

Notation and Equations for Exam 3

r	Sample correlation
m	The number of predictor variables in a regression
X_i	A predictor variable in a regression. The subscript i represents any number from 1 through m .
Y	The outcome variable that is being predicted or explained in a regression
\hat{Y} (Y-hat)	The estimated outcome value as predicted by the regression equation
b_i	The regression coefficient for predictor X_i (sometimes written as $b_{\text{predictor name}}$)
b_0	The intercept in the regression equation
σ_{b_i}	The standard error of a regression coefficient
SS_Y	The total sum of squares for the outcome in a regression
$SS_{\text{regression}}$	The sum of squares explained by the predictors in a regression
R^2	The proportion of variability explained by a regression
SS_{total}	The total variability in the data for an ANOVA
$SS_{\text{treatment}}$	Variability explainable by differences among groups (simple ANOVA) or measurements (repeated measures)
SS_{factor}	Variability explainable by the main effect of some factor
$SS_{A:B}$	Variability explainable by interaction between factors A and B
SS_{residual}	The residual sum of squares, representing the variability that can't be explained in regression or ANOVA
MS_{effect}	Mean square for any effect we might want to test; the subscript can be regression, treatment, Factor, A:B, etc.
df_{effect}	Degrees of freedom for SS_{effect} and MS_{effect} , where <i>effect</i> is any effect we might want to test
MS_{residual}	The residual mean square; used as an estimate of the population variance, σ^2 or σ_Y^2
df_{residual}	The degrees of freedom for SS_{residual} and MS_{residual}
F	F statistic
k	The number of levels of a factor (treatment) in an ANOVA; written as k_{Factor} when there are multiple factors
M_i	The sample mean of Group i in a simple ANOVA or Measurement i in a repeated-measures ANOVA ($i = 1$ to k)
M_s	The mean of all measurements from Subject s in a repeated-measures ANOVA ($s = 1$ to n)
n_i	The number of data (e.g., subjects) in Group i
\bar{M}	The grand mean, i.e. the mean of all the data in all groups taken together

Formula for correlation	$r = \frac{\sum(z_X \cdot z_Y)}{n - 1}$
Interpreting correlation	$r = -1 \rightarrow$ perfect negative relationship $r < 0 \rightarrow$ negative relationship $r = 0 \rightarrow$ no <u>linear</u> relationship $r > 0 \rightarrow$ positive relationship $r = 1 \rightarrow$ perfect positive relationship
Relationship between correlation and prediction	$z_{\hat{Y}} = r \cdot z_X$
Regression equation	$\hat{Y} = b_0 + b_1X_1 + b_2X_2 + \dots + b_mX_m = b_0 + \sum_{i=1 \text{ to } m} b_iX_i$
Total variability in a regression	$SS_Y = \sum(Y - M_Y)^2$
Residual variability in a regression	$SS_{\text{residual}} = \sum(Y - \hat{Y})^2$
Variability explained by a regression	$SS_{\text{regression}} = SS_Y - SS_{\text{residual}}$
Proportion of variability explained by regression	$R^2 = \frac{SS_{\text{regression}}}{SS_Y}$
Explained variability with one predictor	$R^2 = r^2$
t statistic for a regression coefficient	$t = \frac{b_i}{\sigma_{b_i}}$
Total sum of squares in an ANOVA	$SS_{\text{total}} = \sum(X - \bar{M})^2$
Residual sum of squares in a simple ANOVA	$SS_{\text{residual}} = \sum(X_1 - M_1)^2 + \sum(X_2 - M_2)^2 + \dots + \sum(X_k - M_k)^2 = \sum_i \left(\sum(X_i - M_i)^2 \right)$
Treatment sum of squares for simple ANOVA	$SS_{\text{treatment}} = \sum_i n_i \cdot (M_i - \bar{M})^2$

Mean squares from sum of squares	$MS_{\text{regression}} = \frac{SS_{\text{regression}}}{df_{\text{regression}}}$ $MS_{\text{treatment}} = \frac{SS_{\text{treatment}}}{df_{\text{treatment}}}$ $MS_{\text{factor}} = \frac{SS_{\text{factor}}}{df_{\text{factor}}}$ $MS_{\text{A:B}} = \frac{SS_{\text{A:B}}}{df_{\text{A:B}}}$ $MS_{\text{residual}} = \frac{SS_{\text{residual}}}{df_{\text{residual}}}$
F statistic for explained variability	$F = \frac{MS_{\text{regression}}}{MS_{\text{residual}}}$ $F = \frac{MS_{\text{treatment}}}{MS_{\text{residual}}}$ $F_{\text{factor}} = \frac{MS_{\text{factor}}}{MS_{\text{residual}}}$ $F_{\text{A:B}} = \frac{MS_{\text{A:B}}}{MS_{\text{residual}}}$
p-value for explained variability	$p = p\left(F_{df_{\text{regression}}, df_{\text{residual}}} \geq F\right)$ $p = p\left(F_{df_{\text{treatment}}, df_{\text{residual}}} \geq F\right)$ $p_{\text{factor}} = p\left(F_{df_{\text{factor}}, df_{\text{residual}}} \geq F_{\text{factor}}\right)$ $p_{\text{A:B}} = p\left(F_{df_{\text{A:B}}, df_{\text{residual}}} \geq F_{\text{A:B}}\right)$
Partitioning variability for regression	$SS_Y = SS_{\text{regression}} + SS_{\text{residual}}$
Partitioning variability for simple ANOVA	$SS_{\text{total}} = SS_{\text{treatment}} + SS_{\text{residual}}$
Partitioning variability for repeated measures	$SS_{\text{total}} = SS_{\text{treatment}} + SS_{\text{subject}} + SS_{\text{residual}}$
Partitioning variability for factorial ANOVA	$SS_{\text{total}} = SS_{\text{A}} + SS_{\text{B}} + SS_{\text{C}} + \dots \text{ [every main effect]}$ $+ SS_{\text{A:B}} + SS_{\text{A:C}} + SS_{\text{B:C}} + \dots \text{ [every 2-way interaction]}$ $+ SS_{\text{A:B:C}} + \dots \text{ [every higher-order interaction, up to the total number of factors]}$ $+ SS_{\text{residual}}$
Recognizing an interaction	$M_{a_1, b_1} - M_{a_2, b_1} \neq M_{a_1, b_2} - M_{a_2, b_2} \rightarrow \text{Interaction}$