

Multivariate: MANOVA and repeated measures ANOVA

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1 Goals

1.1 Goals

1.1.1 Goals of this section

- **Multiple measures** of the same thing or related things as an **outcome**
 - Possibly over time
- Want the variables **separate**: Not PCA / FA

- In this section:
 - MANOVA (this week)
 - Repeated measures ANOVA (this week)
 - Mixed models (next week)
 - Mediation (2 weeks)

1.1.2 Goals of this lecture

- Multivariate Analysis of Variance (MANOVA)
 - Outcome is **multivariate**: Several outcome variables
- Repeated measures ANOVA (RM ANOVA)
 - **Univariate**: *Single* outcome variable, measured multiple times
 - **Multivariate**: *Multiple* outcome variables
- Punchline: MANOVA is almost never a good choice
 - But multivariate RM ANOVA is a decent approach

2 MANOVA

2.1 Univariate to multivariate

2.1.1 Extending ANOVA to multiple outcomes

- Frequently interested in more than 1 outcome at a time
 - Anxiety
 - * Test anxiety, minor stressor anxiety, general anxiety
 - Children's school achievement
 - * Reading ability, reasoning ability, math ability
 - Performance on a task
 - * Speed and accuracy

2.1.2 Could do GLM on each outcome but...

- ...you (often) shouldn't
 - **Inflated type I error** due to multiple tests on *correlated* outcomes
 - Sometimes only the **combination** of the outcomes shows an effect
 - Ignore **relations between DVs**

2.1.3 Structure of this section

- Review (univariate) between-subjects ANOVA
 - One outcome
- Extend to multivariate version
 - Multiple related outcomes

2.1.4 Univariate analysis of variance (ANOVA)

- Independent variables (IVs) are **categorical groups**
 - e.g., treatment and control
- Independent variables are called **factors**
 - Not to be confused with latent factors
- **Single** outcome variable (DV)
 - Continuous, normally distributed

2.1.5 ANOVA hypotheses are about the means

- One factor ANOVA
 - k levels of the independent variable
 - Null hypothesis: All k group means are equal
 - * $H_0 : \mu_1 = \mu_2 = \dots = \mu_k$

2.1.6 ANOVA hypotheses are about the means

- Two factor ANOVA
 - k levels of one IV, m levels of other IV
 - 3 null hypotheses
 - * Main effect 1: All k means across factor 1 are equal
 - * Main effect 2: All m means across factor 2 are equal
 - * Interaction: All cell means are equal

2.1.7 Partitioned variation

- Partition the variation in scores into:
 - between-subject portion (group differences, $SS_{between}$)
 - within-subject portion (error, SS_{within})
 - $SS_{total} = SS_{between} + SS_{within}$
- Calculate based on *observed scores, group means, grand mean*
 - X_{fi} = score for subject f in condition i
 - \bar{T}_i = mean for scores in condition i
 - \bar{G} = grand mean of all scores in the study

2.1.8 Partitioned variation

- Between group variation:

$$SS_{between} = n \sum (\bar{T}_i - \bar{G})^2 = n [(\bar{T}_1 - \bar{G})^2 + (\bar{T}_2 - \bar{G})^2 + \dots + (\bar{T}_k - \bar{G})^2]$$

- Within group variation:

$$SS_{within} = \sum (X_{fi} - \bar{T}_i)^2 = (X_{1i} - \bar{T}_i)^2 + (X_{2i} - \bar{T}_i)^2 + \dots + (X_{ni} - \bar{T}_i)^2$$

2.1.9 Testing the hypothesis

$$MS_{between} = \frac{SS_{between}}{k - 1}$$

$$MS_{within} = \frac{SS_{within}}{k(n - 1)}$$

$$F = \frac{MS_{between}}{MS_{within}}$$

- Compare observed F to critical $F(k - 1, k(n - 1))$
 - Significant test = at least one of the k groups is different from the other groups

2.2 MANOVA model

2.2.1 Multivariate analysis of variance (MANOVA)

- Independent variables are **categorical groups**
 - e.g., treatment and control
- Independent variables are called **factors**
 - Not to be confused with latent factors
- **Multiple** outcome variables
 - p outcome variables
 - Continuous, normally distributed

2.2.2 What does MANOVA do with all those outcomes?

- MANOVA creates a **linear combination** of the p outcome variables
 - Constructed to *separate* the k groups as much as possible
 - “Maximally discriminating linear combination”
- Look for group differences on the linear combination
- If you can’t find differences on the **maximally discriminating linear combination** of all the DVs, then there really really aren’t group differences on the DVs

2.2.3 MANOVA questions

- Do the groups differ at all?
 - On the maximally discriminating linear combination
- If yes, post hoc:
 - Which DVs have groups differences?
 - Which groups differ on those DVs?

2.2.4 Covariation matrix of outcomes **P**

- Covariation matrix of the p DVs: $p \times p$ matrix
 - Multivariate extension of SS_{total}
- Just like ANOVA: Partitions into **between** (**H**) and **within** (**E**)

$$\mathbf{P} = \begin{bmatrix} SS_1 & SP_{12} & \cdots & SP_{1p} \\ SP_{21} & SS_2 & \cdots & SP_{2p} \\ \vdots & \vdots & \ddots & \vdots \\ SP_{p1} & SP_{p2} & \cdots & SS_p \end{bmatrix}$$

2.2.5 Hypothesis matrix **H**

- Multivariate extension of $SS_{between}$: $p \times p$ matrix
 - Diagonal: between-group **variation** of each DV
 - Off-diagonal: **covariation** between means for pairs of DVs

$$\mathbf{H} = \begin{bmatrix} SS_{H,1} & SP_{H,12} & \cdots & SP_{H,1p} \\ SP_{H,21} & SS_{H,2} & \cdots & SP_{H,2p} \\ \vdots & \vdots & \ddots & \vdots \\ SP_{H,p1} & SP_{H,p2} & \cdots & SS_{H,p} \end{bmatrix}$$

2.2.6 Aside: **H** matrix for two-factor MANOVA

- For a one-factor MANOVA, there is a single **H** matrix
- For a two-factor MANOVA, there is a single **H** matrix
 - BUT it can be further partitioned into 3 matrices reflecting:
 - * Main effect 1
 - * Main effect 2
 - * Interaction effect

2.2.7 Error matrix **E**

- Multivariate extension of SS_{within} : $p \times p$ matrix
 - Diagonal: within-group **variation** of each DV, added across k grp
 - Off-diagonal: error **covariation**, added across k groups

- No between-group information in this matrix

$$\mathbf{E} = \begin{bmatrix} SS_{E,1} & SP_{E,12} & \cdots & SP_{E,1p} \\ SP_{E,21} & SS_{E,2} & \cdots & SP_{E,2p} \\ \vdots & \vdots & \ddots & \vdots \\ SP_{E,p1} & SP_{E,p2} & \cdots & SS_{E,p} \end{bmatrix}$$

2.2.8 Partitioned variation

- ANOVA
 - $SS_{total} = SS_{between} + SS_{within}$
- MANOVA
 - Total variation = between-group variation + within-group variation
 - One factor: $\mathbf{P} = \mathbf{H} + \mathbf{E}$
 - Two factor: $\mathbf{P} = \mathbf{H}_{factor1} + \mathbf{H}_{factor2} + \mathbf{H}_{factor1*factor2} + \mathbf{E}$

2.2.9 Multivariate hypothesis tests (omnibus)

- ANOVA
 - Divide SS by their degrees of freedom to produce MS (variances)
 - F -statistic is ratio of MS s (variances)
- MANOVA
 - Use matrix equivalent of variance: **Determinant**
 - * Determinant is “generalized variance” for a matrix
 - Create analogues to F -statistics
 - Unfortunately, it’s not straight-forward

2.2.10 Multivariate hypothesis tests

- Four commonly used multivariate tests
 - Different ratio of determinants or eigenvalues
- Wilks’ lambda: within / total
- Pillai’s trace: between / total
- Hotelling’s trace: between / within
- Roy’s largest characteristic root: between / total

2.2.11 Wilks' lambda

- $\Lambda = \frac{|\mathbf{E}|}{|\mathbf{H} + \mathbf{E}|} = \frac{|\mathbf{E}|}{|\mathbf{P}|}$
 - where $|\mathbf{E}|$ is the determinant of \mathbf{E}
- H_0 : no between-group variation, so \mathbf{H} is all zeroes and ratio is 1
 - As group differences increase, $\Lambda \rightarrow 0$
- Effect size = eta squared = $\eta^2 = 1 - \Lambda$
 - η^2 = variance accounted for by the best linear combination of DVs

2.2.12 Pillai's trace

- Pillai's trace = $\text{trace} [\mathbf{H}(\mathbf{H} + \mathbf{E})^{-1}]$
 - where the **trace** of a matrix is the **sum of the diagonal elements**
- Conceptually:
 - Matrix representing **proportion of variation that is between-group**
 - Sum of *eigenvalues* from that matrix

2.2.13 Hotelling's trace

- Hotelling's trace = $\text{trace} [\mathbf{H}(\mathbf{E})^{-1}]$
 - where the **trace** of a matrix is the **sum of the diagonal elements**
- Conceptually:
 - Matrix representing **ratio of between- to within-group variation**
 - Sum of *eigenvalues* from that matrix

2.2.14 Roy's largest characteristic root

- Roy's greatest characteristic root = first eigenvalue of $\mathbf{H}(\mathbf{H} + \mathbf{E})^{-1}$
- Conceptually:
 - Matrix representing **proportion of variation that is between-group**
 - *First eigenvalue* from that matrix

2.2.15 Summary of multivariate tests

Test	Matrix	Range (H_0 to H_A)	In words	Function
Wilks	E/T	1 to 0	Error proportion	Determinant
Pillai	H/T	0 to 1	Between proportion	Trace
Hotelling	H/E	0 to ∞	Between to within ratio	Trace
Roy	H/T	0 to 1	Between proportion	1st eigenvalue

These tests are similar, but they differ in terms of **power** and **robustness to violations** of assumptions

2.2.16 Assumptions of MANOVA

- GLM: Multivariate normality of outcomes, linearity, etc
- “Homogeneity of variance-covariance matrices”
 - Error matrix is same in all groups and \mathbf{E} is average
 - Multivariate extension of homogeneity of variance assumption
- Box’s M test to test this assumption
 - Significant test means that assumption is violated
 - Sensitive: use $p < .001$, ignore unless n s very different across groups

2.2.17 Which test should I use???

- One factor MANOVA with $k = 2$ groups: All tests are identical
- **Recommended:** Pillai’s trace
 - Robust to assumptions, powerful when DVs not highly corr
- **Recommended:** Wilks’ lambda
 - Good power, relatively robust when assumptions probably met
- **Maybe use:** Roy’s greatest characteristic root
 - Powerful when DVs highly corr, not robust to assumptions
- **Not recommended:** Hotelling’s trace
 - OK when sample size is very large

2.3 Summary and alternatives

2.3.1 MANOVA

- Extends ANOVA to multiple outcomes
 - Many omnibus test options
 - Many follow-up options
 - Maximally discriminating linear combination?
 - Missing data, ANOVA framework only, time
- Quantitude says [MANOVA must die](#)

2.3.2 MANOVA questions

- Do the groups differ at all (on max discriminating linear comb.)?
 - This is what Pillai's trace, etc are testing
- **If yes, post hoc:**
 - Which DVs have groups differences?
 - Which groups differ on those DVs?
 - Enders, C. K. (2003). Performing multivariate group comparisons following a statistically significant MANOVA. *Measurement and Evaluation in Counseling and Development*, 36, 40-56.

2.3.3 When to use MANOVA?

- DVs are **highly negatively correlated**
 - Time to complete a task and number of errors on task
- DVs are **all moderately correlated** in either direction
 - Around ± 0.6 correlation
 - Not really high enough to support a latent factor
 - Repeated measures

2.3.4 When not to use MANOVA?

- DVs are **not really correlated**
 - MANOVA is unnecessarily complicated and wasteful
 - You don't gain anything by analyzing them together
- DVs are all **highly positively correlated**
 - MANOVA is unnecessarily complicated and wasteful
 - The variables are all basically the same thing

2.3.5 Alternatives to MANOVA

- Repeated-measures DVs:
 - **Repeated measures ANOVA**
 - **Mixed / multilevel / hierarchical linear models**
 - Latent growth models
- Separate univariate ANOVAs: esp *uncorrelated* DVs
- SEM / path model with multiple DVs
- Latent factor: esp highly *correlated* DVs

3 Repeated measures ANOVA

3.1 Overview / review

3.1.1 Between-subjects ANOVA

- Different subjects in each condition or cell of the design
 - 2 dimensions: subjects and variables

subject	condition	outcome
1	1	3
2	1	4
3	1	3
4	2	5
5	2	3
6	2	3
7	3	1

subject	condition	outcome
8	3	2
9	3	4

3.1.2 Between-subjects ANOVA: Partitioning

- Partition the variation in scores into:
 - between-subject portion (group differences, $SS_{between}$)
 - within-subject portion (error, SS_{within})
 - $SS_{total} = SS_{between} + SS_{within}$

3.1.3 Repeated-measures

1. Measure the **same DV** over **time**
 - e.g., anxiety level at 1 wk intervals after starting medication
2. Measure the **same DV** in each of a **set of related conditions**
 - e.g., anxiety level after CBT, after medication, etc.
- Multiple outcome measures that are **related**
 - Measured on the same person (not independent)
 - MANOVA: **related** dependent variables

3.1.4 Repeated-measures ANOVA

- Subjects are repeatedly measured / same subject in all conditions
 - 3 dimensions: subjects, variables (Y_1), treatment or time (T)

subject	$Y1_T1$	$Y1_T2$	$Y1_T3$	$Y1_T4$
1	3	1	2	5
2	4	5	1	3
3	3	3	3	3
4	5	2	4	2
5	3	4	4	5
6	3	3	4	4
7	1	1	4	5
8	2	5	2	1
9	4	4	5	2

3.1.5 Two ways to do repeated-measures ANOVA

- Univariate:
 - Standard repeated measures ANOVA
 - Treats the outcome as **one variable** that is **measured repeatedly**
- Multivariate:
 - Treats the outcome as a **multivariate outcome**
 - * Single outcome made up of **several (related) variables**
 - Sound familiar?

3.1.6 Univariate: n subjects, k repeated measures

- Single outcome variable Y
 - “Univariate”
- T (time or treatment) is a predictor
 - Specific levels: $1, 2, \dots, k$
- Also called “tall” or “stacked” data format
 - Used in mixed models (next week)

3.1.7 Univariate: n subjects, k repeated measures

subject	T	Y
1	1	Y_{11}
1	2	Y_{12}
1	\vdots	\vdots
1	k	Y_{1k}
2	1	Y_{21}
2	2	Y_{22}
2	\vdots	\vdots
2	k	Y_{2k}
\vdots	3	\vdots
n	1	Y_{n1}
n	2	Y_{n2}
n	\vdots	\vdots
n	k	Y_{nk}

3.1.8 Multivariate: n subjects, k repeated measures

- Several related outcome variables Y
 - “Multivariate”
- T (time or treatment) is not an explicit predictor
 - Treated like waves
- Also called “wide” data format
 - Used in MANOVA

3.1.9 Multivariate: n subjects, k repeated measures

subject	$Y1_T1$	$Y1_T2$...	$Y1_T4$
1	Y_{11}	Y_{12}	...	Y_{1k}
2	Y_{21}	Y_{22}	...	Y_{2k}
3	Y_{31}	Y_{32}	...	Y_{3k}
\vdots	\vdots	\vdots	\ddots	\vdots
n	Y_{n1}	Y_{n2}	...	Y_{nk}

3.2 Univariate RM ANOVA

3.2.1 Univariate approach to repeated measures

- Partition variation in scores into:
 - Between-**subject** variation
 - Within-subject variation, which is further partitioned into:
 - * Treatment (or time) effects for individuals
 - * Residual or random error

3.2.2 Univariate approach to repeated measures

- Y_{ij} = score for person i at time or treatment j
- \bar{T}_j = mean score for treatment or time j
 - Up to k treatments or times
- \bar{P}_i = mean score for person i

- Up to n subjects
- \bar{G} = grand mean of all scores

3.2.3 Between-subjects variation

- Individual subjects' variation around the **grand mean**

$$SS_{between\ subject} = k \sum_{i=1}^n (\bar{P}_i - \bar{G})^2$$

$$= k[(\bar{P}_1 - \bar{G})^2 + (\bar{P}_2 - \bar{G})^2 + \dots + (\bar{P}_k - \bar{G})^2]$$

- Similar to between-**groups** variation in ANOVA, but no groups here
 - People are “groups”

3.2.4 Within-subjects variation

- Individual subjects' variation around **their mean**
- For person i :

$$SS_{within\ person\ i} = \sum_{j=1}^k (Y_{ij} - \bar{P}_i)^2$$

$$(Y_{i1} - \bar{P}_i)^2 + (Y_{i2} - \bar{P}_i)^2 + \dots + (Y_{ik} - \bar{P}_i)^2$$

- Add up across all n subjects: $SS_{within\ subject} = \sum_{i=1}^n \sum_{j=1}^k (Y_{ij} - \bar{P}_i)^2$

3.2.5 Within-subjects variation

- Within-subjects variation = **time (or treatment) + residual**
- Time variation = timepoint mean variation around grand mean

$$SS_{time} = n \sum_{j=1}^k (\bar{T}_j - \bar{G})^2$$

$$= n[(\bar{T}_1 - \bar{G})^2 + (\bar{T}_2 - \bar{G})^2 + \dots + (\bar{T}_k - \bar{G})^2]$$

3.2.6 Within-subjects variation

- Within-subjects variation = **time (or treatment) + residual**
- Residual variation = any remaining variation

$$SS_{residual} = SS_{time \times subject} = SS_{within\ subject} - SS_{time}$$

3.2.7 Full partitioning of variation

- Keep in mind: No groups here at all

$$SS_{total} = SS_{between\ subject} + SS_{time} + SS_{residual}$$

Source	SS	df	MS	F
Between	$SS_{between\ subject}$	$n - 1$	$MS_{between\ subject}$	
Within	$SS_{within\ subject}$	$n(k - 1)$	$MS_{within\ subject}$	
–Time	SS_{time}	$k - 1$	MS_{time}	$\frac{MS_{time}}{MS_{residual}}$
–Residual	$SS_{residual}$	$(n - 1)(k - 1)$	$MS_{residual}$	

3.2.8 Mixed effects ANOVA

- Between-subjects + within-subjects = “mixed ANOVA”
 - Unfortunate: too easy to confuse with “mixed models”
 - * Also have several other names: Next week
- You can have BOTH within and between subjects factors in ANOVA
 - e.g., group (between) and time (within)
- Also look at the interaction
 - Does time effect vary across groups? Or vice versa?

3.2.9 Assumptions of univariate RM ANOVA

- About the covariance matrix of the outcomes

$$\mathbf{S}_{YY} = \begin{bmatrix} \sigma_1^2 & \sigma_{12} & \sigma_{13} & \sigma_{14} \\ & \sigma_2^2 & \sigma_{23} & \sigma_{24} \\ & & \sigma_3^2 & \sigma_{34} \\ & & & \sigma_4^2 \end{bmatrix}$$

- σ_1^2 = variance of outcome at time 1 / treatment 1
- σ_{12} = covariance between outcome at time /treatment 1 and outcome at time / treatment 2

3.2.10 Compound symmetry and sphericity

- **Compound symmetry** of the covariance matrix of outcomes
 - Homogeneity of variances (i.e., variances are all the same):
 - * $\sigma_1^2 = \sigma_2^2 = \sigma_3^2 = \sigma_4^2$
 - Homogeneity of covariances (i.e., covariances are all the same):
 - * $\sigma_{12} = \sigma_{13} = \sigma_{14} = \sigma_{23} = \sigma_{24} = \sigma_{34}$
- Actual assumption: **Sphericity**
 - Compound symmetry holds for **differences** between pairs of scores
 - Slightly weaker assumption

3.2.11 Plausibility of sphericity assumption

- T_1 through T_k are different trials or conditions in a *single session*
 - Sphericity may be plausible
- T_1 through T_k are *different time points*
 - Say, 9th, 10th, 11th, and 12th grades
 - Probably expect T1 and T2 to be more alike than T1 and T4
 - Sphericity is probably not very plausible

3.2.12 Violations of Assumptions

- Even if sphericity is plausible, it still may be **violated**
 - Very small violations can **greatly increase type I error rate**
- How to deal with violation of sphericity?
 - Adjust for violations of sphericity to return alpha to .05
 - Use **multivariate test** of repeated measures (next section)

3.2.13 Adjusting for sphericity violations

- **Lower bound correction:** Most conservative
 - Ignore repeated measures, treat as between subjects
 - Use critical $F(1, n - 1)$
- **Greenhouse-Geisser:** Middle of the road
 - $\hat{\epsilon}$ ranges from $\frac{1}{k-1}$ (severe violation) to 1 (sphericity)
 - Multiply degrees of freedom by $\hat{\epsilon}$
- **Huynh-Feldt:** Least conservative, smallest adjustment
 - Multiply degrees of freedom by $\tilde{\epsilon}$ (function of $\hat{\epsilon}$)

3.3 Multivariate RM ANOVA

3.3.1 Multivariate approach to repeated measures

- Multivariate extension of *paired t-test*
- Basically a MANOVA on specific set of difference scores
 - Multivariate tests (i.e., Wilks' lambda) as in MANOVA

3.3.2 Vector of differences

- $k - 1$ differences between combinations of k repeated scores
 - Must be **linearly independent**
 - Most common: Differences between adjacent pairs of means
- For a single subject i :

$$\underline{Y}'_{id} = \begin{bmatrix} d_{i1} \\ d_{i2} \\ d_{i3} \\ \vdots \\ d_{i,k-1} \end{bmatrix} = \begin{bmatrix} Y_{i1} - Y_{i2} \\ Y_{i2} - Y_{i3} \\ Y_{i3} - Y_{i4} \\ \vdots \\ Y_{i,k-1} - Y_{ik} \end{bmatrix}$$

3.3.3 Matrix of difference scores

- $n \times (k - 1)$ matrix of difference scores is matrix of outcomes
 - Rows are subjects, columns are difference scores
 - k repeated measures: $k - 1$ difference scores

$$\mathbf{Y}_d = \begin{bmatrix} d_{11} & d_{12} & \dots & d_{1,k-1} \\ d_{21} & d_{22} & \dots & d_{2,k-1} \\ \vdots & \vdots & \ddots & \vdots \\ d_{n1} & d_{n2} & \dots & d_{n,k-1} \end{bmatrix}$$

3.3.4 Covariance matrix of differences

- $(k - 1) \times (k - 1)$ covariance matrix of differences
 - $s_{d_1}^2$ is the variance of the $(T1 - T2)$ scores across n subjects
 - $s_{d_1 d_2}$ is the covariance between $(T1 - T2)$ and $(T2 - T3)$
 - Unlike univariate test, **no assumptions about this matrix**
- For 4 timepoints, this is a 3×3 matrix:

$$\mathbf{S}_d = \begin{bmatrix} s_{d_1}^2 & s_{d_1 d_2} & s_{d_1 d_3} \\ & s_{d_2}^2 & s_{d_2 d_3} \\ & & s_{d_3}^2 \end{bmatrix}$$

3.3.5 Null hypothesis

- H_0 : $k - 1$ vectors of **mean differences** are simultaneously
 - All equal to each other AND all equal to 0
- NS test = no differences over time
 - All mean differences are 0
 - No adjacent differences are different from one another
- Significant test = differences over time
 - Some of the mean differences are not 0
 - Some adjacent differences are different from one another

3.3.6 Multivariate hypothesis tests

- Perform a MANOVA on the difference score matrix
 - Multivariate hypothesis tests:
 - * Wilks' lambda
 - * Pillai's trace
 - * Hotelling's trace
 - * Roy's largest characteristic root

3.4 Summary and comparison

3.4.1 Summary

- Univariate RM ANOVA
 - Single, repeatedly measured outcome
 - Sphericity assumption
- Multivariate RM ANOVA
 - Multiple, related outcomes
 - No sphericity assumption

3.4.2 Comparison

- Univariate approach: $F(k - 1, (n - 1)(k - 1))$
 - Assumptions about covariance matrix (sphericity)
 - * But can adjust if assumptions not met
 - Missing data results in loss of entire subject
- Multivariate approach: $F(k - 1, n - k + 1)$
 - No assumptions about structure of covariance matrix
 - * (except that $n \geq k$ so it is invertable)
 - Missing data results in loss of entire subject

3.4.3 Recommendations: univariate vs. multivariate

- Univariate is preferred with small sample sizes
 - Sphericity holds (rare): More powerful, simpler, correct α
 - ALWAYS use univariate (with correction) if $n < k$
- Multivariate is preferred with large sample sizes
 - If sphericity doesn't hold (common): correct α
 - Do not use unless $n \geq k$
 - * With BS factors: n in each BS group needs to be $\geq k$

4 Summary

4.1 Summary

4.1.1 Summary of this week

- MANOVA is a way to analyze multiple outcomes in one model
 - Almost never a good choice
 - Limited utility for repeated measures
- RM ANOVA has univariate and multivariate versions
 - Univariate has some easily violated assumptions
 - Multivariate is good but still limited
 - Missing data, ANOVA framework only, time

4.1.2 Next few weeks

- RM ANOVA (both) have shortcomings
 - Best for short-term or single-session studies
 - Does not capture the TIME aspect of longitudinal data
 - Requires same # of repeated measures for each subject
 - Not informative about individual growth
 - Focus on **average** differences over time and group differences
 - ANOVA framework, so only categorical predictors
- Mixed models, latent growth models solve many of these issues