

Statistical Inference for Omics — t-tests, ANOVA & GLM — Hands-on

Build a solid, practice oriented understanding of statistical inference for omics and biomedical data. This module walks you from basic sampling distributions through t tests, ANOVA and generalized linear models, with a focus on assumptions, effect sizes, confidence intervals and reproducible reporting in R and Python.

Statistical Inference for Omics — t-tests, ANOVA, GLM

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Session 1

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Foundations of Statistical Inference

Omics data types and distributions

[continuous / counts / proportions](#) [mean variance relationship](#) [log and variance stabilizing transforms](#)

Sampling distributions and central limit theorem

[standard error and uncertainty](#) [bootstrap intuition](#)
[finite sample caveats](#)

Confidence intervals and p values

interval estimation vs testing **one sided vs two sided**
practical vs statistical significance

Session 2

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Group Comparisons: t-tests & ANOVA

Two group and paired comparisons

independent and paired t tests **Welch correction** **non**
parametric alternatives

One way and multi factor ANOVA

F statistic and variance decomposition **interaction**
terms **balanced vs unbalanced designs**

Post hoc tests and assumptions

pairwise comparisons **normality and variance checks**
robust and rank based options

Session 3

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GLM for Omics and Count Data

Linear models for omics signals

design matrices and contrasts **covariate adjustment**
batch effects in models

Generalized linear models

logistic regression for binary traits **Poisson and**
negative binomial for counts **link functions and**
interpretation

Diagnostics and goodness of fit

residual plots **overdispersion checks** **influence and leverage**

Session 4

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Omics Inference Clinic & Reporting

Case studies with real omics datasets

gene expression and clinical cohorts

Reproducible analysis in R and Python

R stats and broom **Python statsmodels and pingouin**
scripted workflows and notebooks

Deliverables: analysis report and scripts

PDF or HTML summary **annotated R and Python code**
assumption and decision log