

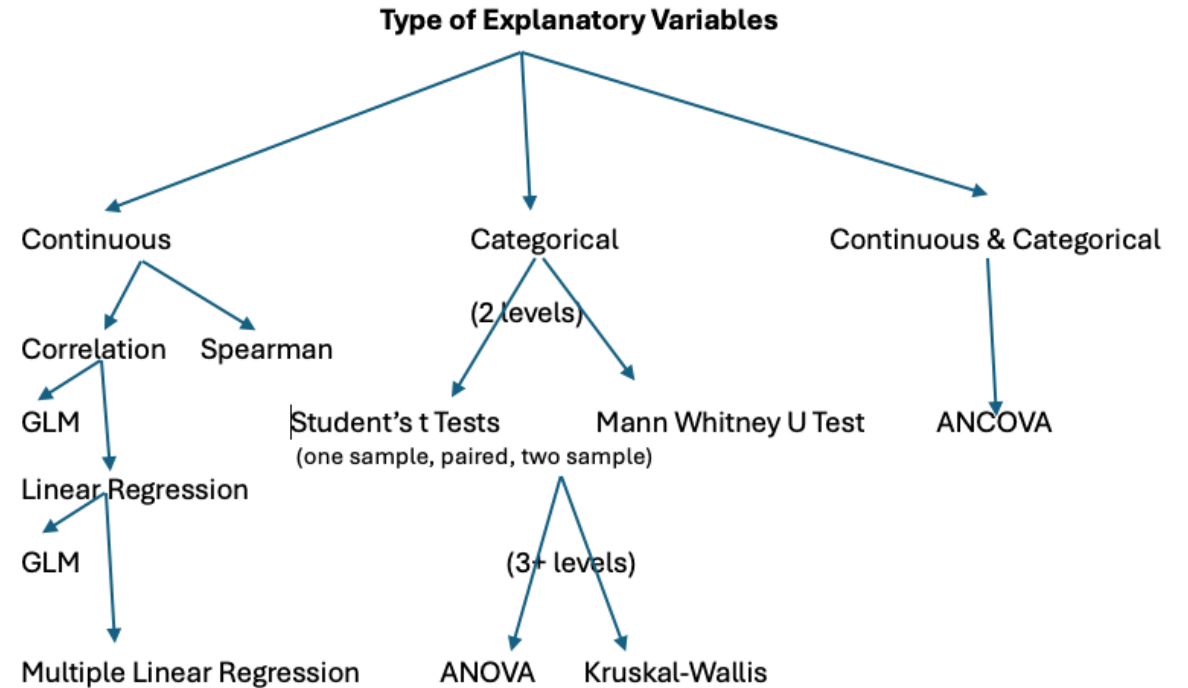
# Module 3C:

# **ANOVA & Correlation**

Assigning signal and noise to variation

# Agenda:

1. ANOVA: Nuts & Bolts
2. Worked Example
  - A. **One way ANOVA**
  - B. Post-hoc tests: Tukey-Kramer
  - C. Kruskal-Wallis (nonparametric)
3. Linear Correlation
  - A. Spearman's rank



## Kruskal-Wallis Test:

- o A non-parametric test similar to a single factor ANOVA
- o Uses the **ranks** of the data points; tests **medians** not means
  - Data points are **not** compared, their ranks are!
  - *Using **ranks** is what frees us from having to assume normality since all distributions have similar predictions about ranks*
  - All group samples are random samples
  - Distribution of the variable has the same shape in every population
  - Small samples lead to little power but when n is large, Kruskal-Wallis has the same power as ANOVA
- o **H**, sampling distribution is  $\chi^2$  with  $df = k - 1$

Researchers want to know whether **activity level** differs across three mouse strains (B6, BALB/c, CAST) after 12 weeks on the same diet. Activity is scored on a 1–10 scale.

Here are the activity scores:

**Strain A: B6 (n = 4)**

M1: 3.0  
M2: 3.5  
M3: 6.0  
M4: 6.5

**Strain B: BALB/c (n = 4)**

M5: 4.0  
M6: 4.5  
M7: 7.0  
M8: 7.5

**Strain C : CAST (n = 4)**

M9: 5.0  
M10: 5.5  
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Step 1: Formulate the null/alternate hypothesis

**H<sub>0</sub>:** The median activity level among the three mouse strains is equal (the independent samples all have the same central tendency and therefore come from the same underlying population)

**H<sub>a</sub>:** The median activity level among the three mouse strains is not equal (at least one of the independent samples does not have the same central tendency and therefore originates from a different population)

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3.0, 3.5, 4.0, 4.5, 5.0, 5.5, 6.0, 6.5, 7.0, 7.5, 8.0, 8.5

$$R_A = 1+2+7+8 = 18$$

$$R_B = 3+4+9+10 = 26$$

$$R_C = 5+6+11+12 = 34$$

$$H = \frac{12}{N(N+1)} \sum_{i=1}^k \frac{R_i^2}{n_i} - 3(N+1)$$

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$$H = \frac{N - 1}{N} \sum_{i=1}^k \frac{n_i(\bar{R} - E_R)^2}{\sigma^2}$$

$$= \frac{11}{12} \sum_{i=1}^k \frac{4(\frac{18}{4} - (12+1)/2)^2}{(12^2-1)/12} + \frac{4(\frac{26}{4} - (12+1)/2)^2}{(12^2-1)/12} + \frac{4(\frac{34}{4} - (12+1)/2)^2}{(12^2-1)/12} = 2.61$$

N=12  
k=3 groups  
 $\overline{R_A}=18/4 = 4.5$   
 $\overline{R_B} = \frac{26}{4} = 6.5$   
 $\overline{R_C} = \frac{34}{4} = 8.5$

$E_R = \frac{N+1}{2} = 13/2 = 6.5$   
 $\sigma^2 = \frac{N^2 - 1}{12} = 143/12$



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**H = 2.61**

Step 3: Critical value

There are **df=k-1=2** and you use a  **$\chi^2$**  table to find the cut-off value.

df	$\chi^2_{.995}$	$\chi^2_{.990}$	$\chi^2_{.975}$	$\chi^2_{.950}$	$\chi^2_{.900}$	$\chi^2_{.100}$	$\chi^2_{.050}$	$\chi^2_{.025}$	$\chi^2_{.010}$	$\chi^2_{.005}$
1	0.000	0.000	0.001	0.004	0.016	2.706	3.841	5.024	6.635	7.879
2	0.010	0.020	0.051	0.103	0.211	4.605	5.991	7.378	9.210	10.597
3	0.072	0.115	0.216	0.352	0.584	6.251	7.815	9.348	11.345	12.838
4	0.207	0.297	0.484	0.711	1.064	7.779	9.488	11.143	13.277	14.860
5	0.410	0.554	0.831	1.145	1.610	9.236	11.070	12.838	15.086	16.750

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Step 3: Critical value

There are **df=k-1=2** and you use a  $\chi^2$  table to find the cut-off value which is **5.991** for  $\alpha=0.05$ .

Step 4: Decision

Fail to reject the null hypothesis

## Fixed Effects: The groups *are* the question

- Also called Model 1 ANOVA
  - What we have been using so far
- Different categories of explanatory variable are predetermined and repeatable
  - **Results cannot be generalizable**
  - Example: specific drug treatment, specific diets, specific season

## Random Effects: The groups *are* a source of noise in the system you are modeling

- Also called Model 2 ANOVA
- Different categories of explanatory variable are *randomly sampled from a larger population of groups*
  - **Results are generalizable**; conclusions reached about difference among groups can be generalized to the whole population
- Example: family in a study about resemblance of IQ
  - Chose a random family in a population of families
  - Family: group
  - Replicates: different children within each family
- **The population and not the particular families involved is the target of study**

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  - What we have been using so far
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  - **Results cannot be generalizable**
  - Example: specific drug treatment, specific diets, specific season

*Do you want to estimate the effect of this specific group (fixed), or treat the group as one random example drawn from a broader population (random)?*

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Example: You compare **Chow vs. HFD vs. Low-Fat** diets.

- You care about *those* diets, not some random sample of all possible diets in the universe.
- You want to estimate:  $\mu_{\text{Chow}}$ ,  $\mu_{\text{LFD}}$ ,  $\mu_{\text{HFD}}$
- These diet means *are the whole point of the study* → **fixed effects**.

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  - **Results are generalizable**; conclusions reached about difference among groups can be generalized to the whole population
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Example: Your mice come from **several litters** or **multiple cages**, but you don't care about comparing litter #1 vs litter #2.

- You treat **litter** or **cage** as a random effect because:
- Litters are not scientifically interesting
- They create real variability
- You want to **control for** that variability without estimating each litter's mean

*Random effects represent “background noise” you want to account for because it changes the variance, not because you want to measure each group*

Example: Suppose you measure **PPARG expression** across three strains: B6, BALB/c, CAST

If your goal is:

**A. “How do PPARG levels differ between these specific strains?”**

**→ Strain = Fixed effect**

**B. “Strain is just a nuisance variable; these 3 strains are a random sample of wild genetic diversity.”**

**→ Strain = Random effect**

The same factor can be fixed or random depending on the scientific question.

## Quick heuristic:

**Ask: “Do I want to estimate the mean of each group?”**

Yes → Fixed effect

No → Random effect

**Ask: “Do these group levels represent all the possibilities or just a sample?”**

All (or all that matter) → Fixed effect

Just a sample of many possible levels → Random effect