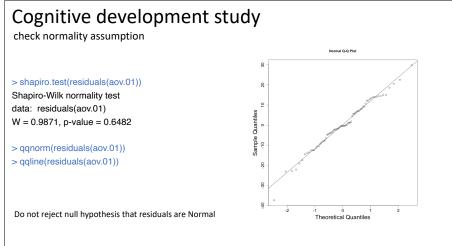
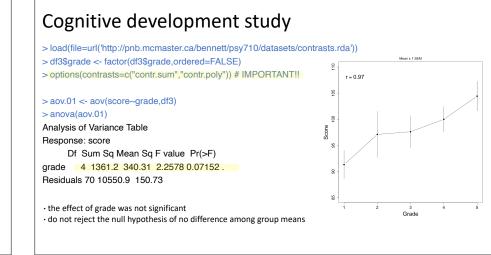
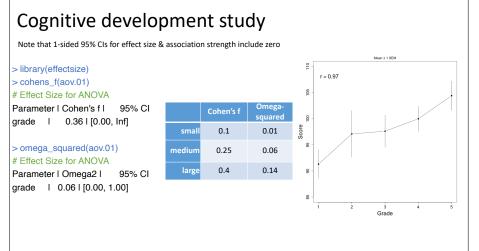
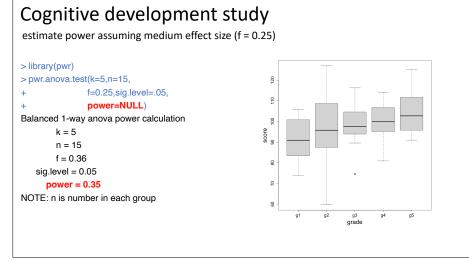


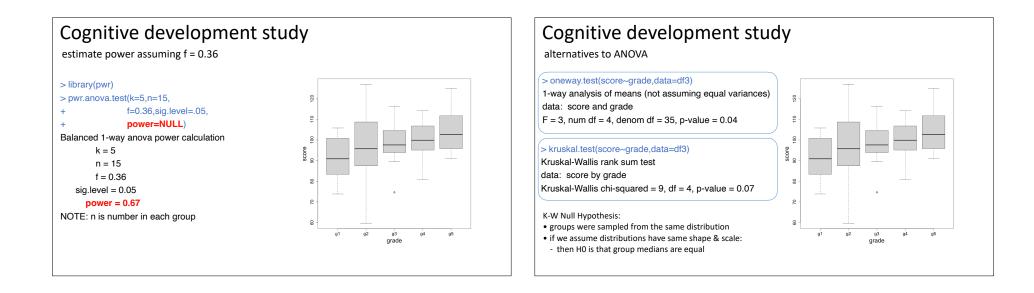
# Cognitive development study check constant variance assumption > bartlett.test(score--grade,df3) Bartlett test of homogeneity of variances data: score by grade Bartlett's K-squared = 6.6227, df = 4, p-value = 0.1572 Do not reject null hypothesis that variances are equal







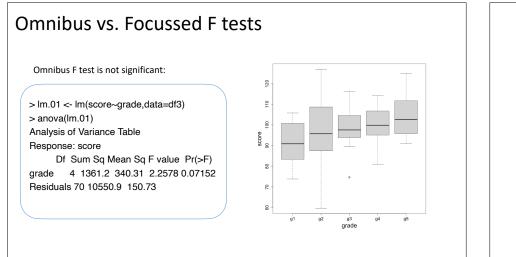






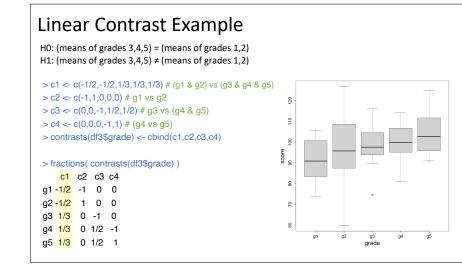
## Omnibus vs. Focussed F tests

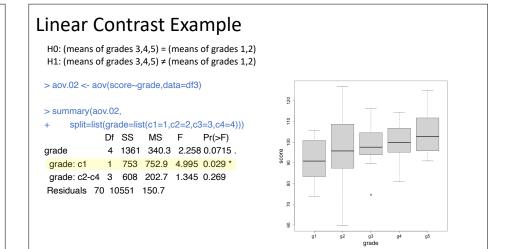
- $H0: \quad \alpha_1 = \alpha_2 = \dots = \alpha_a = 0$  $H1: \qquad \qquad \alpha_j \neq 0$
- A significant omnibus F test tests a very general hypothesis
- H0: all group means are equal; H1: not all means are equal
- H0: all group effects are zero; H1: not all group effects are zero
- Significant F doesn't tell us how group means differ
- Generality of omnibus F often comes at cost of reduced power

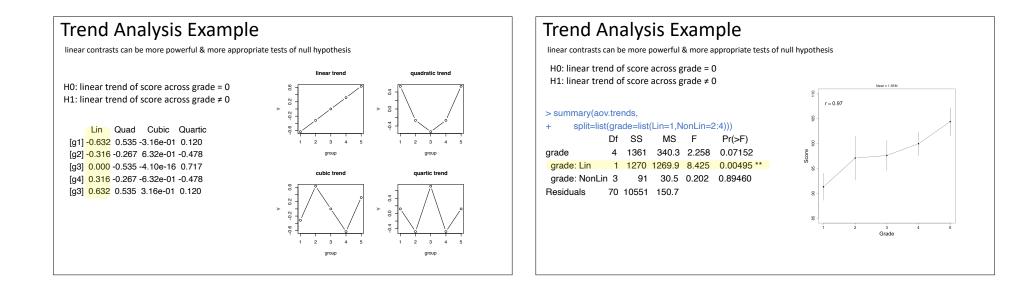


# focussed tests provide more power

linear contrasts often more appropriate & more powerful





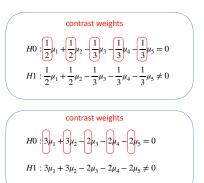


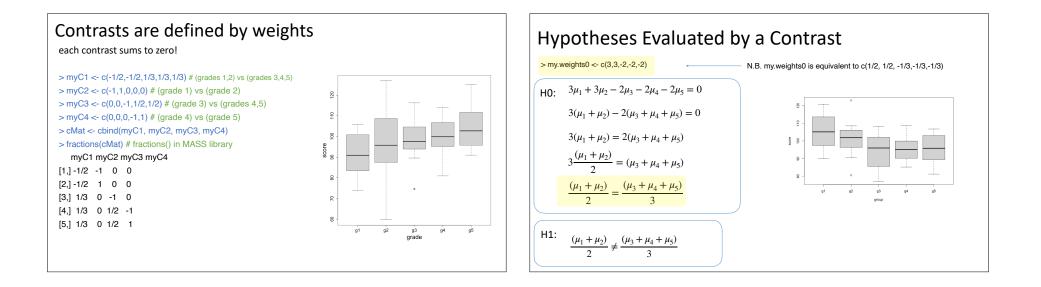
# Linear Contrasts (Comparisons)

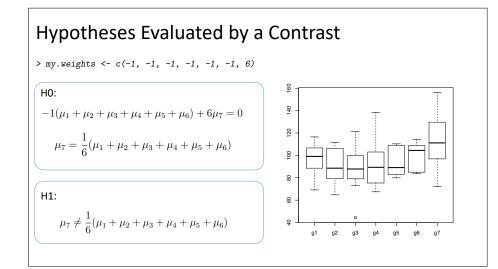
- · Contrasts allow us to evaluate focussed hypotheses
- evaluate specific pattern of differences among group means
- Each contrast is defined by a set of contrast weights
- weights (c1, c2, ... ca) specify a pattern of group means
- value of contrast,  $\boldsymbol{\psi},$  is a weighted combination of group means
- $\bullet \quad \psi = c_1 \overline{Y}_1 + c_2 \overline{Y}_2 + c_3 \overline{Y}_3 + c_4 \overline{Y}_4 + \dots c_a \overline{Y}_a$

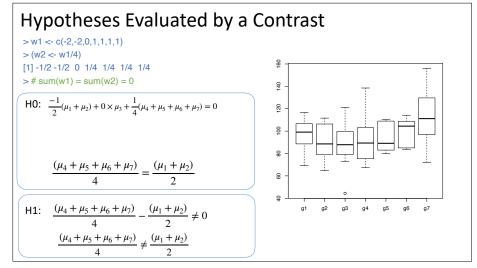
# Hypotheses tested with Linear Contrasts

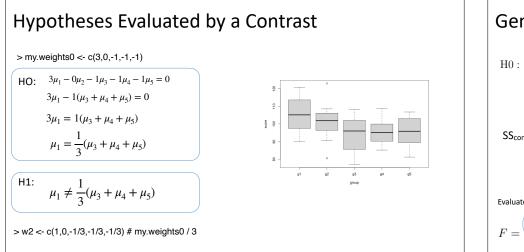
- Linear contrasts are defined by weights
- must sum to zero
- sum(1/2, 1/2, -1/3, -1/3, -1/3) = 0
- Multiplying weights by constant produces an equivalent linear contrast
- $w_1 = (1/2, 1/2, -1/3, -1/3, -1/3)$
- w<sub>2</sub> = 6 x w<sub>1</sub> = (3,3,-2,-2,-2)
- w<sub>1</sub> is equivalent to w<sub>2</sub>

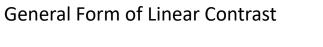


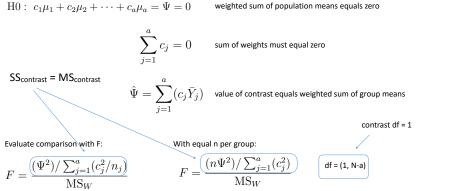


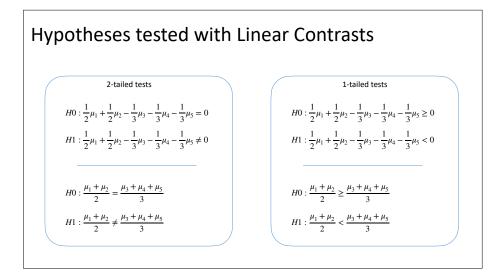


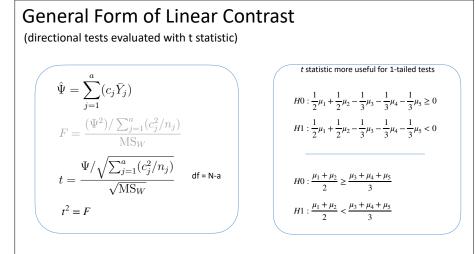


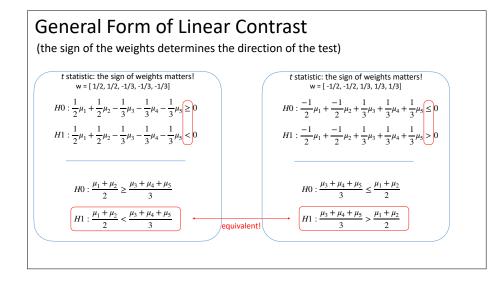


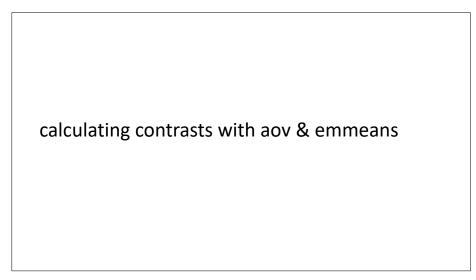












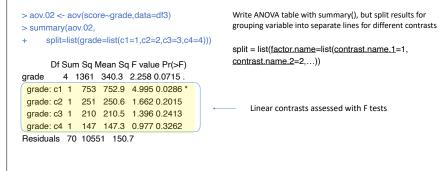
# Conducting Contrasts with R aov()

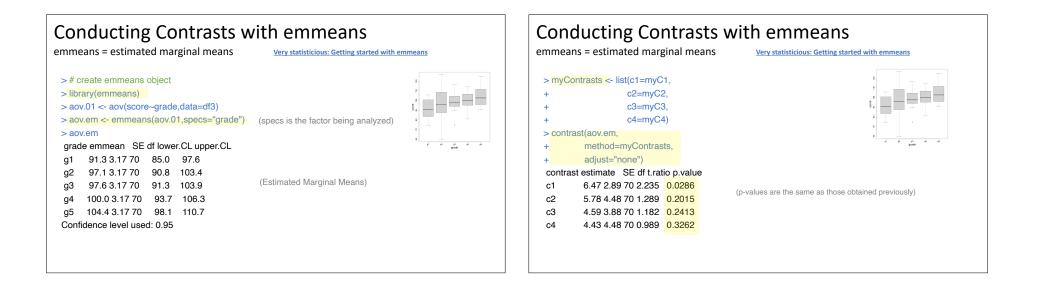
> contrasts(df3\$grade) <- cMat > fractions( contrasts(df3\$grade) ) myC1 myC2 myC3 myC4 g1 -1/2 -1 0 0 g2 -1/2 1 0 0 g3 1/3 0 -1 0 g4 1/3 0 1/2 -1 g5 1/3 0 1/2 1 > aov.02 <- aov(score~grade,data=df3)</pre>

Store contrast weights as columns in a matrix & then assign contrast weights to grouping variable

Perform ANOVA with aov

# Conducting Contrasts with R aov()





Conducting Contrasts with linear.comparison		
<pre>&gt; source(url("http://pnb.mcmaster.ca/bennett/psy710/Rscripts/linear_contrast_v2.R")) [1] "loading function linear.comparison" &gt; y &lt;&gt; df3\$score &gt; g &lt;&gt; df3\$grade &gt; myContrast1 &lt;&gt; linear.comparison(y,g,c.weights = myContrasts,var.equal=T)) [1] "computing linear comparisons assuming equal variances among groups" [1] "C 1: [F=4.995, t=2.235, p=0.029, psi=6.467, Cl=(0.367,12.568), adj.Cl= (-0.952,13.887)" [1] "C 2: F=1.662, t=1.289, p=0.202, psi=5.780, Cl=(-4.615,16.175), adj.Cl= (-5.714,17.274)" [1] "C 3: F=1.396, t=1.182, p=0.241, psi=4.588, Cl=(-2.544,11.719), adj.Cl= (-5.366,14.542)" [1] "C 4: F=0.977, t=0.989, p=0.326, psi=4.432, Cl=(-2.955,11.819), adj.Cl= (-7.062,15.926)"</pre>	trend analysis	

### **Trend Analysis**

trends are linear contrasts

- the analysis of trends uses the same methods as linear contrasts
- weights are designed to evaluate specific differences across groups:
- linear, quadratic, cubic, etc.
- weights must sum to zero
- weights can be calculated using R's contr.poly function
- useful when differences between levels on group variable are not constant

### > contr.poly(n=5,scores=c(8,9,10,11,12))

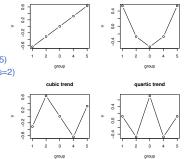
.L .Q. .C ^4 [1,] -0.6324555 0.5345225 -3.162278e-01 0.1195229 [2,] -0.3162278 -0.2672612 6.324555e-01 -0.4780914 [3,] 0.000000 -0.5345225 -4.095972e-16 0.7171372 [4,] 0.3162278 -0.2672612 -6.324555e-01 -0.4780914 [5,] 0.6324555 0.5345225 3.162278e-01 0.1195229

#### > contr.poly(n=5,scores=c(8,9,10,12,15)) .L .Q .C ^4

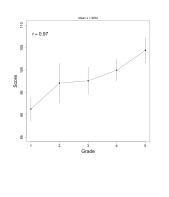
[1,] -0.5045250 0.54194676 -0.4466312 0.22862383 [2,] -0.3243375 -0.01290349 0.4344281 -0.71127414 [3,] -0.1441500 -0.38710483 0.4685966 0.64014672 [4,] 0.2162250 -0.59356074 -0.6077113 -0.17781853 [5,] 0.7567875 0.45162230 0.1513177 0.02032212

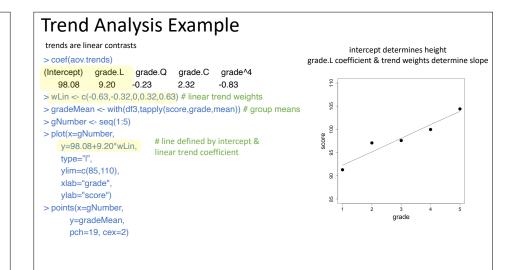
#### **Trend Analysis Example** trends are linear contrasts > # set polynomial contrasts as default for ordered factors: > options(contrasts=c("contr.sum","contr.poly") > load(file=url('http://pnb.mcmaster.ca/bennett/psy710/labs/L3/hw3-2021.rda')) > sapply(df3,class) linear trend quadratic trend \$grade [1] "ordered" "factor" \$score Trend weights are orthogonal [1] "numeric" > polyWeights <- contr.poly(n=5) > contrasts(df3\$grade) > round(cor(polyWeights).digits=2) .L .Q .C ^4

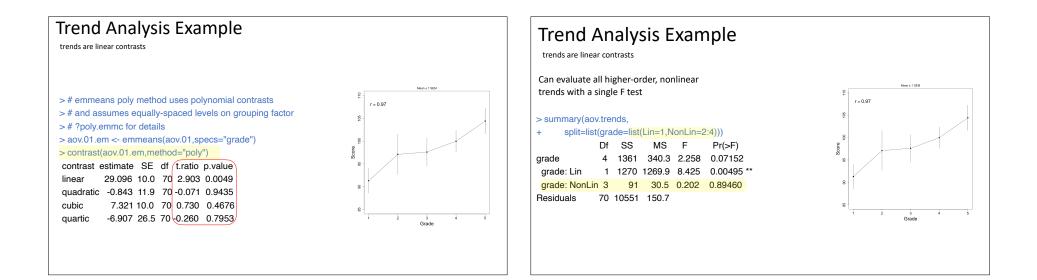
.L .Q .C ^4 g1 -0.63 0.53 -0.32 0.12 L 1.00 0.00 0.00 0.00 g2 -0.32 -0.27 0.63 -0.48 .Q 0.00 1.00 0.00 0.00 g3 0.00 -0.53 0.00 0.72 .C 0.00 0.00 1.00 0.00 g4 0.32 -0.27 -0.63 -0.48 ^4 0.00 0.00 0.00 1.00 q5 0.63 0.53 0.32 0.12



#### Trend Analysis Example trends are linear contrasts > contrasts(df3\$grade) <- contr.poly(n=5,scores=1:5)</pre> > aov.trends <- aov(score~grade,data=df3) r = 0.97 > summary(aov.trends, + split=list(grade=list(Lin=1,Quad=2,Cube=3,Quart=4))) Df SS MS F Pr(>F)4 1361 340.3 2.258 0.07152 grade grade: Lin 1 1270 1269.9 8.425 0.00495 \*\* grade: Quad 1 1 0.8 0.005 0.94354 grade: Cube 1 80 80.4 0.533 0.46760 0.068 0.79530 grade: Quart 1 10 10.2 Residuals 70 10551 150.7 Grade H0 & H1 defined by trend weights: H0: $-0.63\mu_1 - 0.32\mu_2 + 0.32\mu_4 + 0.63\mu_5 = 0$ H1: $-0.63\mu_1 - 0.32\mu_2 + 0.32\mu_4 + 0.63\mu_5 \neq 0$







effect size & association strength

# Effect Size for a Linear Comparison

linear contrasts are used to compare two weighted means, so Cohen's d is approprate

Cohen's d (for a contrast)

$$d = 2\Psi / \left( \sigma_e \left[ \sum_{j=1}^a |c_j| \right] \right)$$

$$d = 2\hat{\Psi} / \left(\sqrt{\mathrm{MS}_W}\left[\sum_{j=1}^a |c_j|\right]\right)$$

Expresses  $\Psi$  in terms of the number of standard deviations of population error distribution

# Effect Size

Cohen's d calculation with emmeans & linear.comparison

> library(	(emmeans	s)		
> aov.01	<- aov(so	core~grade	,data=df3)	
> sigma	<- sigma(	aov.01) # s	sqrt(MS.res	sid)
> edf <- (	df.residua	l(aov.01) #	residual d	lf
> aov.em	n <- emme	eans(aov.0	1,specs="g	grade")
> mvCor	ntrasts <-	list(c1=mv	C1.c2=mv(	C2,c3=myC3,c4=myC4)
		en's d for e	1.1	
_	· · · · ·			yContrasts)
contrast	t effect.siz	e SE df	lower.CL	upper.CL
c1	0.53	0.24 70	0.05	1.01
c2	0.47	0.37 70	-0.26	1.20
сЗ	0.37	0.32 70	-0.26	1.01
c4	0.36	0.37 70	-0.37	1.09
	$\square$			

### Effect Size Cohen's d calculation with emmeans & linear.comparison > y <- df3\$score > g <- df3\$grade > myContrast1 <- linear.comparison(y,g,c.weights = myContrasts,var.equal=T) [1] "computing linear comparisons assuming equal variances among groups" [1] "C 1: F=4.995, t=2.235, p=0.029, psi=6.467, Cl=(0.367,12.568), adj.Cl= (-0.952,13.887)" [1] "C 2: F=1.662, t=1.289, p=0.202, psi=5.780, Cl=(-4.615,16.175), adj.Cl= (-5.714,17.274)" [1] "C 3: F=1.396, t=1.182, p=0.241, psi=4.588, CI=(-2.544,11.719), adj.CI= (-5.366,14.542)" [1] "C 4: F=0.977, t=0.989, p=0.326, psi=4.432, Cl=(-2.955,11.819), adj.Cl= (-7.062,15.926)" > myContrast1[[1]]\$d.effect.size [1] 0.53 > myContrast1[[2]]\$d.effect.size Note double brackets [[x]]! [1] 0.47 > myContrast1[[3]]\$d.effect.size [1] 0.37 > myContrast1[[4]]\$d.effect.size [1] 0.36

## Association Strength for a Linear Comparison - Proportion of Between-Groups variation accounted for by contrast $R_{alerting}^2 = SS_{contrast}/SS_B \qquad - \frac{Proportion of between or output variation}{2} \text{ with equal n, equals squared correlation between contrast weights \& }$ group means $R_{effectsize}^2 = SS_{contrast}/SS_{Total}$ - <u>Proportion of total variation</u> accounted for by contrast - Variation accounted for by contrast relative to the sum of $R_{contrast}^2 = SS_{contrast} / (SS_{contrast} + SS_W)$ contrast-variation and within-group (error) variation - Not affected by groups that are weighted zero - More resistant to changes in experimental design (e.g., adding or removing groups).

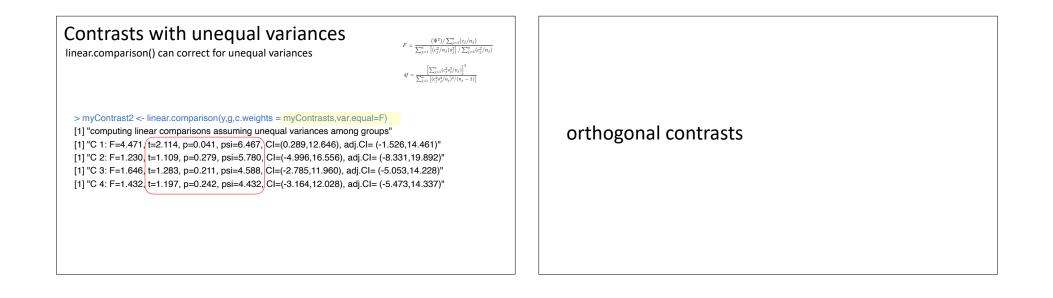
Association Streng	$R_{alerting}^2 = SS_{contrast}/SS_B$
• str(myContrast1[[1]]) ist of 15 \$ contrast : num [1:5] -0.5 -0.5 \$ F : num 4.99 \$ t : num 2.23 \$ df1 : num 1 \$ df2 : int 70 \$ p.2tailed : num 0.0286 \$ psi : num 0.0286 \$ psi : num 6.47 \$ confinterval : num [1:2] 0.367 1 \$ adj.confint : num [1:2] 0.952 1 \$ adj.confint : num [1:2] -0.952 1 \$ adj.confint : num 753 \$ d.effect.size : num 0.553 \$ R2.effect.size : num 0.553 \$ R2.effect.size : num 0.0632 \$ R2.effect.size : num 0.0632 \$ R2.effect.size : num 0.0632	$R_{effectsize}^2 = SS_{contrast}/SS_{Total}$ $R^2 = -SS_{contrast}/SS_{Total}$



# **Unequal Group Variances**

- So far our tests assume equal variance in different groups
- F/t tests for contrasts are not robust to violation of equal variance assumption
- When groups have unequal variances, use a different method to calculate F/t denominator, which is an estimate of population error variance
- Correcting for unequal var reduces denominator df (and, hence, power)

$$=\frac{(\Psi^2)/\sum_{j=1}^a (c_j/n_j)}{\sum_{j=1}^a \left[ (c_j^2/n_j) s_j^2 \right] / \sum_{j=1}^a (c_j^2/n_j)} \qquad df = \frac{\left[ \sum_{j=1}^a (c_j^2 s_j^2/n_j) \right]^2}{\sum_{j=1}^a \left[ (c_j^2 s_j^2/n_j)^2 / (n_j - 1) \right]}$$



F

# **Orthogonal Contrasts**

Equal n:

Unequal n:

$$\sum_{j=1}^{a} (c_{1j}c_{2j}) = 0 \qquad \qquad \sum_{j=1}^{a} (c_{1j}c_{2j}/n_j) = 0$$

A set of contrasts is <u>mutually orthogonal</u> if all pairs of contrasts are orthogonal

Orthogonal contrasts evaluate independent questions about group means

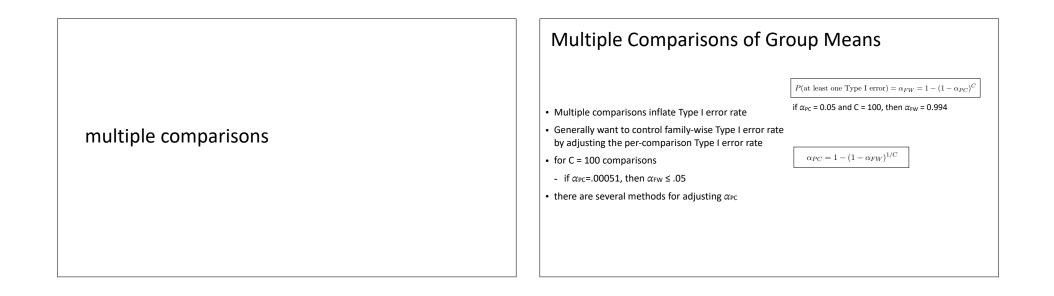
### Complete Set of Mutually Orthogonal Contrasts

If there are a groups, then the largest set of mutually orthogonal contrasts will have (a-1) contrasts, and:

$$\sum_{j=1}^{a-1} \mathrm{SS}_{contrast,j} = \mathrm{SS}_B$$

A complete set of orthogonal contrasts divides SS<sub>B</sub> into independent pieces of variation,
the sum of the (*a-1*) SS<sub>contrasts</sub> will equal SS<sub>B</sub>,
and the average of the contrast F values will equal the omnibus F.

Complete set of orthogonal contrasts
breaks SS <sub>group</sub> into separate pieces
$ > cMat <- contrasts(df3Sgrade) > fractions(cMat) myC1 myC2 myC3 myC4 g3 1/2 -1 0 0 g3 1/3 0 -1 0 g4 1/3 0 1/2 -1 g5 1/3 0 1/2 1 > fmeter contrasts/columns are mutually orthogonal: > round(tfMat) e^{S} cMat.digits=2)myC1 myC2 myC3 myC4myC1 0.83 0 0.0 0myC2 0.00 2 0.0 0myC2 0.00 1.5 0myC4 0.00 0 0.0 2$



# Controlling False Discovery Rate

- Instead of controlling  $\alpha_{FW}$ , control **False Discovery Rate** (FDR):
- Q = (# of false H0 rejections) / (total # H0 rejections)
- FDR = Expected Value[Q]
- When all H0 are true, controlling  $\alpha_{\text{FW}}$  and FDR are equivalent
- When some H0 are false, FDR-based methods are more powerful

# **Corrections for Multiple Comparisons**

- Controlling  $\alpha_{FW}$  by adjusting  $\alpha_{PC}$ :
- Bonferroni Adjustment (aka Dunn's Procedure)
- Holm's Sequential Bonferroni Test
- Controlling False Discovery Rate (FDR):
- Benjamini & Hochberg's (1995) Linear Step-Up Procedure (FDR)
- Relative Power: FDR > Holm's > Bonferroni

Multiple Comparisons in R adjust p values with p.adjust()	Controlling Type I error rate p.adjust()
<pre>&gt; my.p.values &lt;- c(.127,.08,.03,.032,.02,.001,.01,.005,.025) &gt; sort(my.p.values) [1] 0.001 0.005 0.010 0.020 0.025 0.030 0.032 0.080 0.127 &gt; p.adjust(sort(my.p.values),method='bonferroni') [1] 0.009 0.045 0.090 0.180 0.225 0.270 0.288 0.720 1.000 &gt; p.adjust(sort(my.p.values),method='holm') [1] 0.009 0.040 0.070 0.120 0.125 0.125 0.125 0.160 0.160 &gt; p.adjust(sort(my.p.values),method='fdr')</pre>	<pre>&gt; summary(aov.vp,split=list(complexity=list(L=1,Q=2,C=3,q4=4)))</pre>
<ul> <li>[1] 0.009 0.0225 0.0300 0.04114 0.04114 0.04114 0.04114 0.090 0.127</li> <li>Significant tests (alpha/FDR = .05) are highlighted in orange font.</li> <li>N.B. Sorting p-values is not required.</li> </ul>	<pre>&gt; p.adjust(p=c(0.000532,0.006100,0.340714,0.363286),method="bonferroni") [1] 0.002128 0.024400 1.000000 1.000000 &gt; p.adjust(p=c(0.000532,0.006100,0.340714,0.363286),method="holm") [1] 0.002128 0.018300 0.681428 0.681428 &gt; p.adjust(p=c(0.000532,0.006100,0.340714,0.363286),method="fdr") [1] 0.002128 0.012200 0.363286 0.363286</pre>

# Controlling Type I error rate

### emmeans

> aov.vp <- aov(visPref~complexity,data=df4)
> vp.em <- emmeans(aov.vp,specs="complexity")</pre>

### > contrast(vp.em,method="poly",adjust="fdr")

 contrast
 estimate
 SE
 df
 t
 p

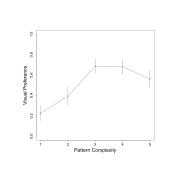
 linear
 0.964
 0.253
 35
 3.815
 0.0021

 quadratic
 0.873
 0.299
 35
 -2.919
 0.0122

 cubic
 -0.244
 0.253
 35
 -0.966
 0.3633

 quartic
 0.616
 0.669
 35
 0.921
 0.3633

P value adjustment: fdr method for 4 tests



# Setting family-wise alpha and FDR

- Generally,  $\alpha_{\text{FW}}$  and FDR are set to 0.01 or 0.05
- larger α<sub>FW</sub> may be justified for small number of orthogonal comparisons
- Bonferroni & Holm tests may reduce power too much
- perhaps set  $\alpha_{PC}$  to 0.05 or 0.01
- family-wise Type I error will increase but Type II error will decrease
- Note: we do this with factorial ANOVA already...

# All pairwise tests (Tukey HSD)

- Tukey HSD evaluates all pairwise differences between groups
- Is more powerful than Bonferroni method (for between-subj designs)
- Tukey HSD:
- NOT necessary to evaluate omnibus F prior to Tukey test
- assumes equal n per group & equal variances
- Tukey-Kramer is valid with sample sizes are unequal
- Dunnett's T3 test is better with unequal n & unequal variances [see Kirk (1995, pp. 146-50) for more details]

# Tukey HSD (all pairwise differences)

optimal method for evaluating all pairwise differences

### assumes equal variances

### > TukeyHSD(aov.vp,which="complexity")

p4-p2 0.2906 -0.035 0.62 0.10

p5-p2 0.1706 -0.154 0.50 0.56 p4-p3 -0.0051 -0.330 0.32 1.00

p5-p3 -0.1250 -0.450 0.20 0.80 p5-p4 -0.1199 -0.445 0.21 0.83

Tukey multiple comparisons of means 95% family-wise confidence level Fit: aov(formula = visPref ~ complexity, data = df4) \$complexity diff lwr upr p adj p2-p1 0.1663 -0.159 0.49 0.59 p3-p1 0.4620 0.137 0.79 0.00 p4-p1 0.4569 0.012 0.66 0.00 p5-p1 0.3569 0.012 0.66 0.00 p3-p2 0.2957 -0.029 0.62 0.09

### does not assume equal variances

> library(PMCMRplus)
> dunnettT3Test(x=df4\$visPref,g=df4\$complexity)

Pairwise comparisons using Dunnett's T3 test for multiple comparisons with unequal variances

data: df4\$visPref and df4\$complexity

 p1
 p2
 p3
 p4

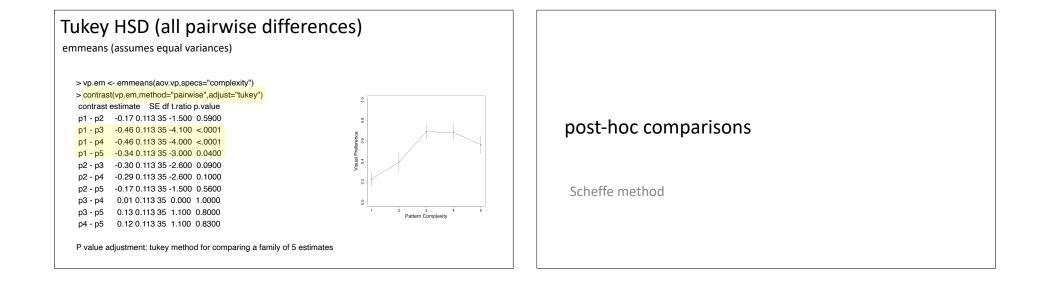
 p2
 0.8081

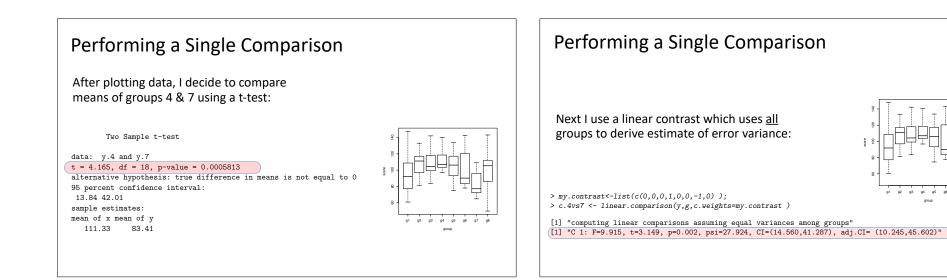
 p3
 0.0037
 0.1934

 p4
 0.0057
 0.2340
 1.0000

 p5
 0.0834
 0.8475
 0.9321
 0.9563

P value adjustment method: single-step alternative hypothesis: two.sided





### What was wrong with the preceding analyses?

**Answer:** I performed the analyses after inspecting the data and choosing to compare groups 4 & 7 <u>because they looked different</u> which, obviously, inflates Type I error

## Planned vs. Post-hoc Comparisons

- Previous comparisons were planned
- Last 2 comparisons, made after looking at data, were post-hoc
- Scheffe method is preferred for post-hoc linear contrasts
- compute contrast with normal procedures
- evaluate observed F with new critical value:
- F<sub>Scheffe</sub> = (a-1) x F<sub>α(FW)</sub> (df1= a-1; df2 = N-a)
- ▶ a = number of groups
- +  $F_{\alpha(FW)}$  is the F value required for desired alpha
- F<sub>Scheffe</sub> is "normal" omnibus F x (a-1)
- alternatively, keep standard F & adjust p values using Scheffe adjustment
- Scheffe method and omnibus F test are mutually consistent

for post-hoc comparisons > c1 <- c(-3, -3, 2, 2, 2) > c2 <- c(-1, 1, 0, 0) > (con1 <- contrast(vp.em,method=list(c1, c2),adjust="none")) contrast estimate SE df t.ratio p.value c(-3, -3, 2, 2, 2) 2.013 0.438 35 4.596 0.0001 c(-1, 0, 0, 0) = 0.166 0.143 25 4.471 0.4502	These methods compute normal F and adjust the p
<pre>&gt; c2 &lt;- c(-1,1,0,0,0) &gt; (con1 &lt;- contrast(vp.em,method=list(c1,c2),adjust="none"))) contrast estimate SE df t.ratio p.value c(-3, -3, 2, 2, 2) 2.013 0.438 35 4.596 0.0001</pre>	These methods compute normal F and adjust the p
c(-1, 1, 0, 0, 0) 0.166 0.113 35 1.471 0.1503 > summary(con1,adjust="scheffe",scheffe.rank=4) contrast estimate SE df t.ratio p.value c(-3, -3, 2, 2, 2) 2.013 0.438 35 4.596 0.0019 c(-1, 1, 0, 0, 0) 0.166 0.113 35 1.471 0.7068 P value adjustment: scheffe method with rank 4	value to be consistent with Scheffe method Scheffe.rank should be set to degrees of freedom fo grouping factor (i.e., a-1)
	contrast         estimate         SE df t.ratio p.value           c(-3, -3, 2, 2, 2)         2.013 0.438 35         4.596         0.0019           c(-1, 1, 0, 0, 0)         0.166 0.113 35         1.471         0.7068