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Abstract

Multiple linear regression (MLR) remains a mainstay analysis in organizational research, yet inter-correlations between predictors (multicollinearity) undermine the interpretation of MLR weights in terms of predictor contributions to the criterion. Alternative indices include validity coefficients, structure coefficients, product measures, relative weights, all-possible-subsets regression, dominance weights, and commonality coefficients. This article reviews these indices, and uniquely, it offers freely available software that (a) computes and compares all of these indices with one another, (b) computes associated bootstrapped confidence intervals, and (c) does so for any number of predictors so long as the correlation matrix is positive definite. Other available software is limited in all of these respects. We invite researchers to use this software to increase their insights when applying MLR to a data set. Avenues for future research and application are discussed.

Keywords

multiple regression, quantitative research, exploratory, research design

A continued goal of organizational researchers conducting regression analysis is to make inferences about the relative importance of predictor variables (cf. Nimon, Gavrilova, & Roberts, 2010; Zientek, Capraro, & Capraro, 2008), yet it is all too common to rely heavily (if not solely) on the regression coefficients from the analysis which optimize sample-specific prediction (minimize sum of squared errors). Instead, other metrics that operationalize relative importance in ways that are consistent with such researchers' goals would seem more appropriate, and a range of metrics and approaches exists. In addition to regression weights and zero-order correlation coefficients that researchers likely report, MLR interpretation may be further informed by considering structure

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coefficients, product measures, relative weights, all-possible-subsets regression, dominance weights, and commonality coefficients. The current article offers software that cogently summarizes these metrics so that researchers can make more sophisticated judgments about the nature and meaningfulness of variables in a linear regression model than judgments from regression weights or any single metric in isolation. As noted by Nathans, Oswald, and Nimon (2012), each metric serves a different purpose and has certain features that support interpreting specific aspects of a multiple linear regression (MLR) model (see Table 1).

The aforementioned metrics are reviewed here briefly, but for more details we refer the reader to other work, both recent and historical in nature (e.g., Budescu, 1993; Darlington, 1968; Johnson, 2000; Johnson & LeBreton, 2004; Kraha, Turner, Nimon, Zientek, & Henson, 2012; Krasikova, LeBreton, & Tonidandel, 2011; Lindeman, Merenda, & Gold, 1980; Tonidandel & LeBreton, 2011). This allows the current article to take a more practical focus on the software tool and its capabilities, with the support of two empirical examples.

Regression Weights

In MLR models, raw data yield *unstandardized (raw) regression weights*, and standardized data yield *standardized regression weights*. Regardless of whether or not the data are standardized, the values residing in the vector $\mathbf{b} = (b_1, b_2, b_3, \dots, b_p)$ are chosen in such a way that the weighted composite $\mathbf{b}^T \mathbf{X}$ is maximally correlated with the dependent variable, Y , which is external to the composite (i.e., choose \mathbf{b} so that $r[\mathbf{b}^T \mathbf{X}, Y]$ is maximized). This is the same as choosing \mathbf{b} to minimize the variance of the errors in prediction (i.e., choose \mathbf{b} so that $[\text{var}(Y) - \mathbf{b}^T \mathbf{R}_{\mathbf{X}\mathbf{X}} \mathbf{b}]$ is minimized). If we assume that we are using standardized regression coefficients, then each coefficient b_k indicates the expected change in Y , in standard deviation units, given a corresponding 1 standard deviation change in X_k , when all the other predictors in the model in \mathbf{X} (i.e., $(X_1, X_2, \dots, X_{k-1}, X_{k+1}, \dots, X_p)^T$) are fixed or controlled for.

It may be reasonable to assume that predictors with larger standardized coefficients (called betas henceforth) are more important than other predictors with smaller coefficients. Certainly this is true when variables are uncorrelated, because in that case, betas are exactly equal to the zero-order correlations between \mathbf{X} and Y . However, when the predictors in \mathbf{X} are correlated, standardizing does not disentangle the effects of \mathbf{X} on Y from the standard deviations of \mathbf{X} ; in fact, it confounds them in the service of placing all weights on a z-score metric.

Zero-Order Correlation Coefficients

Obviously, prediction of Y from each independent variable X_i is found in $\mathbf{R}_{\mathbf{X}\mathbf{Y}}$, the vector of zero-order correlation coefficients (often called validity coefficients). If all variables in \mathbf{X} were completely uncorrelated (i.e., $\mathbf{R}_{\mathbf{X}\mathbf{X}} = \mathbf{I}$, the $p \times p$ identity matrix), then the contribution of each X_i to Y can be clearly represented as the squared values in $\mathbf{R}_{\mathbf{X}\mathbf{Y}}$, because these squared values are independent of one another, meaning that they partition R^2 or, equivalently, they add up to the model R^2 (i.e., $\mathbf{R}'_{\mathbf{X}\mathbf{Y}} \mathbf{R}_{\mathbf{X}\mathbf{Y}} = R^2$). However, when predictors are correlated, the sum of the elements in $\mathbf{r}_{\mathbf{X}\mathbf{Y}}^2$ will not add up to the model R^2 , necessitating other metrics to determine how the MLR model is affected by multicollinearity.

Structure Coefficients

One relatively simple approach to determining the contribution of p independent variables in linear regression is through calculating a $p \times 1$ vector of *structure coefficients (sr)*, which are the correlations between each predictor X and \hat{Y} , the latter being the predicted values of Y . Equivalently,

Table 1. Regression Metric/Analysis and Their Associated Purpose and Features.

Regression Metric/Analysis	Purpose	Variable Importance ^a	Partition R^2	Identify Presence of Multicollinearity and Suppression?
Regression weight	Identifies contribution of each IV to the regression equation, holding all other IVs constant	Total	Into p partitions, with standardized data and uncorrelated IVs	With zero-order correlation or structure coefficients
Zero-order correlation coefficient	Identifies magnitude and direction of the relationship between the IV and DV, without considering any other IVs in the MLR model	Direct	Into p partitions, when squared and with uncorrelated IVs	With regression weights
Squared structure coefficient	Identifies how much variance in the predicted scores (\hat{Y}) for the DV can be attributed to each IV	Direct	Into p partitions, with uncorrelated IVs	With regression weights and when summed and compared to unity
Pratt measure	Partitions the regression effect into nonoverlapping partitions based on multiplying the beta weight of each independent variable with its respective zero-order correlation with the dependent variable	Direct and total	Into p partitions	May identify presence of multicollinearity or suppression
Relative weight	Determines how IVs contribute to the DV when expressed as a joint function of (a) how highly related IVs are to their uncorrelated counterpart and (b) how highly related the uncorrelated counterparts relate to the DV	Total	Into p partitions	Do not identify presence of multicollinearity or suppression
All-possible-subsets regression	Determines the R^2 for MLRs from all possible $2^p - 1$ subsets of IVs; serves as base for commonality and dominance analysis	N/A	N/A	May identify presence of multicollinearity or suppression
Commonality analysis	Identifies how much variance in the DV is uniquely explained by each possible IV set in the MLR; yields unique and common effects that, respectively, identify variance unique to one IV and multiple IVs	Total	Into $2^p - 1$ partitions.	Identify presence, location, and magnitude of multicollinearity and suppression
Dominance analysis	Indicates whether one IV contributes more unique variance than another IV, either (a) across all possible MLR submodels (i.e., complete dominance) or (b) on average across models of all-possible-subset sizes (i.e., conditional dominance); averaging conditional dominance weights yields general dominance weights	Direct, total, and partial	Into p partitions with general dominance weights	Conditional dominance weights may identify presence of suppression

Note: Adapted from Nathans, Oswald, and Nimon (2012). p = number of predictor; IV = independent variable; DV = dependent variable; MLR = multiple linear regression.

^aQuantifies IV importance when: (a) measured in isolation from other IVs (direct effect), (b) contributions of all other IVs have been accounted for (total effect), and (c) contributions to regression models of a specific subset or subsets of other IVs have been accounted for (partial effect). See Nathans et al. (2012) for more on advantages, disadvantages, and recommendations for practice.

structure coefficients can be calculated by dividing the $p \times 1$ vector of zero-order validity coefficients R_{XY} by the multiple R ; thus,

$$sr = \frac{R_{XY}}{\sqrt{b^T R_{XY}}} = \frac{R_{XY}}{R} = r_{X\hat{Y}}. \quad (1)$$

Structure coefficients have been used to indicate the relative contributions of each X_i to the prediction of Y , but they cannot be interpreted in a straightforward manner when variables in X are correlated or when there are suppression effects (i.e., the sum of the elements in $r^2_{X\hat{Y}}$ does not equal unity).

Pratt Measure

The Pratt measure, or product measure, of a predictor variable's relative importance was proposed by Pratt (1987) and is defined simply as $m_i = b_{YX_i} r_{YX_i}$; the product of the standardized regression coefficient and the zero-order validity coefficient for X_i . Pratt measures divide the model R^2 across predictor variables, meaning that

$$R^2 = b^T R_{XY} = \sum_{i=1}^p m_i, \quad (2)$$

where each component of the sum is a Pratt measure (m_i). The Pratt measure can be viewed as problematic in cases where individual values are negative or zero values, because those might be products of suppression or multicollinearity that require further explanation (see Thomas, Hughes, & Zumbo, 1998). However, this decomposition remains in the literature as a common method for decomposing R^2 , and we include it in our software package, if only to compare it with more modern methods such as general dominance and relative weights.

Relative Weights

Relative weights are another way to partition an MLR model R^2 across predictors. They are computed by first transforming p predictors into a new set of p variables that are uncorrelated with one another, yet are correlated as highly as possible with the original predictors; that is, given the data matrix $X = P\Delta Q^T$, then create the new variables $Z = PQ^T$. Fabbris (1980) noted that treating X as a set of dependent variables and regressing X onto Z creates M , the $p \times p$ correlation matrix between the original predictors and their orthogonal counterparts; this was rediscovered by Genizi (1993) and then in the organizational literature by Johnson (2000). It may be somewhat surprising to know that M is also equal to the matrix square root of the predictor intercorrelation matrix ($M = (Z'Z)^{-1}Z^T X = Z^T X = Q\Delta Q^T = R_{xx}^{1/2}$). Also surprising might be that the squared elements in each column (or each row) of M (or $R_{xx}^{1/2}$) sum to 1.

These elements for each column are used to weight each predictor variable according to its independence from other predictor variables (i.e., a higher weight will mean greater independence from that variable). In addition to this matrix, regressing Y on Z yields a $p \times 1$ vector of orthogonal weights we call V (i.e., $V = (Z'Z)^{-1}Z^T y = Z^T y$). Because the components of Z are uncorrelated and all variables are standardized, the squared values in V also sum to 1, where each element provides the proportional contribution of each orthogonalized predictor Z_i to Y .

Thus, a relative weight multiplies or "glues" like elements from the two orthogonal vectors of M and V weights just described, namely, (a) the i th squared regression weight for the given predictor regressed onto its orthogonal counterpart multiplied by (b) the i th squared weight regressing the

criterion on that given predictor's orthogonal counterpart. Mathematically, the relative weight for predictor i is called ε_i^2 and equals

$$\varepsilon_i^2 = \sum_{j=1}^p \{M_{ij}^2 V_j^2\}, \quad (3)$$

and each of the p relative weights across is an independent part of the total model R^2 , meaning the weights add to R^2 ($R^2 = \sum_{i=1}^p \varepsilon_i^2$) even when the independent variables are correlated (see Fabbris, 1980; Genizi, 1993; Johnson, 2000). In this way, relative weights are easy to explain in the same way as general dominance weights because they sum to the model R^2 . Alternatively, the relative weights can be re-expressed as proportions or percentages of R^2 .

Relative weights are thus defined as the contribution of a given predictor to criterion variance, considering the predictor's contribution alone as well as jointly with the other predictors in the model. Note that relative weights and general dominance weights have often been found to rank predictors similarly in terms of relative importance (Johnson, 2000); however, the current software program can verify where results converge or diverge.

All-Possible-Subsets Regression

As the name implies, all-possible-subsets (APS) regression involves running linear regression models on all $2^p - 1$ subsets of predictors. In doing so, one often takes either a predictor-based approach or model-based approach to the set of results. APS is best compared to other metrics when taking a predictor-based approach, where a predictor deemed more important within a regression model will tend to be one that is more important across submodels in APS regression. APS regressions can be analyzed in this manner for each predictor, or one can use commonality coefficients or dominance weights, to be discussed next, because these are based on the results of APS regressions.

In addition to the predictor-based approach, researchers and practitioners also may take a model-based approach to APS, because it is an exploratory approach to determining the tradeoff between model parsimony and model fit, where a submodel with fewer predictors retains a model R^2 that is either similar to the full model R^2 or is above some practical minimum established by the researcher or practitioner. Obviously, the model-based approach can be related to the variable-based approach when there is a consistent recommendation to include and/or exclude specific predictors across models.

Commonality Analysis

Commonality analysis partitions the R^2 explained by all predictors in multiple regression into variance unique to each predictor and variance shared between each combination of predictors (see Mayeske et al., 1969; Mood, 1969, 1971; Newton & Spurrell, 1967; Onwuegbuzie & Daniel, 2003; Rowell, 1996). These components of variance are called *commonality coefficients* that can then be evaluated in terms of their magnitude, and they can be compared with one another. As partitions, commonality coefficients sum to the total R^2 for the regression model.

There are two general types of commonality coefficients: *unique effects* and *common effects*. The unique effect of a predictor (also called the *uniqueness coefficient*) is the square of the semipartial correlation between a given predictor and the criterion. Thus, if predictors are all uncorrelated, then predictor importance can be entirely determined by ranking the unique effects. When predictors are correlated, as is usually the case, the common effects can indicate the extent and pattern of the predictors' shared variance in predicting variance in the criterion (Mood, 1971).

Consider the hypothetical situation discussed by Hedges and Olkin (1981) in which two variables, X1 and X2, are used to predict a variable X0. For this regression equation, the explained variance ($R^2_{0.12}$) can be partitioned into three components:

$$\gamma_1 = \text{unique contribution of X1 to } R^2_{0.12}, \quad (4)$$

$$\gamma_2 = \text{unique contribution of X2 to } R^2_{0.12}, \quad (5)$$

$$\gamma_{12} = \text{common contribution of X1 and X2 to } R^2_{0.12}, \quad (6)$$

and they are computed as follows:

$$\gamma_1 = R^2_{0.12} - R^2_{0.2}, \quad (7)$$

$$\gamma_2 = R^2_{0.12} - R^2_{0.1}, \quad (8)$$

$$\gamma_{12} = R^2_{0.1} + R^2_{0.2} - R^2_{0.12}. \quad (9)$$

Commonality coefficients provide researchers with rich detail about how independent variables operate together in a given regression model. The coefficients are more specific than regression weights, relative weights, or general dominance weights. As Seibold and McPhee (1979) noted, “[Only by] determining the extent to which . . . independent variables, singly and in all combinations, share variance with the dependent variable . . . can we fully know the relative importance of independent variables with regard to the dependent variable in question” (p. 355).

Note that negative values of commonality coefficients generally indicate that a predictor exerts a suppressor effect, where it is removing (partialing out) the irrelevant variance in the other predictor(s) to increase the latter’s contributions to the model R^2 (Zientek & Thompson, 2006). Unlike other metrics, commonality coefficients are uniquely able to pinpoint the predictors involved in a suppressor relationship and the specific nature of that relationship. Summing all negative common effects for a regression equation can quantify the amount of suppression present in the regression model as a whole.

Dominance Analysis

A regression metric was originally proposed by Chevan and Sutherland (1991), then elaborated upon by Budescu (1993), who detailed the procedure for conducting a dominance analysis. Dominance analysis involves computing each predictor’s incremental validity across all possible submodels that involve that predictor and using the incremental validity coefficients to evaluate complete dominance, conditional dominance, and general dominance. To make the dominance analysis procedure more concrete, see the dominance analysis formulas in Table 2 that support a three-predictor MLR model.

If incremental validity is always higher for X_i than for X_j for every submodel, then X_i is said to show *complete dominance* over X_j . Complete dominance is a restricted form of dominance that may rarely occur. There is a more relaxed form of dominance called *conditional dominance*, which occurs when the average incremental variance within each submodel of sizes 0 to $p - 1$ is greater for one predictor than another across all model sizes. The average incremental variance components used to evaluate conditional dominance are called conditional dominance weights. Conditional dominance weights are averaged across p predictors to create general dominance weights that partition the model R^2 across predictors. A predictor is said to show general dominance if it has the highest overall average incremental validity across regression submodels of sizes 0 to $p - 1$.

Dominance weights have two appealing properties: First, like relative weights, each general dominance weight is the average contribution of a predictor to a criterion, both on its own and when

Table 2. Dominance Analysis Formulas for Three Predictor Case.

Subset Model (X)	$R^2_{Y \cdot X}$	Additional Contribution of:		
		X1	X2	X3
CD, k = 0 average				
X1	$R^2_{Y \cdot X1}$	$R^2_{Y \cdot X2}$	$R^2_{Y \cdot X3}$	$R^2_{Y \cdot X3}$
X2	$R^2_{Y \cdot X2}$	$R^2_{Y \cdot X1X2} - R^2_{Y \cdot X1}$	$R^2_{Y \cdot X1X3} - R^2_{Y \cdot X1}$	$R^2_{Y \cdot X1X3} - R^2_{Y \cdot X1}$
X3	$R^2_{Y \cdot X3}$	$R^2_{Y \cdot X1X2} - R^2_{Y \cdot X2}$ $R^2_{Y \cdot X1X3} - R^2_{Y \cdot X3}$	$R^2_{Y \cdot X2X3} - R^2_{Y \cdot X3}$	$R^2_{Y \cdot X2X3} - R^2_{Y \cdot X2}$
CD, k = 1 average		$\frac{(R^2_{Y \cdot X1X2} - R^2_{Y \cdot X2}) + (R^2_{Y \cdot X1X3} - R^2_{Y \cdot X3})}{2}$	$\frac{(R^2_{Y \cdot X1X2} - R^2_{Y \cdot X1}) + (R^2_{Y \cdot X2X3} - R^2_{Y \cdot X3})}{2}$	$\frac{(R^2_{Y \cdot X1X3} - R^2_{Y \cdot X1}) + (R^2_{Y \cdot X2X3} - R^2_{Y \cdot X2})}{2}$
X1X2	$R^2_{Y \cdot X1X2}$			$R^2_{Y \cdot X1X2X3} - R^2_{Y \cdot X1X2}$
X1X3	$R^2_{Y \cdot X1X3}$		$R^2_{Y \cdot X1X2X3} - R^2_{Y \cdot X1X3}$	
X2X3	$R^2_{Y \cdot X2X3}$	$R^2_{Y \cdot X1X2X3} - R^2_{Y \cdot X2X3}$		
CD, k = 2 average		$\frac{R^2_{Y \cdot X1X2X3} - R^2_{Y \cdot X2X3}}{2}$	$R^2_{Y \cdot X1X2X3} - R^2_{Y \cdot X1X3}$	$R^2_{Y \cdot X1X2X3} - R^2_{Y \cdot X1X2}$
X1X2X3	$R^2_{Y \cdot X1X2X3}$			
GD, overall average		$\frac{\sum_{k=0}^2 CD_{X1}}{3}$	$\frac{\sum_{k=0}^2 CD_{X2}}{3}$	$\frac{\sum_{k=0}^2 CD_{X3}}{3}$

Note: Blank cells are not applicable. CD = conditional dominance; GD = general dominance.

taking all other predictors in the model into account. Second, general dominance weights always sum to the overall model R^2 . Third, conditional dominance weights have the potential to illuminate the properties of model predictors that can get lost in commonality coefficients (which are numerous and can be difficult to interpret even when $p = 3$) or averaged away in more general metrics (e.g., relative weights, general dominance weights).

Software for Exploring the Predictor Space

The software tool we introduce was developed in R (R Development Core Team, 2013), and it offers a number of incremental contributions over both the literature and applications of the past addressing this topic of the relative importance of variables in linear regression analysis. First and most importantly, the software analyzes a wide variety of regression metrics within a single program. Recently, researchers have developed and provided computer programs, macros, and software packages to compute the metrics in linear regression to be discussed here, such as dominance analysis (Azen & Budescu, 2003) and relative weights (Grömping, 2006; Tonidandel, LeBreton, & Johnson, 2009). Although other researchers have integrated multiple metrics into their programs (Braun & Oswald, 2011), the current software is much more comprehensive in integrating all regression metrics reviewed here. Second, the software allows for some computational efficiencies; for instance, all-possible-subsets regression is computed and applied to multiple metrics rather than having to be computed each time within different programs. Third, the software computes confidence intervals (CIs) for all coefficients and for differences between specific pairs of coefficients. Confidence intervals are based on bootstrapping procedures and estimate an interval that 95% of the time, for example, contains the corresponding population parameter. Although this has been accomplished for some metrics (e.g., Algina, Keselman, & Penfield, 2010, for squared semipartial correlations; Lorenzo-Seva, Ferrando, & Chico, 2010, for bootstrapped regression coefficients, structure coefficients, and relative importance weights; and Tonidandel & LeBreton at the website <http://relativeimportance.davidson.edu/> for bootstrapped relative importance weights in regression, multivariate regression [regression with multiple criteria], and logistic regression), this has not been accomplished to date across such a wide array of metrics within a more integrated software package as we have done here. To conduct statistical significance tests of metrics such as relative weights and general dominance weights, as in Tonidandel et al. (2009), one can include a randomly generated variable as an additional predictor in the model. This software will then automatically generate CIs on the differences between each predictor metric and the metric associated with the randomly generated variable, which can then be used to assess statistical significance. Our software program also graphs CIs, keeping in mind that the statistical differences between statistics may still be significant even when their respective CIs overlap (Cumming & Finch, 2005). Fourth, some software solutions have limited the number of predictors that are allowed; by contrast, our software computes metrics for as many predictors as are allowed by computer memory, by the positive definiteness of the predictor correlation matrix, and by the patience of the user. Fifth and finally, the software is available in R code, meaning that it is free to use, and anyone can read, learn from, revise, and extend the program code.

Method

We developed software to compute and bootstrap the regression metrics reviewed in this article, using R as our underlying platform. R is a “cutting-edge, free, open source statistical package” that runs on all commonly used operating systems (see R Development Core Team, 2013) and is gaining popularity across research disciplines (Culpepper & Aguinis, 2011). One major benefit of R platform is the opportunity for researchers to develop and update programs or “packages” to the

R repository that extend the functions of the base system. To date, the R platform is supported by 4,461 user-contributed packages.

Extending the work of one of the user-contributed packages, `yhat` (Nimon & Roberts, 2012), the current article presents software that (a) calculates all of the regression results discussed in this article (`calc.yhat`), (b) bootstraps the results (`boot.yhat`), (c) evaluates the bootstrapped results (`booteval.yhat`), and (d) plots relevant CIs (`plotCI.yhat`). These extensions are reviewed in the sections that follow. We believe they represent a meaningful update to the `yhat` package.

`calc.yhat`. The `calc.yhat` function produces four sets of regression metrics. The first set is called `PredictorMetrics` and contains the predictor metrics reviewed in this article: b weights, beta weights, validity coefficients, structure coefficients, squared structure coefficients, unique coefficients, the sum of common coefficients, conditional dominance weights, general dominance weights, product measures, and relative weights.

The second set is called `OrderedPredictorMetrics`, and for each metric simply ranks the absolute values of the values contained in the `PredictorMetrics` set.

The third set is called `PairedDominanceMetrics` and contains the complete, conditional, and general dominance values (D_{ij}) for each pair of predictors. As in Azen and Budescu (2003), a value of 1 in D_{ij} indicates that X_i dominates X_j , 0 indicates that X_j dominates X_i , and .5 indicates that dominance cannot be established between X_i and X_j .

The fourth set is called `APSRRelatedMetrics` and contains commonality coefficients, which is the criterion variance partitioned across all subsets of independent variables. Also included are the multiple R^2 s resulting from an APS regression and the incremental variance for each predictor (the latter of which is used with uniqueness coefficients to determine general, conditional, and complete dominance).

`boot.yhat`. The `boot.yhat` function is based on the R package `boot` (Canty & Ripley, 2011) and supports the bootstrapping of all the metrics computed by `calc.yhat`. This function also computes a Kendall's tau correlation for each type of predictor metric (e.g., r , β , r_s), correlating bootstrap metrics with sample metrics to indicate stability of the ranking of each type of predictor metric, where high correlations imply agreement in the rank ordering. This feature would most likely be used to correlate the rank order of predictor metrics between like sample and bootstrap statistics. However, it is also possible for the user to pass a set of statistics such that each set of bootstrap statistics is correlated to the same set of sample statistics, as in LeBreton, Ployhart, and Ladd (2004), where the rank order of predictors from general dominance analysis was correlated with the rank order of predictors generated from alternative methods (e.g., product measure, r^2 , relative weights analysis).

`booteval.yhat`. The `booteval.yhat` function takes the `boot.yhat` output generated across bootstrap iterations and calculates descriptive statistics and CIs based on them. Bootstrapped CIs avoid the need to rely on multivariate normality assumptions in the data or in the statistics generated from them. The accelerated bootstrap interval ("bca") is the default CI. The basic bootstrap interval ("basic"), the accelerated bootstrap interval ("bca"), the first order normal approximation ("norm"), and the bootstrap percentile interval ("perc") are also supported.

Descriptive Statistics

Regarding descriptive statistics, `booteval.yhat` produces `tauDS`, which contains the M and SD of Kendall's tau (described earlier) across bootstrap iterations from `boot.yhat`. These results

indicate how consistent the observed predictor ordering is between each bootstrapped sample and the actual sample (high mean and low variance indicates a consistent ordering). Also, for each pair of predictors, the M and SE of D_{ij} for each type of dominance is reported alongside the proportion of $D_{ij} = 1$ ($P_{i,j}$), $D_{ij} = 0$ ($P_{\bar{i},j}$), and $D_{ij} = .5$ (P_{noij}) across bootstrap replications, as well as the reproducibility of D_{ij} ($Reprod$). These values are in keeping with Azen and Budescu (2003). The SE of

$(D_{ij}) = \sqrt{\frac{1}{S-1} \sum_{s=1}^S (D_{ij}^s - \bar{D}_{ij})^2}$, where S equals the number of bootstrap samples and the reproducibility of D_{ij} is the “proportion of bootstrap samples that agree with the sample results” (p. 141).

Confidence Intervals

For data in the `PredictorMetrics` and `APSRelatedMetrics` object, CIs are provided alongside and apart from sample statistics in objects, respectively, named `combCIpm`, `lowerCIpm`, `upperCIpm` and `combCIaps`, `lowerCIaps`, `upperCIaps`. The data in these objects allow researchers to make confidence statements regarding the associated metrics and to identify whether pairs of metrics differ to a statistically significant degree (i.e., the related CIs do not overlap).

However, when comparing the metric of one variable to another, overlapping CIs do not necessarily indicate a statistically nonsignificant difference between parameter estimates (see Cumming & Finch, 2001; Zientek, Yetkiner, & Thompson, 2010). One has to examine the distribution of differences between the two bootstrapped estimates of interest across replications. To that end, `booteval.yhat` also provides CIs around differences between select pairs of metrics. Confidence intervals around differences between pairs of predictors for each predictor metric are reported with sample statistics in `combCIpmDiff` and separately in `lowerCIpmDiff` and `upperCIpmDiff`. For each order of predictor combinations (e.g., 1st order, 2nd order, ... k th order, where k = number of predictors), CIs around pairs of APS and commonality coefficients are reported with sample statistics in `combCIapsDiff` and separately in `lowerCIapsDiff` and `upperCIapsDiff`. Data in these objects allow researchers to determine whether the R^2 s or the commonality coefficients associated with a particular number of predictors are statistically significantly different from one another, which may be useful for identifying equivalent fitting models for a particular number of predictors (cf. Thompson, 2006) and for statistically evaluating the difference between joint variance components (cf. Schoen, DeSimone, & James, 2011). For each pair of predictors, CIs around differences between incremental variance are reported with sample statistics in `combCIincDiff` and separately in `lowerCIincDiff` and `upperCIincDiff`. Data in these objects provide insight into the determination of complete dominance across bootstrapped samples.

`plotCI.yhat`. The `plotCI.yhat` function plots the sample statistics and the upper and lower CI for associated objects that are passed to it, such as the sample statistics in `PredictorMetrics`. The function could also be used to plot the CI around sample statistics in `APSRelatedMetrics` when the number of predictors is small.

Other Functions

Several other new functions were also written to support these main functions, including but not limited to functions to conduct commonality analysis, dominance analysis, and relative weights analysis. Although the `yhat` package (Nimon & Roberts, 2012) supports the calculation of commonality coefficients, the associated function (`commonalityCoefficients`) performs an APS

regression internally and does not provide its results, whereas the current program produces APS results directly. Furthermore, because the current program computes and reports results for both commonality analysis and dominance analysis, and because both analyses are based upon an APS regression, the current software conducts APS regression only once, thus enhancing the efficiency and comprehensiveness of the current solution.

Illustrative Example 1

To first illustrate the use of `calc.yhat`, `boot.yhat`, `booteval.yhat`, and `plotCI.yhat`, we conducted a secondary data analysis on the four-predictor regression example discussed by Azen and Budescu (2003). We selected their primary four-predictor model as an example since dominance analysis did “not reveal the complex suppression effects” (Beckstead, 2012, p. 243), and thus we were interested in seeing what the other regression metrics discussed in this article might reveal about the regression model. Replicating the example was accomplished in six steps (see the Appendix for details).

Results for Illustrative Example 1

calc.yhat

In reviewing the results from `calc.yhat`, we first note similarities and differences in the `PredictorMetrics` and `OrderedPredictorMetrics` presented in Figure 1. In sum, the `bs`, `betas`, uniqueness coefficients, and product measures identify the predictor order as X_1 , X_2 , X_3 , and X_4 , whereas the validity coefficients (and their derivatives), general dominance weights, and relative weights identify the predictor order as X_1 , X_4 , X_3 , and X_2 , thus swapping X_4 and X_2 . Conditional dominance weights also suggest a predictor order of X_1 , X_4 , X_3 , and X_2 , except the conditional dominance weights associated with two predictors (i.e., `CD:2`) and the last set of conditional dominance weights (i.e., `CD:3`) that are mathematically identical to uniqueness coefficients and thus share the same results.

The `bs` and `betas` values and ranks are identical, as expected with a data set whose variables have been standardized to have M_s of 0 and SD_s of 1. The uniqueness coefficients and the conditional dominance weights for $k = 3$ (i.e., `CD:3`) are always identical, because both identify the amount of predictive variance that is unique to a predictor. Also note that the order of the `b` and `betas` agrees with the order of the uniqueness coefficients, and the order of the validity (r), structure (r_S), squared structure coefficients (r_S^2), and squared validity (`CD:0`) are identical. The former agreement is in line with Nunnally and Bernstein (1994), who noted that unique coefficients and `b` weights generate the same p values. The latter agreement is aligned with Courville and Thompson (2001), who point out that structure coefficients are rescaled validity coefficients. The similarity between the general dominance (`GenDom`) and relative weights (`RLW`) is in keeping with research that indicates that these coefficients typically yield almost identical values, despite different definitions and computational strategies (see Johnson, 2000; LeBreton et al., 2004).

The Pratt weights diverge from the general dominance and relative weights, even though they also sum to equal the R^2 . This divergence likely stems from a disagreement in the signs of X_2 's beta and validity coefficient, which is suggestive of a suppression effect (cf. Thompson, 2006). However, the conditional dominance weights do not highlight X_2 as a suppressor because the conditional dominance weights for X_2 (i.e., `CD:0`, `CD:1`, `CD:2`, `CD:3`) do not increase as subset models become more complex (cf. Azen & Budescu, 2003; Beckstead, 2012).

The D_{ij} values in the `PairedDominanceMetrics` set reflect some of the information revealed by the `PredictorMetrics` set. The D_{ij} values for general dominance (`Gen`) are reflective of the order of the general dominance weights (i.e., $X_1 > X_4 > X_3 > X_2$). The D_{ij} values for conditional

```

regrOut
$PredictorMetrics
      b  Beta  r   rs  rs2 Unique Common CD:0  CD:1  CD:2  CD:3 GenDom
X1    0.905 0.905 0.6 0.785 0.616 0.246 0.114 0.36 0.300 0.263 0.246 0.292
X2   -0.466 -0.466 0.3 0.392 0.154 0.071 0.019 0.09 0.074 0.069 0.071 0.076
X3    0.291 0.291 0.4 0.523 0.274 0.061 0.099 0.16 0.095 0.063 0.061 0.095
X4    0.129 0.129 0.5 0.654 0.428 0.010 0.240 0.25 0.152 0.073 0.010 0.121
Total   NA    NA  NA    NA 1.472 0.388 0.472 0.86 0.621 0.468 0.388 0.584

      Pratt  RLW
X1    0.543 0.291
X2   -0.140 0.076
X3    0.117 0.098
X4    0.065 0.120
Total  0.585 0.585

$OrderedPredictorMetric
      b Beta r rs rs2 Unique Common CD:0  CD:1  CD:2  CD:3 GenDom Pratt  RLW
X1 1 1 1 1 1 1 2 1 1 1 1 1 1 1 1
X2 2 2 4 4 4 2 4 4 4 3 2 4 2 4
X3 3 3 3 3 3 3 3 3 3 4 3 3 3 3
X4 4 4 2 2 2 4 1 2 2 2 4 2 4 2

$PairedDominanceMetrics
      Comp Cond Gen
X1>X2 1.0 1.0 1
X1>X3 1.0 1.0 1
X1>X4 1.0 1.0 1
X2>X3 0.5 0.5 0
X2>X4 0.5 0.5 0
X3>X4 0.5 0.5 0

$APSRRelatedMetrics
      Commonality  % Total  R2 X1.Inc X2.Inc X3.Inc X4.Inc
X1          0.246    0.421 0.360      NA  0.090  0.117  0.113
X2          0.071    0.121 0.090  0.360      NA  0.138  0.223
X3          0.061    0.105 0.160  0.317  0.068      NA  0.120
X4          0.010    0.018 0.250  0.223  0.063  0.030      NA
X1,X2       -0.013   -0.022 0.450      NA      NA  0.124  0.073
X1,X3       -0.036   -0.062 0.477      NA  0.097      NA  0.037
X2,X3       -0.020   -0.035 0.228  0.346      NA      NA  0.110
X1,X4        0.100    0.170 0.473      NA  0.051  0.041      NA
X2,X4        0.026    0.045 0.313  0.210      NA  0.025      NA
X3,X4        0.063    0.107 0.280  0.233  0.058      NA      NA
X1,X2,X3     0.025    0.043 0.574      NA      NA      NA  0.010
X1,X2,X4    -0.016   -0.028 0.523      NA      NA  0.061      NA
X1,X3,X4     0.050    0.086 0.513      NA  0.071      NA      NA
X2,X3,X4     0.013    0.023 0.338  0.246      NA      NA      NA
X1,X2,X3,X4 0.004    0.006 0.584      NA      NA      NA      NA
Total        0.584    1.000      NA      NA      NA      NA      NA
    
```

Figure 1. Output from calc.yhat for illustrative example.

dominance (Cond) indicate that across regression models of different subset sizes (i.e., k), X_1 conditionally dominates the other predictors (i.e., $X_1 > X_2, X_3, X_4$) and contributes more incremental variance on average to models of different subsets than the other predictors. The D_{ij} values for complete dominance reflect information not previously presented and indicate that X_1 completely dominates all other predictors (i.e., $X_1 > X_2, X_3, X_4$).

In reviewing the APSRRelatedMetrics set, note that the uniqueness coefficients and squared validity coefficients are redundant with what is reported in the PredictorMetrics set. This is done for ease of interpretation in either set. The commonality coefficients (Commonality) in the

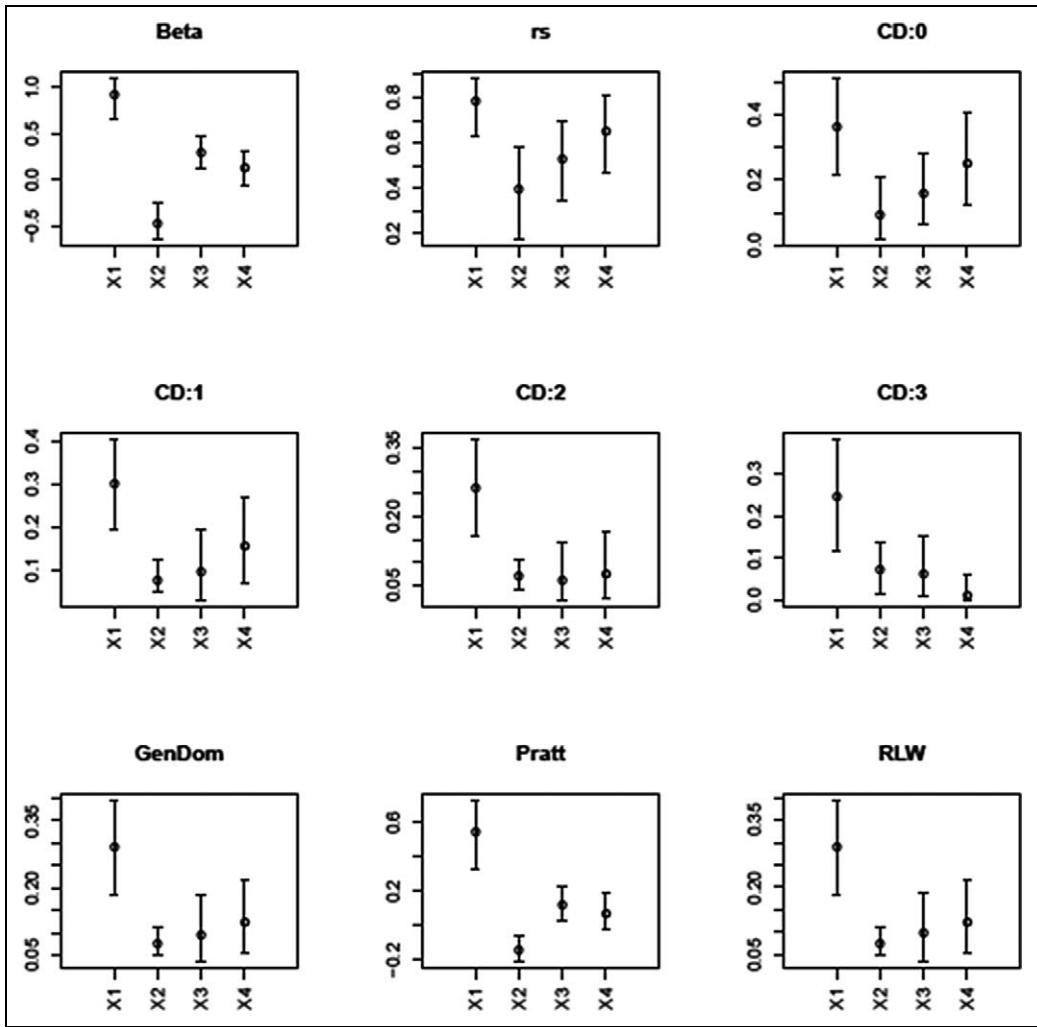


Figure 2. Output from plot.yhat for select predictor metrics from illustrative example.

APSRelatedMetrics set confirm suppression in the regression model. In particular, they identify suppression involving the following subsets of independent variables: X_1 and X_2 , X_1 and X_3 , X_2 and X_3 , and X_1, X_2 , and X_4 . The APSRelatedMetrics set contains the results of the APS regression and shows, for example, that there is little difference in the multiple R^2 produced from the regression model with variables X_1, X_2 , and X_3 ($R^2 = .523$) and the regression model with variables X_1, X_3 , and X_4 ($R^2 = .513$). Finally, the APSRelatedMetrics identify the amount of variance each predictor adds to each subset model, which can be used to understand the determination of complete dominance. One sees, for example, that X_4 does not completely dominate X_2 because X_2 (i.e., .097) adds less variance to the regression model with X_1 and X_3 than does X_4 (i.e., .037).

booteval.yhat and plotCI.yhat

Figure 2 presents the bootstrapped CIs around select coefficients from the PredictorMetrics set. Note that with few exceptions (e.g., betas for X_1 and X_2 , product measures for X_1 and X_2 , relative

```

combCipmDiff[,c("Beta", "rs", "CD:0", "CD:1", "CD:2", "CD:3", "GenDom", "Pratt", "RLW")]

```

	Beta	rs	CD:0
X1-X2	1.371 (0.925, 1.713) *	0.393 (0.235, 0.526) *	0.270 (0.151, 0.372) *
X1-X3	0.614 (0.333, 0.831) *	0.262 (0.000, 0.492) *	0.200 (0.000, 0.377) *
X1-X4	0.776 (0.400, 1.100) *	0.131 (-0.099, 0.336)	0.110 (-0.084, 0.283)
X2-X3	-0.757 (-1.018, -0.460) *	-0.131 (-0.435, 0.159)	-0.070 (-0.220, 0.091)
X2-X4	-0.595 (-0.805, -0.355) *	-0.262 (-0.530, -0.009) *	-0.160 (-0.326, -0.005) *
X3-X4	0.162 (-0.171, 0.490)	-0.131 (-0.380, 0.113)	-0.090 (-0.262, 0.079)
	CD:1	CD:2	CD:3
X1-X2	0.226 (0.130, 0.312) *	0.194 (0.110, 0.276) *	0.175 (0.088, 0.261) *
X1-X3	0.205 (0.032, 0.338) *	0.200 (0.043, 0.326) *	0.185 (0.035, 0.315) *
X1-X4	0.148 (-0.019, 0.299)	0.190 (0.035, 0.328) *	0.236 (0.075, 0.374) *
X2-X3	-0.021 (-0.133, 0.068)	0.006 (-0.084, 0.068)	0.010 (-0.088, 0.086)
X2-X4	-0.078 (-0.197, 0.013)	-0.004 (-0.098, 0.064)	0.061 (-0.028, 0.129)
X3-X4	-0.057 (-0.189, 0.080)	-0.010 (-0.116, 0.105)	0.051 (-0.043, 0.143)
	GenDom	Pratt	RLW
X1-X2	0.216 (0.125, 0.298) *	0.683 (0.410, 0.925) *	0.215 (0.123, 0.297) *
X1-X3	0.197 (0.040, 0.322) *	0.426 (0.176, 0.626) *	0.193 (0.033, 0.319) *
X1-X4	0.171 (0.013, 0.310) *	0.478 (0.202, 0.698) *	0.171 (0.010, 0.317) *
X2-X3	-0.019 (-0.122, 0.057)	-0.257 (-0.372, -0.141) *	-0.022 (-0.126, 0.056)
X2-X4	-0.045 (-0.149, 0.030)	-0.205 (-0.327, -0.088) *	-0.044 (-0.149, 0.035)
X3-X4	-0.026 (-0.148, 0.097)	0.052 (-0.134, 0.236)	-0.022 (-0.148, 0.105)

Figure 3. Select predictor metric differences from booteval.yhat for illustrative example.

weights for X_1 and X_2), it is difficult to determine visually whether or not a pair of predictor metrics differs by a statistically significant amount. However, differences between predictor coefficients can easily be seen in Figure 3. The CIs that are indicated with an asterisk (*) identify differences between pairs of predictor coefficients that are statistically significantly different from zero. One sees that across metrics, X_1 tends to produce coefficients that are statistically significantly greater than the remaining predictors. The exception to this rule involves X_4 where its structure coefficient (rs), squared validity coefficient ($CD:0$), conditional dominance weight for $k = 1$ ($CD:1$) are statistically equivalent to X_1 's. Also note that with the exception of X_3 and X_4 , all betas are statistically significant from one another and that the structure coefficient, squared validity coefficient, and product measure ($Pratt$) for X_2 are statistically significantly different from X_4 's.

Figure 4 presents the descriptive statistics of the bootstrapped Kendall's tau correlation between the sample predictor metrics and the bootstrap statistics of like metrics. Across metrics, the order of b and betas ($Ms = .948, .946$) was most reproducible across bootstrapped samples. Note that the variance of the correlations across bootstraps is lower as the mean of the correlations is higher (in fact, the correlation between the mean and variance across metrics was $-.98$). Figure 4 also presents the sample D_{ij} values along with their Ms , SEs , and probabilities and reproducibility over the 1,000 bootstrap samples. It is interesting to note that X_1 completely dominated X_2 in each of the bootstrapped samples. Given the hierarchical nature of dominance analyses, X_1 also conditionally and generally dominated X_2 in each of the bootstrapped samples.

Figure 5 presents the bootstrapped CIs around the coefficients in the `APSRelatedMetrics` set. Note that the CIs around the uniqueness coefficients and the squared validity coefficients found in Figure 3 are also presented in Figure 5, for ease of interpretation. With only a few nonoverlapping CIs, it is difficult to visually identify statistically significant differences among pairs of coefficients. However, statistically significant differences between APS-related coefficients can readily be seen in Figures 6 and 7, indicated by an asterisk next to the CIs for pairs of APS-related coefficients that are statistically significantly different from zero. One sees, for example, that a majority of the joint variance components involving two predictors are statistically significant different from one another


```

result$tauDS
      b  Beta    r    rs   rs2 Unique Common  CD:0  CD:1  CD:2  CD:3 GenDom Pratt  RLW
Mean 0.948 0.946 0.835 0.835 0.834 0.775 0.776 0.836 0.784 0.559 0.775 0.717 0.905 0.706
SD   0.121 0.123 0.202 0.202 0.202 0.236 0.206 0.202 0.218 0.377 0.236 0.257 0.150 0.257

result$domBoot
      Dij Mean  SE  Pij  Pji Pijno  Reprod
Com_X1>X2 1.0 1.000 0.000 1.000 0.000 0.000 1.000
Com_X1>X3 1.0 0.982 0.092 0.965 0.000 0.035 0.965
Com_X1>X4 1.0 0.916 0.187 0.831 0.000 0.169 0.831
Com_X2>X3 0.5 0.470 0.230 0.077 0.138 0.785 0.785
Com_X2>X4 0.5 0.458 0.151 0.007 0.091 0.902 0.902
Com_X3>X4 0.5 0.486 0.255 0.117 0.144 0.739 0.739
Con_X1>X2 1.0 1.000 0.000 1.000 0.000 0.000 1.000
Con_X1>X3 1.0 0.984 0.096 0.971 0.003 0.026 0.971
Con_X1>X4 1.0 0.932 0.171 0.864 0.000 0.136 0.864
Con_X2>X3 0.5 0.390 0.310 0.107 0.326 0.567 0.567
Con_X2>X4 0.5 0.460 0.162 0.016 0.095 0.889 0.889
Con_X3>X4 0.5 0.496 0.265 0.137 0.144 0.719 0.719
Gen_X1>X2 1.0 1.000 0.000 1.000 0.000 0.000 1.000
Gen_X1>X3 1.0 0.990 0.100 0.990 0.010 0.000 0.990
Gen_X1>X4 1.0 0.983 0.129 0.983 0.017 0.000 0.983
Gen_X2>X3 0.0 0.335 0.472 0.335 0.665 0.000 0.665
Gen_X2>X4 0.0 0.159 0.366 0.159 0.841 0.000 0.841
Gen_X3>X4 0.0 0.330 0.470 0.330 0.670 0.000 0.670
    
```

Figure 4. Descriptive statistics output from booteval.yhat for illustrative example.

(e.g., X_1, X_3 vs. X_1, X_4 ; X_2, X_3 vs. X_3, X_4). One also sees out of the 15 comparisons involving regression models with $k = 2$, 7 contained statistically significant differences in multiple R^2 s; for the 6 comparisons involving regression models with $k = 3$, 3 contained statistically significant differences in multiple R^2 s. One can also see that X_1 did not completely dominate X_4 across bootstrapped samples because of statistically nonsignificant differences in incremental variance in the cases where $k = 0$ and where $k = 1$.

Illustrative Example 2

We also conducted a secondary data analysis on the correlation matrix reported in Podsakoff, Ahearne, and MacKenzie (1997) to provide an illustrative example of how one might write up the results from the software presented in this article to be suitable for publication. We examined the model that regressed work crew ($n = 40$) quantity on “crew members’ assessments of their crews’ helping behavior, civic virtue, and sportsmanship . . . aggregated at the work group level” (p. 265). We selected the model as it has been previously identified in the literature (e.g., Courville & Thompson, 2001; Nimon & Reio, 2011) as a model that benefitted from examining multiple metrics. Although prior literature has examined the regression model using bs, betas, r_s s, and commonality coefficients, it does not appear that the model has been examined using other metrics presented in this article, including dominance analysis or relative weights. To analyze their model, we modified the example code previously presented to accommodate their correlation matrix, which we present in Table 3.

Results for Illustrative Example 2

The model that regressed aggregated *civic virtue*, *sportsmanship*, and *helping behavior* on *work crew quality* was statistically and practically significant, $F(3, 36) = 3.886, p = .017, R^2 = .247$. The aggregated organizational citizenship behaviors explained $\sim 25\%$ of the variance in work crew

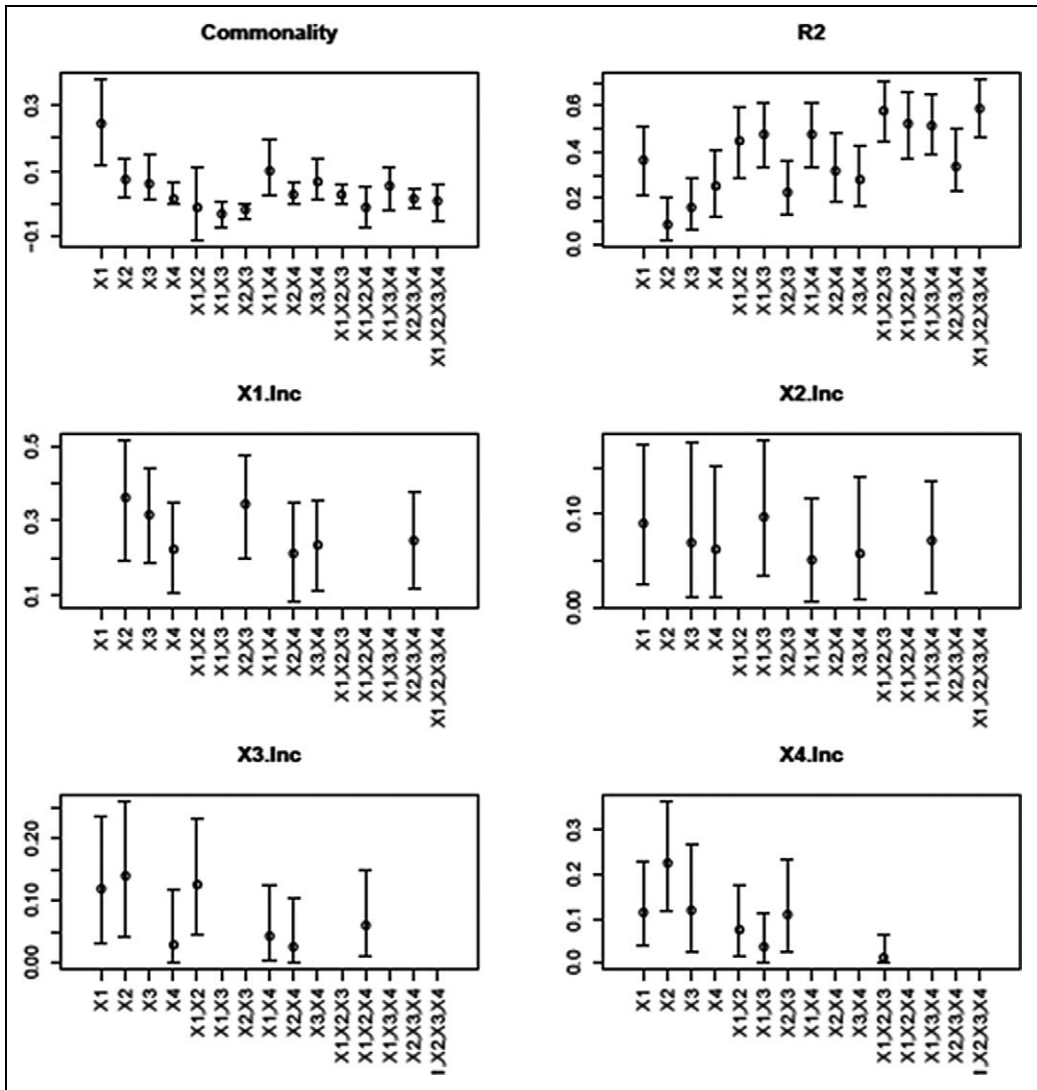


Figure 5. Output from plot.yhat for all-possible-subsets (APS)–related metrics from illustrative example.

quality. Tables 4 and 5, respectively, present the predictor and APS-related metrics, including 95% accelerated bootstrap confidence intervals that were produced over 1,000 iterations.

With the exception of betas that identify helping behavior as the most important predictor, followed by sportsmanship and civic virtue, the remaining predictor metrics identify sportsmanship as the most important predictor and helping behavior as the second most important predictor. This means that although sportsmanship (a) shares the most variance with the work crew quality and predicted work crew quality, (b) contributes the most unique and common variance to work crew quality, (c) adds the most incremental variance, on average, to models of different subsizes, and (d) accounts for the largest partition of R^2 as computed with general dominance weights, Pratt measures, and relative weights, helping behavior is given the greatest credit in the regression equation. The predictor metrics also identify civic virtue as a suppressor variable. In addition to it contributing

	Commonality	R2
X1-X2	0.175 (0.088, 0.261) *	0.270 (0.151, 0.372) *
X1-X3	0.185 (0.035, 0.315) *	0.200 (0.000, 0.377) *
X1-X4	0.236 (0.075, 0.374) *	0.110 (-0.084, 0.283)
X2-X3	0.010 (-0.088, 0.086)	-0.070 (-0.220, 0.091)
X2-X4	0.061 (-0.028, 0.129)	-0.160 (-0.326, -0.005) *
X3-X4	0.051 (-0.043, 0.143)	-0.090 (-0.262, 0.079)
X1, X2-X1, X3	0.023 (-0.091, 0.147)	-0.027 (-0.165, 0.089)
X1, X2-X2, X3	0.007 (-0.095, 0.127)	0.222 (0.023, 0.397) *
X1, X2-X1, X4	-0.113 (-0.257, 0.038)	-0.023 (-0.150, 0.077)
X1, X2-X2, X4	-0.039 (-0.155, 0.092)	0.137 (-0.067, 0.309)
X1, X2-X3, X4	-0.076 (-0.195, 0.050)	0.170 (-0.029, 0.341)
X1, X3-X2, X3	-0.016 (-0.043, 0.013)	0.249 (0.149, 0.333) *
X1, X3-X1, X4	-0.136 (-0.231, -0.043) *	0.004 (-0.101, 0.114)
X1, X3-X2, X4	-0.062 (-0.109, -0.009) *	0.164 (0.014, 0.301) *
X1, X3-X3, X4	-0.099 (-0.187, -0.020) *	0.197 (0.030, 0.334) *
X2, X3-X1, X4	-0.120 (-0.217, -0.037) *	-0.245 (-0.384, -0.084) *
X2, X3-X2, X4	-0.046 (-0.091, -0.009) *	-0.085 (-0.230, 0.062)
X2, X3-X3, X4	-0.083 (-0.159, -0.026) *	-0.052 (-0.197, 0.070)
X1, X4-X2, X4	0.074 (0.018, 0.144) *	0.160 (0.071, 0.255) *
X1, X4-X3, X4	0.037 (-0.054, 0.143)	0.193 (0.049, 0.314) *
X2, X4-X3, X4	-0.037 (-0.105, 0.019)	0.033 (-0.065, 0.125)
X1, X2, X3-X1, X2, X4	0.041 (-0.044, 0.108)	0.051 (-0.043, 0.143)
X1, X2, X3-X1, X3, X4	-0.025 (-0.100, 0.040)	0.061 (-0.028, 0.130)
X1, X2, X3-X2, X3, X4	0.012 (-0.038, 0.052)	0.236 (0.075, 0.375) *
X1, X2, X4-X1, X3, X4	-0.066 (-0.145, 0.036)	0.010 (-0.088, 0.086)
X1, X2, X4-X2, X3, X4	-0.029 (-0.097, 0.052)	0.185 (0.036, 0.315) *
X1, X3, X4-X2, X3, X4	0.037 (-0.007, 0.079)	0.175 (0.087, 0.261) *

Figure 6. Commonality coefficient and R² differences from booteval.yhat for illustrative example.

	X1.Inc-X2.Inc	X1.Inc-X3.Inc	X1.Inc-X4.Inc
.	0.270 (0.151, 0.372) *	0.200 (0.000, 0.377) *	0.110 (-0.084, 0.283)
X1	NA	NA	NA
X2	NA	0.222 (0.022, 0.397) *	0.137 (-0.067, 0.309)
X3	0.249 (0.149, 0.333) *	NA	0.197 (0.030, 0.333) *
X4	0.160 (0.071, 0.255) *	0.193 (0.049, 0.315) *	NA
X1, X2	NA	NA	NA
X1, X3	NA	NA	NA
X2, X3	NA	NA	0.236 (0.075, 0.375) *
X1, X4	NA	NA	NA
X2, X4	NA	0.185 (0.035, 0.315) *	NA
X3, X4	0.175 (0.088, 0.261) *	NA	NA
	X2.Inc-X3.Inc	X2.Inc-X4.Inc	X3.Inc-X4.Inc
.	-0.070 (-0.220, 0.091)	-0.160 (-0.326, -0.005) *	-0.090 (-0.262, 0.079)
X1	-0.027 (-0.165, 0.089)	-0.023 (-0.150, 0.077)	0.004 (-0.101, 0.114)
X2	NA	NA	-0.085 (-0.229, 0.062)
X3	NA	-0.052 (-0.197, 0.070)	NA
X4	0.033 (-0.065, 0.126)	NA	NA
X1, X2	NA	NA	0.051 (-0.043, 0.143)
X1, X3	NA	0.060 (-0.028, 0.130)	NA
X2, X3	NA	NA	NA
X1, X4	0.010 (-0.088, 0.087)	NA	NA
X2, X4	NA	NA	NA
X3, X4	NA	NA	NA

Figure 7. Incremental predictor variance differences from booteval.yhat for illustrative example.

Table 3. Correlation Matrix for Example 2.

	Helping	Civic Virtue	Sportsmanship
Civic virtue	.69		
Sportsmanship	.46	.54	
Quantity	.36	.17	.40

Table 4. Predictor Metrics for Example 2.

Metric	Civic Virtue	Sportsmanship	Helping Behavior
Beta	-.314 (-.785, .087)	.386 (-.011, .713)	.399 (.023, .826)
<i>r</i>	.170 (-.092, .480)	.400 (.074, .639)	.360 (.035, .605)
<i>r_s</i>	.344 (-.280, .803)	.809 (.247, .982)	.728 (.172, .982)
<i>r_s²</i>	.118 (.000, .605)	.654 (.061, .964)	.530 (.045, .969)
Unique	.045 (.000, .235)	.103 (.001, .353)	.082 (.003, .331)
Common	-.016 (-.225, .170)	.057 (-.064, .249)	.048 (-.079, .280)
CD:0	.029 (.000, .226)	.160 (.006, .408)	.130 (.003, .367)
CD:1	.007 (.000, .034)	.102 (.004, .332)	.076 (.003, .264)
CD:2	.045 (.000, .235)	.103 (.001, .353)	.082 (.003, .331)
GenDom	.027 (.002, .059)	.122 (.006, .336)	.096 (.006, .287)
Pratt	-.053 (-.242, .003)	.154 (-.001, .419)	.144 (.003, .447)
RLW	.024 (.002, .051)	.124 (.007, .332)	.097 (.006, .278)

Note: Unique = uniqueness coefficient; common = r^2 - uniqueness; CD = conditional dominance weights; GenDom = general dominance weights; Pratt = Pratt measure; RLW = relative weights.

Table 5. All-Possible-Subset-Related Metrics for Example 2.

Subset	Commonality	R^2	Civ.Inc	Sprt.Inc	Help.Inc
Civ	.045 (.000, .235)	.029 (.000, .226)	NA	.134 (.002, .389)	.112 (.003, .350)
Sprt	.103 (.001, .353)	.160 (.006, .408)	.003 (0, .030)	NA	.039 (.000, .232)
Help	.082 (.003, .331)	.130 (.003, .367)	.012 (0, .112)	.070 (.000, .301)	NA
CivSprt	-.034 (-.166, .005)	.163 (.009, .388)	NA	NA	.082 (.003, .331)
CivHelp	-.042 (-.208, .016)	.141 (.007, .361)	NA	.103 (.001, .353)	NA
SprtHelp	.031 (-.011, .173)	.199 (.013, .424)	.045 (0, .235)	NA	NA
CivSprtHelp	.060 (-.007, .225)	.245 (.024, .435)	NA	NA	NA

Note: Civ = civic virtue; Sprt = sportsmanship; Help = helping behavior.

more unique variance to the regression effect than it has in common with the work crew quality and yielding a negative Pratt's measure (as a result of inconsistent signs between its beta and validity coefficient), the conditional dominance weights for civic virtue do not decrease monotonically with more complex models (cf. Azen & Budescu, 2003).

The APS-related metrics show that civic virtue contributes the most incremental variance when added to a regression model that contains sportsmanship and helping behavior and that its addition suppresses irrelevant variance in sportsmanship and helping behavior, making them better predictors than if civic virtue was not included. This means that if sportsmanship and helping behavior are to have maximum impact in predicting work crew quality, the measures should be refined to eliminate irrelevant variance related to civic virtue. Analysis of the bivariate correlations and the incremental validity coefficients reported in Table 4 indicates that

Table 6. Descriptive Statistics for Kendall's Tau Across Bootstrap Iterations.

Metric	<i>M</i>	<i>SD</i>
Beta	.641	.367
<i>r</i>	.668	.415
r_s^2	.667	.415
r_s^2	.645	.435
Unique	.353	.540
Common	.554	.521
CD:0	.646	.434
CD:1	.531	.492
CD:2	.353	.540
GenDom	.530	.490
Pratt	.634	.402
RLW	.566	.480

Note: Unique = uniqueness coefficient; common = r^2 - uniqueness; CD = conditional dominance weights; GenDom = general dominance weights; Pratt = Pratt measure; RLW = relative weights.

Table 7. Differences in Predictor Metrics for Example 2.

Metric	Civ-Sprt	Civ-Help	Sprt-Help
Beta	-.700 (-1.367, -.065)*	-.713 (-1.543, -.042)*	-.013 (-.738, .489)
<i>r</i>	-.230 (-.523, .039)	-.190 (-.457, .012)	.040 (-.313, .354)
r_s^2	-.465 (-.977, .050)	-.384 (-.846, -.013)*	.081 (-.601, .710)
r_s^2	-.536 (-.906, -.026)*	-.412 (-.806, -.024)*	.124 (-.715, .813)
Unique	-.058 (-.308, .093)	-.037 (-.246, .056)	.021 (-.275, .298)
Common	-.073 (-.301, .005)	-.064 (-.230, .003)	.009 (-.042, .166)
CD:0	-.131 (-.388, .009)	-.101 (-.284, .003)	.030 (-.213, .288)
CD:1	-.095 (-.340, .030)	-.069 (-.250, .017)	.026 (-.232, .293)
CD:2	-.058 (-.308, .093)	-.037 (-.246, .056)	.021 (-.275, .298)
GenDom	-.095 (-.340, .030)	-.069 (-.249, .016)	.026 (-.231, .294)
Pratt	-.207 (-.557, -.004)*	-.197 (-.626, .001)	.010 (-.415, .330)
RLW	-.100 (-.340, .021)	-.073 (-.245, .014)	.027 (-.219, .288)

Note: Civ = civic virtue; Sprt = sportsmanship; Help = helping behavior. * indicates that the confidence interval does not contain 0.

sportsmanship completely dominates helping behavior, which completely dominates civic virtue. This means that across all possible subset models, (a) sportsmanship adds more incremental variance than helping behavior and civic virtue and (b) helping behavior adds more incremental variance than civic virtue.

It is important to note that many of the study's findings may not be replicable, given its small sample size. Across bootstrapped samples, the predictor order based on betas was most consistent with the sample data. As presented in Table 6, the average Kendall's tau, correlating beta weight-based predictor order from bootstraps to sample data, was .641 ($SD = .367$). Predictor order based on unique variance was least replicable ($M = .353$, $SD = .540$). Bootstrap analysis of differences between predictor metrics found only a few statistically significant differences (see Table 7). Bootstrap analyses of differences between appropriate APS metrics found no statistically significant differences. For each order of predictor combinations, there were no statistically significant differences among the multiple R^2 or commonality coefficient produced, nor were there statistically significant differences among the predictors' incremental validity coefficients.

Discussion

Although the interpretation of linear regression weights is straightforward when the goal is prediction, it has long been known that when the goal of a regression analysis is instead to make some conclusions about the relative importance of the predictors in the model, the intercorrelations between predictors (multicollinearity) undermine the use of regression weights for this purpose. Alternative metrics are required—and perhaps more than one set of metrics is interesting to consider because each is operationalized differently and carries different assumptions. Specific to the current article, we review regression weights, zero-order validity coefficients, structure coefficients, Pratt measures, relative importance weights, all-possible-subsets regression, commonality coefficients, and dominance weights.

Perhaps more important than this review of a wide range of metrics relevant to linear regression, our article offers a freely available software package in R code that (a) computes all of these indices at once, (b) computes associated bootstrapped confidence intervals, (c) compares pairs of predictors against one another in terms of their bootstrapped metric, and (d) performs these contributions for any number of predictors so long as the correlation matrix is positive definite (invertible). Other software is limited in all four respects.

We hope researchers will use this software to change their fundamental approach to conducting and interpreting linear regression analysis as applied to their data. Given the variety of weights available, it can be informative to consider an array of weights and to report the most appropriate importance weights, or to examine how they converge and diverge, rather than merely focus on the weights that are the most popular or typically available. The program we offer obviously requires the expertise of the researcher or practitioner to determine which set or sets of importance weights are most appropriate to report. Fortunately, Nathans et al. (2012) provided an accessible treatment of the metrics reported by the software presented along with strengths, limitations, and recommendations for practice. This along with other works that also address predictor importance in detail (e.g., Budescu & Azen, 2004; LeBreton et al., 2004) and the examples in the current article should provide researchers a general template for their own work in interpreting and reporting MLR models.

Although the regression indices we have reviewed can be informative, and the software can be a useful tool to make use of these indices, there are several avenues for future research that extend beyond the current purview. First, it is possible that the study of some psychological phenomena requires multiple criteria as well as multiple predictors, leading to a complex canonical prediction problem (e.g., Azen & Budescu, 2006; LeBreton & Tonidandel, 2008; Nimon, Henson, & Gates, 2010). Second, another frequent concern is the reliability or stability of importance-weight estimates across independent samples to which a regression model is supposed to generalize (e.g., Azen & Budescu, 2003; Johnson, 2004). We implemented bootstrapping to compute standard errors of the coefficients for all metrics and to address directly the problem of stability in a random sample; however, there is no guarantee that similar results would be obtained in a nonrandom sample, in particular a sample that is supposed to exhibit the same pattern of relationships and variable importance but that is a substantively different sample from the first one (e.g., an Army sample vs. an Air Force sample). Thus, research could examine replication and generalizability of these MLR metrics across samples of varying degrees of generalizability.

Third, we provide information on the submodels from all-possible-subsets regression for further investigation, and future research could quickly apply statistical and graphical exploratory tools to represent and test patterns within APS that go beyond the submodel summaries provided by general and conditional dominance analysis. Fourth, future research could consider MLR models that remain linear in their parameter estimates but include interaction terms and/or nonlinear terms, which would likely raise additional considerations (Dalal & Zickar, 2012). Perhaps many of these metrics could be extended to the hierarchical regression analysis framework; this might be something like APS yet

would impose constraints or a structure on the sets of models to be tested. Fifth and finally, although we have covered a wide range of metrics, we also realize that other metrics could eventually be incorporated into the `yhat` package (e.g., the Lasso method and its variants, Tibshirani, 1996; Bayesian variable selection, Mitchell & Beauchamp, 1988; Bayesian model averaging, Raftery, Madigan, & Hoeting, 1997).

Although all of these suggestions might prove worthy of consideration in future research, it was beyond the scope of this article to address them in detail. Again, our main focus is in providing a comprehensive and freely available program useful for bringing together and generating different types of predictor importance metrics in multiple regression analysis and to provide empirical examples that accompany the program, both of which we hope will allow researchers to think about and conduct regression analysis in a fundamentally different manner. Previously, such metrics were examined in isolation, often without much consideration of the other metrics available. Thus, regular use of the program in the future will hopefully provide researchers with new insights and guidance for the use of regression metrics that nobody—including ourselves—has yet offered.

Consider the work of Seibold and McPhee (1979) who examined the impact that cognition and social affect had in minority women's intent to get a cancer screening test. Had the researchers only considered betas, they would have missed identifying cognition as a suppressor variable and the need to purify cognitive relevance from screening messages aimed at addressing social affect in order to have the maximum impact on intentions. To generalize from this example, we believe and envision that the software described can become an essential tool in substantive research, to understand the predictive relationships and interrelationships among variables in regression models more closely and from different perspectives, as well as in simulation research, to understand and appreciate statistical conditions that cause convergence and divergence among different regression metrics (extending the foundational work of LeBreton et al., 2004). Without conducting such detailed analyses, researchers may miss detecting and interpreting valuable relationships in their data.

Appendix

Steps to Replicate Illustrative Example 1

First, we adapted the R software in Kraha, Turner, Nimon, Zientek, and Henson (2012) to generate a data set (`exdata`) from Azen and Budescu's (2003) correlation matrix reported in Table 3.

```
library(MASS)
library(corpcor)
covm<-c( 1, .6, .3, .4, .5,
        .6, 1, .8, .1, .3,
        .3, .8, 1, .1, .1,
        .4, .1, .1, 1, .5,
        .5, .3, .1, .5, 1)
covm<-matrix(covm,5,5)
varlist<-c("Y", "X1", "X2", "X3", "X4")
dimnames(covm)<-list(varlist,varlist)
exdata<-mvrnorm(n=100,rep(0,5),covm,empirical=TRUE)
exdata<-data.frame(exdata)
```

Second, we applied the regression function `lm` in R to create `lm.out`, an object that contained the primary results of the regression model:

```
library(yhat)
library(miscTools)
lm.out<-lm(Y~X1+X2+X3+X4,data=exdata)
```

Third, we used both `calc.yhat` on `lm.out` and saved the results (e.g., predictor metrics, dominance metrics, all-possible-subsets (APS)–related metrics) in an object named `regrOut`:

```
regrOut<-calc.yhat(lm.out)
```

Fourth, we bootstrapped the results produced from `calc.yhat`, where the boot function operated off of the sample data (`exdata`), the `boot.yhat` function, the number of bootstrap samples (1,000), the regression output from `lm` (`lm.out`), and the output from `calc.yhat` (`regrOut`). Results were saved in an object named `boot.out`.

```
library(boot)
boot.out<- boot(exdata,boot.yhat,1000,lmOut=lm.out, regrout0=regrOut)
```

Fifth, we used the output of `boot` (`boot.out`) and `calc.yhat` (`regrOut`) to create summary statistics of the bootstrap data. We saved these summary data in an object called `result`.

```
result<-booteval.yhat(regrOut,bty="perc",boot.out)
```

Sixth, we reviewed results and created plots of relevant data.

```
library(plotrix)
regrOut
result$tauDS
result$domBoot
plotCI.yhat(regrOut$PredictorMetrics[-
  nrow(regrOut$PredictorMetrics),],result$upperCIpm,result$lowerCIpm,
  pid=which(colnames(regrOut$PredictorMetrics) %in%
  c("Beta","rs","CD:0","CD:1","CD:2","CD:3","GenDom","Pratt","RLW") ==
  TRUE), nr=3,nc=3)
result$combCIpmDiff[,c("Beta","rs","CD:0","CD:1","CD:2","CD:3","GenDom","Prat
  t","RLW")]
plotCI.yhat(regrOut$APSRelatedMetrics[-nrow(regrOut$APSRelatedMetrics),-
  2],result$upperCIaps,result$lowerCIaps,nr=3,nc=2)
result$combCIapsDiff
result$combCIincDiff
```

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