## NUTRITIONAL TRIALS AND DATA ANALYSIS



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## STATISTICS

- Research tool
- Deals with the collection, organization, analysis, and interpretation of data.
- Useful in drawing meaningful conclusions from a set of data
- You can make the data look the way you like, if you know how to do it ( misuse)
- Should know why you are using?
- Should have focused questions to answer rather than reporting all possible relationships among all possible treatments


## ANOVA

Analysis of Variance / Partitioning of variance
Understanding Variance:

- All individuals in a population are not similar
- They differ from each other.
- Population forms a bell shape curve
- We want to know whether this dissimilarity (variation) is a chance variation or otherwise
- Inherent variation of due to other factors
- Proportion of variation due to known variables is analysed


## NUTRITIONAL TRIALS

- Growth Trials
- Production Trials
- Testing different treatments on any other aspect(s)


## EXPERIMENTAL DESIGNS

- Completely Randomized Designs (CRD)
- Randomized Complete Block Designs (RCBD)
- Latin Square Designs (LSD)
- Factorial Experiments


## LIMITATIONS OF EACH DESIGN

- CRD

When experimental units are homogenous
Have less variation
Randomization carried out using Random Number Tables

- RCBD
when experimental units can meaningfully grouped
Such groups are called blocks
- LSD

Double grouping
Where two major sources of variation are present

## ANOVA for CRD

When we have 4 treatments and 4 replicates

| Source of variation | Degree of freedom | Degree of <br> freedom |
| :--- | :---: | :---: |
| Treatment | $(\mathrm{t}-1)$ | 3 |
| Error | $\mathrm{t}(\mathrm{r}-1)$ | 12 |
| Total | $(\mathrm{n}-1)$ | 15 |

## ANOVA for RCBD

## When we have 4 treatments and 4 replicates

| Source of variation | Degree of freedom | Degree of <br> freedom |
| :--- | :---: | :---: |
| Treatment | $(\mathrm{t}-1)$ | 3 |
| Blocks | $(\mathrm{b}-1)$ | 3 |
| Error | $(\mathrm{t}-1)(\mathrm{b}-1)$ | 9 |
| Total | $(\mathrm{n}-1)$ | 15 |

## ANOVA for $2 \times 2$ factorial arrangement

 When we have 4 treatments and 4 replicates| Source of variation | Degree of freedom | Degree of <br> freedom |
| :--- | :---: | :---: |
| Treatment | $(\mathrm{t}-1)$ | 3 |
| Factor A | $(\mathrm{a}-1)$ | 1 |
| Factor B | $(\mathrm{b}-1)$ | 1 |
| $\mathbf{A x B}$ | $(\mathrm{a}-1)(\mathrm{b}-1)$ | 1 |
| Error | $\mathrm{ab}(\mathrm{r}-1)$ | 12 |
| Total | $(\mathrm{n}-1)$ | 15 |

## ANOVA for factorial experiment

## Two factor factorial $2 \times 2$ with 12 replicates each

| Source of variation | Degree of freedom | Degree of <br> freedom |
| :--- | :---: | :---: |
| Factor A | $(\mathrm{a}-1)$ | 1 |
| Factor B | $(\mathrm{b}-1)$ | 1 |
| Interaction AB | $(\mathrm{a}-1)(\mathrm{b}-1)$ | 1 |
| Error | $\mathrm{ab}(\mathrm{r}-1)$ | 44 |
| Total | $(\mathrm{n}-1)$ | 47 |

## ANOVA for Latin Square Design

## When we have 4 treatments and 4 replicates

| Source of variation | Degree of freedom | Degree of <br> freedom |
| :--- | :---: | :---: |
| Treatments | $(\mathrm{r}-1)$ | 3 |
| Blocks (animals) | $(\mathrm{r}-1)$ | 3 |
| Periods | $(\mathrm{r}-1)$ | 3 |
| Error | $(\mathrm{r}-1)(\mathrm{r}-2)$ | 6 |
| Total | $(\mathrm{n}-1)$ | 15 |

## ANOVA for Latin Square Design

Four treatments and 4 replicates with $2 \times 2$ factorial arrangement

| Source of variation | Degree of freedom | Degree of <br> freedom |
| :--- | :---: | :---: |
| Treatments | $(\mathrm{r}-1)$ | 3 |
| Factor A <br> Factor B <br> A x B | $(\mathrm{a}-1)$ <br> $(\mathrm{b}-1)$ <br> $(\mathrm{a}-1)(\mathrm{b}-1)$ | 1 |
| Blocks (animals) | $(\mathrm{r}-1)$ | 1 |
| Periods | $(\mathrm{r}-1)$ | 3 |
| Error | $(\mathrm{r}-1)(\mathrm{r}-2)$ | 3 |
| Total | $(\mathrm{n}-1)$ | 6 |

## Example I

## Effect of bST and Enzose on DMI and production performance of buffaloes

Abu Bakar Sufyan's MSc Data
Two levels of bST: bST0 and bST1 ( 250 mg )
Three levels of Enzose: ENZ1, ENZ2 and ENZ3 $(0,20,40)$
Two replicates per treatment
Data analysis
$2 \times 2$ factorial arrangements



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|  |  | －$f_{x}$ |  |  |  |  |  |  |  |  |  |
|  | E | F | G | H | I | J | K | L | M | N | ヘ |
| 31 |  | bST1 |  |  |  |  |  |  |  |  |  |
| 32 |  | Count | 2 | 2 | 2 | 6 |  |  |  |  |  |
| 33 |  | Sum | 25.99 | 26.72 | 17.66 | 70.37 |  |  |  |  |  |
| 34 |  | Average | 12.995 | 13.36 | 8.83 | 11.72833 |  |  |  |  |  |
| 35 |  | Variance | 0.49005 | 0.1152 | 0.5202 | 5.291937 |  |  |  |  |  |
| 36 |  |  |  |  |  |  |  |  |  |  |  |
| 37 |  | Tota！ |  |  |  |  |  |  |  |  |  |
| 38 |  | Count | 4 | 4 | 4 |  |  |  |  |  |  |
| 39 |  | Sum | 49.14 | 50.24 | 33.53 |  |  |  |  |  |  |
| 40 |  | Average | 12.285 | 12.56 | 8.3825 |  |  |  |  |  |  |
| 41 |  | Variance | 0.998833 | 1.2 | 0.660825 |  |  |  |  |  |  |
| 42 |  |  |  |  |  |  |  |  |  |  |  |
| 43 |  |  |  |  |  |  |  |  |  |  |  |
| 44 |  | ANOVA |  |  |  |  |  |  |  |  |  |
| 45 |  | Source of Variation | SS | $d f$ | MS | $F$ | $P$－value | F crit |  |  |  |
| 46 |  | Sample | 5.109075 | 1 | 5.109075 | 9.574878 | 0.02127 | 5.987374 |  |  |  |
| 47 |  | Columns | 43.67552 | 2 | 21.83776 | 40.92597 | 0.000319 | 5.143249 |  |  |  |
| 48 |  | Interaction | 0.26835 | 2 | 0.134175 | 0.251456 | 0.785471 | 5.143249 |  |  |  |
| 49 |  | Within | 3.20155 | 6 | 0.533592 |  |  |  |  |  |  |
| 50 |  |  |  |  |  |  |  |  |  |  |  |
| 51 |  | Total | 52.25449 | 11 |  |  |  |  |  |  |  |
| 52 |  |  |  |  |  |  |  |  |  |  |  |
| 53 |  |  |  |  |  |  |  |  |  |  | $v$ |
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- 



Jata one;
Input ENLEVEL\$ BSTLEVEL\$ DMI;
cards;

| Enzose1 | bST0 | 12.07 |
| :--- | :--- | :--- |
| Enzose1 | bST0 | 11.08 |
| Enzose1 | bST1 | 12.5 |
| Enzose1 | bST1 | 13.49 |
| Enzose2 | bST0 | 12.44 |
| Enzose2 | bST0 | 11.08 |
| Enzose2 | bST1 | 13.6 |
| Enzose2 | bST1 | 13.12 |
| Enzose3 | bST0 | 8.51 |
| Enzose3 | bST0 | 7.36 |
| Enzose3 | bST1 | 8.32 |
| Enzose3 | bST1 | 9.34 |

; ${ }^{\text {Proc GLM; }}$
Class ENLEVEL BSTLEVEL;
Model DMI = ENLEVEL BSTLEVEL ENLEVEL*BSTLEVEL;
means ENLEVEL BSTLEVEL ENLEVEL*BSTLEVEL /duncan;
1smeans ENLEVEL BSTLEVEL ENLEVEL*BSTLEVE/stderr;
Title 'Stat Analysis using factorial arrangements';
run;


```
F SAS - [OUTPUT - (Untitled) PROC GLM running]
```


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$\square$

Dependent Variable: DMI

| Source | DF |
| :--- | ---: |
| Model | 5 |
| Error | 6 |
| Corrected Total | 11 |
|  | R-Square |
|  | 0.938732 |


| Source | DF | Type I SS |
| :--- | ---: | ---: |
| ENLEVEL | 2 | 43.67551667 |
| BSTLEVEL | 1 | 5.10907500 |
| ENLEVEL*BSTLEVEL | 2 | 0.26835000 |
| Source | DF |  |
|  |  |  |
| ENLEVEL | 2 | 43.67551667 |
| BSTLEVEL | 1 | 5.10907500 |
| ENLEVEL*BSTLEVEL | 2 | 0.26835000 |

Sum of Squares
49.05294167
3.20155000
52.25449167
C.V.
6.595202

Type I SS
43.67551667 5.10907500 0.26835000

Type III SS
5.109075
0.26835000

| Mean Square | F Value | Pr >F |
| ---: | ---: | ---: |
| $\mathbf{9 . 8 1 0 5 8 8 3 3}$ | 18.39 | 0.0014 |
| 0.53359167 |  |  |
|  |  |  |
| Root MSE |  | DMI Mean |
| 0.7304735 | 11.07583333 |  |


| Mean Square | F Value | Pr >F |
| ---: | ---: | ---: |
| 21.83775833 | 40.93 | 0.0003 |
| 5.10907500 | 9.57 | 0.0213 |
| 0.13417500 | 0.25 | 0.7855 |
|  |  |  |
| Mean Square | F Value | Pr >F |
|  |  |  |
| 21.83775833 | 40.93 | 0.0003 |
| 5.10907500 | 9.57 | 0.0213 |
| 0.13417500 | 0.25 | 0.7855 |

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## Effect of BST and Enzose on DMI in

 buffaloes

## Example II

## Effect of Intake level and forage source on kinetics of fibre digestion



## $4 \times 4$ Latin Square design

$2 \times 2$ factorial Arrangement
Factor I: Forage Source
Factor II: Intake level

## Model Statement

proc glm;
class anim per trmt;
model dm = anim per trmt;
contrast 'grass vs leg+grass' trmt +1-1 +1 -1; contrast 'restrict vs ad lib' trmt +1 +1-1 -1; contrast 'interaction' trmt +1-1-1 +1;
Ismeans trmt/stderr;
means trmt/duncan;
run;
File: Latin Square Factorial Intake
-

* ${ }^{*}$ File Edit View Locals Globals Options Window Help

81 AGA $17.625 \quad 16.04820 \quad 14.92092 \quad 10.13611 \quad 6.606400 \quad 1.206767 \quad 5.3996343 .529713$
72 AGA 22.5520 .5434719 .0875212 .783228 .5387551 .5072387 .0315174 .244469
63 AGA $20 \quad 18.15679 \quad 16.83420 \quad 11.15296 \quad 7.3534341 .305841 \quad 6.047593 \quad 3.799528$
54 AGA 22.320 .0008217 .9529312 .828118 .3577111 .4050496 .9526624 .470405
$\begin{array}{lllllllllllllllll}6 & 1 & \text { RG } & 10 & 9.127073 & 8.590611 & 6.192421 & 3.992915 & 0.689663 & 3.303252 & 2.199505\end{array}$
52 RG 109.0791988 .5650626 .2403964 .0795630 .6605863 .4189772 .160832
83 RG $109.0875658 .5216126 .1234383 .937581 \quad 0.6370843 .3004972 .185855$
74 RG $108.9168708 .0509916 .4135084 .066170 \quad 0.6176523 .448518 \quad 2.347338$
51 RGA 109.0935758 .4552325 .821648 3.785081 $0.678893 \quad 3.1061882 .036566$
82 RGA $109.0620348 .447763 \quad 5.8205073 .8392570 .6594153 .1798411 .981248$
73 RGA 109.0499978 .4058735 .7194213 .7422540 .6514823 .0907711 .977165
64 RGA $108.9218618 .0414545 .8832353 .8104600 .618828 \quad 3.1916322 .072775$ ;
proc glm;
class anim per trmt;
model dm = anim per trmt;
contrast 'grass vs leg+grass' trmt +1 -1 +1 -1;
contrast 'restrict vs ad lib' trmt +1 +1 -1 -1;
contrast 'interaction' trmt +1-1-1 +1;
lsmeans trmt/stderr:
means trmt/duncan;
run;
$\square$


## Effect of feed intake level and forage source on Kinetics of fibre digestion.. in Beef cattle



## Example III

## Effect of different feeding regimens on the growth performance of Sahiwal Calves



## STAT ANALYSIS

## DIFFERENT OPTIONS:

CRD
Trtmt I = Milk and SR
Trtmt II = Milk and Hay
Trtmt III = MR and SR
Trtmt IV = MR and Hay
Birth weight as Covariance????

## RCBD

## Milk and Milk Replacer

## Sex as Blocks

## CRD

$2 \times 2$ Factorial Arrangement
Factor I: Liquid Diet, Milk vs milk replacer
Factor II: Starter ration+ Hay vs Hay only

## MODEL STATEMENTS IN SAS

## CRD

Effect of different feeding regimens: milk and MR with or without SR
Proc GLM;
Class trt sex;
Model wwt TWGain DWGain TMilk FCR = trt sex bwt;
contrast 'Milk vs CMR' TRT -1-1 +1 +1;
contrast 'Fodder Vs Concen' TRT +1-1 +1-1;
means trt sex /duncan;
Ismeans trt sex /stderr;
Title 'stat analysis using CRD';
run;

## Model Finally used

## Yijkl $=\mu+$ F1i +F2j + (F1 xF2)ij + BWTk + calfl + eijkl

## Model Statement in SAS

proc mixed;
class fone ftwo id;
model DWGain = fone|ftwo bwt;
random id(fone*ftwo);
Ismeans fone|ftwo / bylevel om pdiff;
run;

File: SWL PI mixed models weight etc.sas
Output STAT mixed models.excel

## Performance of Sahiwal calves given different dietary treatments

| Parameters | Milk vs MR |  | SR vs Hay |  | Milk |  | MR |  | F1 | F2 | F1*F2 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Milk | MR | SR | Hay | SR | Hay | SR | Hay |  |  |  |
| Weaning weights (kg) | $52 \pm .8$ | $35 \pm .8$ | $49 \pm .8$ | $38 \pm .8$ | $56 \pm 1$ | 47さ1 | $40 \pm 1$ | $30 \pm 1$ | 0.0001 | 0.0001 | 0.66 |
| Total weight gain (kg) | $30.0 \pm .8$ | $14 \pm .8$ | 26 $\pm .8$ | $18 \pm .8$ | $34 \pm 1$ | $26 \pm 1$ | $18 \pm 1$ | $10 \pm 1$ | 0.0001 | 0.0001 | 0.66 |
| Daily growth rate (g/d) | $357 \pm 9$ | 162 $\ddagger 9$ | 311さ9 | 208 $\pm 9$ | $401 \pm 13$ | $310 \pm 13$ | $214 \pm 13$ | $115 \pm 13$ | 0.0001 | 0.0001 | 0.67 |

## GROWTH TRIALS

- Repeated measure analysis
- What does it mean?


## Model Statement

Yijklm $=\mu+$ Sex $i+F 1 j+F 2 k+W /+(S E X \times F 1 \times F 2 \times W) i j k l+C a l f m+$ eijklm

## Model Statement in SAS

proc mixed;
class sex fone ftwo id wk;
model wt = sex|fone|ftwo|wk;
random id(fone*ftwo);
repeated wk / sub=id(fone*ftwo) type = ar(1);
Ismeans sex|fone|ftwo|wk / bylevel om pdiff;
run;
File: SWL PI mixed model growth curve

Growth Curve of Sahiwal Calves on different preweaning dietary regimens


## Example IV

Economic feasibility of raising Lohi sheep and Beetal goats for meat production under high input system

Effect of different protein levels on the performance of Lohi Sheep with or without ionophores and Probiotics

Treatments
Fodder
Concentrate
LP MP HP
With or without lonophores
With or without Probiotics

## Treatment plan

| Fodder |  |  |  | Ionophores |  |  | Probiotics |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | LP | MP | HP | LP | MP | HP | LP | MP | HP |

## How to analyze this data?

- Analyze separately: delete Fodder and analyze the rest using $2 \times 3$ factorial design
- Imbalance design?
- CRD?
- Nested design?
- Fodder Vs Concentrate
- Ionophores vs probiotics
- Concentrate vS lonophores or Probiotics
- Linear Response?
- Quadratic Response?


## Model Statement in SAS

- proc glm;
- class trmt;
- model TDMI DMI DMIBW CPI NDFI ADFI TGAIN DGAIN FCR FEEDC ECONO = trmt;
- contrast 'Fodder vs concentrates' trmt +1 +1 +1 +1 +1 +1 +1 +1 +1 -9;
- contrast 'Conc vs l+P' trmt -2 -2 -2 +1 +1 +1 +1 +1 +1 0;
- contrast 'I vs P' trmt $000-1-1-1+1+1+10$;
- contrast 'Linear conc' trmt -1 $0+10000000$;
- contrast 'Quadratic conc' trmt +1-2 +1 0000000 ;
- contrast 'Linear l' trmt 000-10 +10000;
- contrast 'Quadratic I' trmt $000+1-2+10000$;
- contrast 'Linear P' trmt 000000-10+10;
- contrast 'Quadratic P' trmt $000000+1$-2 +10;
- means trmt/duncan;
- Ismeans trmt/stderr;
- File: Linear Quadratic response


## Linear, quadratic and cubic curves








Figure 19.1 General types of curves.

## Calculation of digestion rate of fibre or protein

Fractional digestion rate?
Example of a Tank filled with water
Model fitting
$\mathrm{Ct}=\mathrm{CO} . \mathrm{e}(-\mathrm{kt})$
Where
Ct = amount of potentially digestible fibre remaining at any time.
C0 = amount of substrate remaining at time zero
e = exponential
k $\quad=$ fractional digestion rate
t = time

To solve the above equation,
take natural logarithm (In) of both the sides.
The above equation then becomes like the following:
In Ct = $\mathbf{l n} \mathbf{C O} \mathbf{- k t}$

Lag time= (In 100-intercept)/rate of digestion.

Example: digestion rate calculation. excel
Non linear Model in SAS:
Example: nonlinear model for digestion rate.sas

## As a Nutritionist you should know

- What you want to do?
- You can draw the desired conclusions by changing a design
- Precision and accuracy
- Coefficient of variation
- Probability level
- Type I and Type II Error
- Standard Deviation vs Standard Error
- Sample size
- Treatments well apart to detect the difference
- Stat significance vs practical significance
- Interpretation of data: regression and correlation example
- Drawing conclusions


## HOPE YOU UNDERSTOOD IT



## Additional slides

## Type I Error: <br> Rejecting the null hypothesis when it is true

## Type II

Accepting the null hypothesis when it is false

## Precision and accuracy

## Precision

the magnitude of difference between two treatments that an experiment is capable of detecting at a given level of significance

Accuracy
The degree of closeness with which a measurement can be made

The measurement can be accurate but not precise
Examples: Watch, Balance, Any equipment that change its results with calibration

## Standard Deviation and Standard Error of mean

 Standard Deviation:Average Squared Deviation: Variance

$$
s^{2}=\frac{\Sigma\left(Y_{1}-Y\right)^{2}}{(n-1)}
$$

Root mean square Deviation:
Represented by small s for a sample and $\sigma$ for a population

Deviation from mean of a Sample/ population

## Standard Deviation of Mean or Standard Error



Standard Deviation applies to observation and Standard Error applies to means

## Co efficient of variation:

A quantity used for evaluating results from different experiments

$$
=\frac{100 s}{\bar{Y}} \text { percent }
$$

## Interpretation of Results

## Describing results

## Explaining results

## Regression

The magnitude of change in a dependant variable as a result of per unit change in an independent variable

## Or

Increase of decrease in a dependant variable as a result of per unit increase or decrease in an independent variable

## Example: FCR

## Correlation:

Measurement of relationship between two variables
Relationship could be positive or negative

Relationship between the number of storks flown over Tokyo city and number of births


Number of storks flown over Tokyo city

