parameters?

### Summary of analysis

Source	d.f.	s.s.	m.s.	v.r.
Regression	1	6.080	6.08032	
Residual	20	1.806	0.09032	
Total	21	7.887	0.37556	67.32

Percentage variance accounted for 76.0  
Standard error of observations is estimated to be 0.301.

Message: the following units have large standardized residuals.

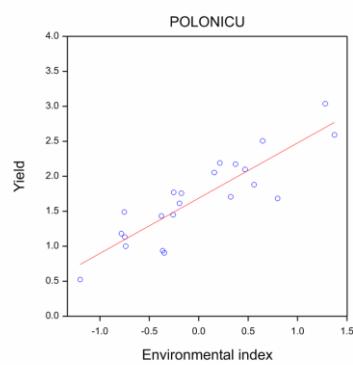
Unit	Response	Residual
2	1.683	-2.24

Message: the following units have high leverage.

Unit	Response	Leverage
1	2.592	0.239
10	0.524	0.193
18	3.038	0.214

### Estimates of parameters

Parameter	estimate	s.e.	t(20)
Constant	1.6871	0.0641	26.33
EnvIndex	0.7897	0.0962	8.20



## POLONICU VS ARDENTE

**Regression analysis**

Response: ARDENTE higher stability (smaller var( $\varepsilon$ ))

Source	d.f.	s.s.	m.s.	v.r.
Regression	1	6.080	6.08032	67.32
Residual	20	1.806	0.09032	
Total	21	7.887	0.37556	

Percentage variance accounted for 76.0  
Standard error of observations is estimated to be 0.301.

Source	d.f.	s.s.	m.s.	v.r.
Regression	1	11.671	11.67059	225.58
Residual	20	1.035	0.05174	
Total	21	12.705	0.60501	

Percentage variance accounted for 91.4  
Standard error of observations is estimated to be 0.227.

*Message: the I* Unit ARDENTE (slightly) higher general adaptation ( $\mu_i$ ) *individuals.*

*Message: the following units have high leverage.*

Unit	Response	Leverage
1	2.592	0.239
10	0.524	0.193
18	3.038	0.214

*Message: the following units have high leverage.*

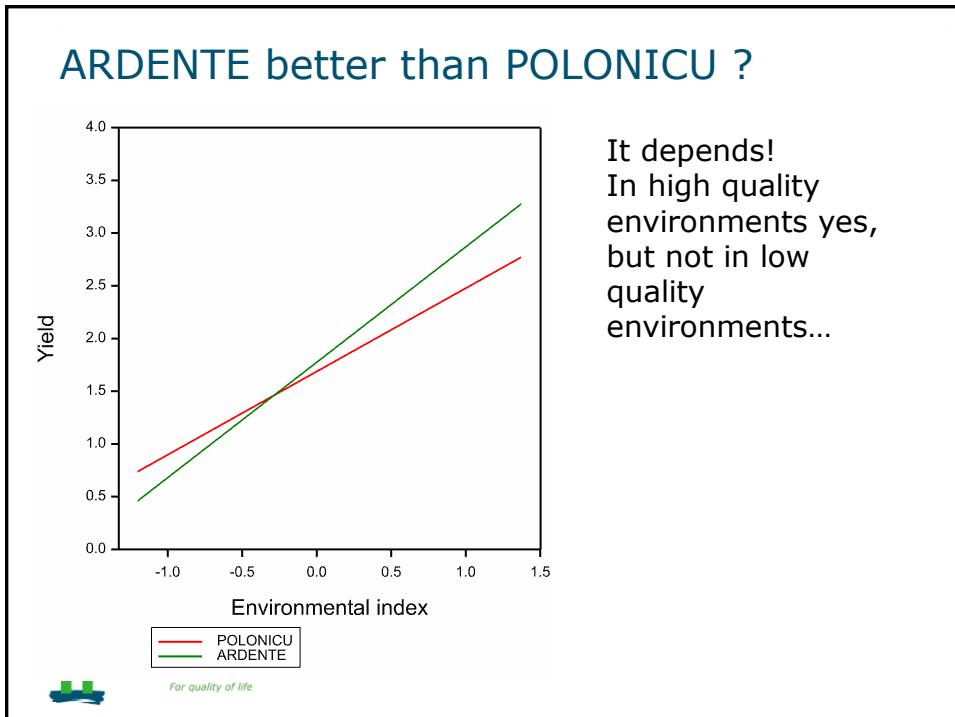
Unit	Response	Leverage
1	3.641	0.239
10	0.646	0.193
18	3.140	0.214

**Estimates of parameters**

Parameter	estimate	s.e.	t(20)
Constant	1.6871	0.0641	26.33
EnvIndex	0.7897	0.0962	8.20

Parameter	estimate	s.e.	t(20)
Constant	1.7744	0.0485	36.59
EnvIndex	1.0940	0.0728	15.02

WAGENINGEN  ARDENTE higher adaptability ( $\beta$ )



## From ANOVA to FW model

### Accumulated analysis of variance

	d.f.	s.s.	m.s.	v.r.	F pr.
Change	23	13.47438	0.58584	10.34	<.001
+ Geno	21	233.96713	11.14129	196.63	<.001
+ Env			0.05666		
Residual	483	27.36739			
Total	527	274.80890	0.52146		

### Analysis of variance

Source	d.f.	s.s.	m.s.	v.r.	F pr.
Genotypes	23	13.4744	0.5858	10.51	<0.001
Environments	21	233.9671	11.1413	199.95	<0.001
Sensitivities	23	1.7364	0.0755	1.35	0.127
Residual	460	25.6310	0.0557		
Total	527	274.8089	0.5215		

- Partition of the ANOVA “residual” (GxE+error) into:
  - Heterogeneity of slopes (=“sensitivities”)
  - Residual
- (a little) part of the GxE has become predictable



## Full results

$$\underline{y}_{ij} = \mu_i + \beta_i E_j + \varepsilon_{ij}$$

### Sorted sensitivity estimates

Genotype	Sensitivity	s.e.	Mean	s.e.	Var( $\varepsilon$ )
1) POLONICU	0.7897	0.07559	1.687	0.05033	0.09032
2) O.ZENADI	0.8336	0.07559	1.575	0.05033	0.09884
3) HEBDA03	0.8628	0.07559	1.617	0.05033	0.06455
4) MBBACHIR	0.8679	0.07559	1.611	0.05033	0.08581
5) BIDI17	0.9206	0.07559	1.685	0.05033	0.05923
6) POLON/ZB	0.9407	0.07559	1.661	0.05033	0.05514
7) OFANTO	0.9724	0.07559	1.990	0.05033	0.04635
8) EIDER	0.9863	0.07559	1.929	0.05033	0.02086
9) OUMRAB19	0.9935	0.07559	1.940	0.05033	0.02562
10) INRAT69	1.0092	0.07559	1.812	0.05033	0.03523
11) MEXICALI	1.0138	0.07559	1.969	0.05033	0.07472
12) DUILIO	1.0158	0.07559	1.973	0.05033	0.05176
13) KEBIR	1.0223	0.07559	1.832	0.05033	0.02185
14) SAHEL77	1.0292	0.07559	2.076	0.05033	0.11352
15) BELIKH02	1.0326	0.07559	1.934	0.05033	0.03979
16) B.DUR194	1.0464	0.07559	1.986	0.05033	0.06401
17) CHENS	1.0647	0.07559	2.056	0.05033	0.02646
18) WAHA	1.0663	0.07559	2.001	0.05033	0.04165
19) SIMETO	1.0685	0.07559	2.015	0.05033	0.07589
20) BIDI/WAHA	1.0694	0.07559	2.021	0.05033	0.02367
21) HEBD/GDO	1.0752	0.07559	1.746	0.05033	0.04052
22) VITRON	1.0812	0.07559	1.965	0.05033	0.04987
23) ARDENTE	1.0940	0.07559	1.774	0.05033	0.05174
24) GTADUR	1.1395	0.07559	2.084	0.05033	0.02416



## Summary: Finlay-Wilkinson model

- ❑ In FW-regression the environments is the mean performance of all genotypes in that environment ( $E$ )
- ❑ Genotypes are characterized in terms of:
  - intercept (general performance,  $\mu_i$ )
  - slope (adaptability,  $\beta$ )
  - deviations from regression (stability,  $\text{var}(\varepsilon)$ )
- ❑ Prediction of variety performance in unobserved environment is possible when  $E$  is known

