

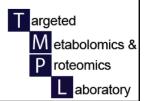
UAB Metabolomics Workshop July 17, 2017

Knowledge that will change your world

Designing a Metabolomics Experiment

Xiangqin Cui, PhD





Experimental Design

- Experimental design: is a term used about efficient methods for planning the collection of data, in order to obtain the maximum amount of information for the least amount of work. Anyone collecting and analyzing data, be it in the lab, the field or the production plant, can benefit from knowledge about experimental design. http://www.stat.sdu.dk/matstat/Design/index.html
- Good experimental design is the foundation for valid answer to research questions

Key Questions for Experimental Design

- What are the research questions to be answered?
- How will the data be analyzed?
- What is the best design of the experiment to answer the questions using the analysis methods?

General Statistical Principals of Experimental Design

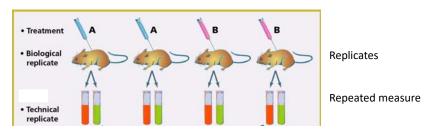
- Replication
- Randomization
- Blocking
- Use of factorial experiments instead of the one-factor-at-a-time methods.
- Orthogonality

Replication

 Replication is repeating the creation of a phenomenon (or redo your experiment), so that the <u>variability</u> associated with the phenomenon can be estimated. (no replication, no way to know the variability)

Replications should not be confused with repeated measurements which refer to taking several measurements of a single occurrence of a phenomenon (single experiment).

• Replications should not be confused with repeated measurements.



More terms saying the same things

- What to replicate?
 - Biological replicates (replicates at the experimental unit level, e.g. mouse, plant, pot of plants...)
 - Experimental unit is the unit that the experiment treatment or condition is directly applied to, e.g. a plant if hormone is sprayed to individual plants; a pot of seedlings if different fertilizers are applied to different pots.
 - Technical replicates
 - Any replicates below the experimental unit, e.g. different leaves from the same plant sprayed with one hormone level; different seedlings from the same pot; Different aliquots of the same RNA extraction; multiple arrays hybridized to the same RNA; multiple spots on the same array.

Randomization

• The experimental treatments are assigned to the experimental units (subjects) in a random fashion. It helps to eliminate effect of "lurking variables", uncontrolled factors which might vary over the length of the experiment. Randomization is essential for making causal inferences.

Example

 Number the objects to be randomized and then randomly draw the numbers.

Example: Assign two treatments, Special Diet and control, to 6 mice



Special Diet: 1, 3, 4 Control: 2, 5, 6

Blocking

- Some identified uninteresting but varying factors can be controlled through blocking.
 - COMPLETELY RANDOMIZED DESIGN
 - COMPLETE BLOCK DESIGN
 - INCOMPLETELY BLOCK DESIGNS

Completely Randomized Design

There is no blocking

- → Example
 - Compare two hormone treatments (trt and control) using 6 Arabidopsis plants (or mice or human).



Hormone trt: (1,3,4); (1,2,6) Control:

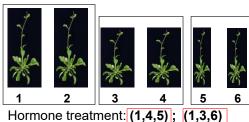
(2,5,6); (3,4,5)

Complete Block Design

➡ There is blocking and the block size is equal to the number of treatments.

Example:

• Compare two hormone treatments (trt and control) using 6 Arabidopsis plants. For some reason plant 1 and 2 are taller, plant 5 and 6 are thinner.



Control: (2,3,6); (2,4,5)

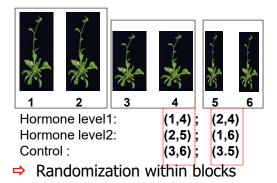
Randomization within blocks

Incomplete Block Design

→ There is blocking and the block size is smaller than the number of treatments.

Example:

 Compare three hormone treatments (hormone level 1, hormone level 2, and control) using 6 Arabidopsis plants. For some reason plant 1 and 2 are taller, plant 5 and 6 are thinner.



Applications of these principals in metabolomics studies

General Statistical Principals of Experimental Design

- Replication
- Randomization
- Blocking
- Use of factorial experiments instead of the onefactor-at-a-time methods.
- Orthogonality

Replication in Metabolomics Experiments

- Number of replication can go from a few to tens, but rarely hundreds.
- The larger the number of replicates the better, but budget is always limited.
- One sample per treatment/condition is not OK.
 What is wrong?

Sample Size and Statistical Power Calculation

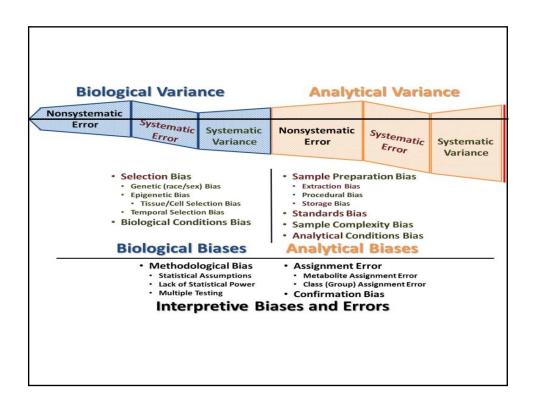
•Sample size for a general two group comparison

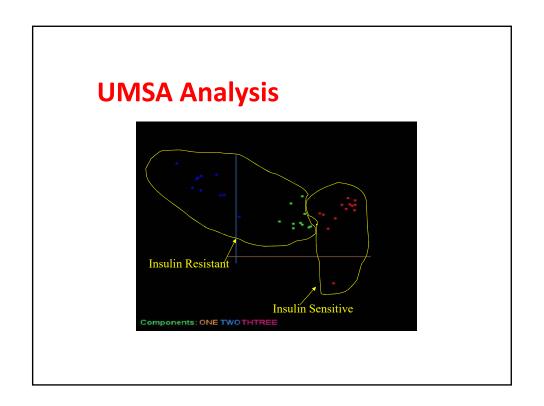
$$n = \frac{2(z_{(1-\alpha/2)} + z_{(1-\beta)})^{2}}{(\delta/\sigma)^{2}}$$

- n increases as variability, σ , increases.
- n increases as the difference between two means, δ , decreases.
- n increases as the significant level of the test, α , decreases.
- n increases as the power of the test, 1- β , increases.

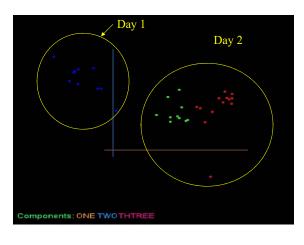
General Statistical Principals of Experimental Design

- Replication
- Randomization
- Blocking
- Use of factorial experiments instead of the onefactor-at-a-time methods.
- Orthogonality



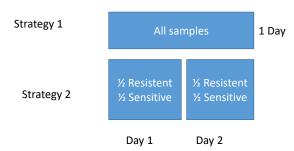


UMSA Analysis



How to Solve This Problem?

- Process all samples in the same day.
- Process half sensitive samples and half resistant samples in each day (balance sample groups against days—"treat each day as a block" in statistical terms).



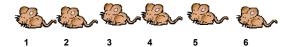
Known sources of non-biological biases (not exhaustive) that must be addressed

- Technician / post-doc
- Reagent lot
- Temperature
- Protocol
- Date
- Location
- Cage/ Field positions

Too Many factors to balance? -- Randomize

 Number the objects to be randomized and then randomly draw the numbers.

Example: Assign two treatments, Special Diet and control, to 6 mice



Special Diet: 1, 3, 4 Control: 2, 5, 6

Too Many factors to balance? -- Randomize

- Randomize samples in respect to treatments
- Randomize the order of handling samples.
- Randomize batches/runs/days in respect to samples
- Randomize over any other variable procedures.

References

Churchill GA. Fundamentals of experimental design for cDNA microarrays. Nature Genet. 32: 490-495, 2002.

Cui X and Churchill GA. How many mice and how many arrays? Replication in mouse cDNA microarray experiments, in "Methods of Microarray Data Analysis III", Edited by KF Johnson and SM Lin. Kluwer Academic Publishers, Norwell, MA. pp 139-154, 2003.

Gadbury GL, et al. Power and sample size estimation in high dimensional biology. Stat Meth Med Res 13: 325-338, 2004.

Kerr MK. Design considerations for efficient and effective microarray studies. Biometrics 59: 822-828, 2003.

 $\hbox{Kerr MK and Churchill GA. Statistical design and the analysis of gene expression microarray data. Genet. Res. 77: 123-128, 2001.$

Kuehl RO. Design of experiments: statistical principles of research design and analysis, 2^{nd} ed., 1994, (Brooks/cole) Duxbry Press, Pacific Grove, CA.

Page GP et al. The PowerAtlas: a power and sample size atlas for microarray experimental design and research. BMC Bioinformatics. 2006 Feb 22;7:84.

Rosa GJM, et al. Reassessing design and analysis of two-colour microarray experiments using mixed effects models. Comp. Funct. Genomics 6: 123-131. 2005.

Wit E, et al. Near-optimal designs for dual channel microarray studies. Appl. Statist. 54: 817-830, 2005.

Yand YH and Speed T. Design issues for cDNA microarray experiments. Nat. Rev. Genet. 3: 570-588, 2002

Thank you – Questions?