Bad Statistics



Visualization and Presentation of Statistics, Open University, 18 May 2011 ... John Gower was head of the Statistics Department at Rothamsted Experimental Station (and I was a member of the department).

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John disagreed rather strongly.

From a conference poster in December 2010



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Is this always appropriate?

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An experiment was conducted to compare two protective dyes (B and C) for metal, both with each other and with 'no dye' (A).

Ten braided metal cords were broken into three pieces. The three pieces of each cord were randomly allocated to the three treatments. Thus the cords were blocks.

After the dyes had been applied, the cords were left to weather for a fixed time, then their strengths were measured.

Simple display of results for dyes on metal cords



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Simple display of results for dyes on metal cords



But these standard deviations include the variability between cords!

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raw data	mean	96.67	99.29	101.62
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Isn't this a more useful visual summary?

The "antenna" part of the "bar-and-antenna" diagram is completely misleading.

Then the estimate of each treatment mean is no longer the same as the mean of the data for that treatment,

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We can summarize the ANOVA table graphically by scaling the lines in the Hasse diagram.

Metal cords: Hasse diagram of models



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Experimental unit = jar.

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RichnessTreatmentLevelA, ..., Fmonoculture12 of type A1AB, ..., EFduoculture6 of A, 6 of B2ABC, ..., DEFtriculture4 of A, 4 of B, 4 of C3

Experimental unit = jar.

				Richness
	Treatment			Level
6	A, \ldots, F	monoculture	12 of type A	1
15	AB, \ldots, EF	duoculture	6 of A, 6 of B	2
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D' 1

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The experiment was carried out in 4 blocks of 41 jars.

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The editor accepted our arguments.

What models did we fit?

The biologist fitted the model 'Richness' with 3 parameters, one for each level of richness,

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I suggested the model 'Type' with 6 parameters $\alpha_A, \ldots, \alpha_F$:



In other words, if there are x_i shrimps of type *i* then

$$\mathbb{E}(Y) = \sum_{i=1}^{6} a_i x_i \qquad \text{where } 12a_i = \alpha_i$$

 $(\sum x_i = 12 \text{ always, so no need for intercept.})$









Treatment Fichness * Type Richness + Type Scale: $3 \times$ residual mean square Richness Constant

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Treatment Richness	s * Type
Richness + Type • Type	Conclusions: The model Richness does not explain the data. The model Type explains the data well.
	Scale: 3 × residual mean square
Richness • Constar	」 と・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・

Treatment Fichness	s * Type
Richness + Type • Type	Conclusions: The model Richness does not explain the data. The model Type explains the data well. There is no evidence that any larger model does any better.
	Scale:
	5 × residuai mean square
Richness 🔥 Constar	t ← □ ▷ < 클 ▷ < 클 ▷ < 클 ▷ < 클 > < 클 > < 3,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0,0

Treatment Richness * Type Is such a scaled Hasse diagram a good way of displaying the ANOVA table when there is only one relevant residual mean square? Scale: $3 \times$ residual mean square ▲□▶ ▲□▶ ▲ □▶ ▲ □▶ ▲□ ● ● ● Richness Constant

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