A Generalized Missing-Indicator Approach to Regression with Imputed Covariates

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Abstract

This paper considers estimation of a linear regression model using data where some covariate values are missing but imputations are available to fill-in the missing values. The availability of imputations generates a trade-off between bias and precision in the estimators of the regression parameters: the complete cases are often too few, so precision is lost, but filling-in the missing values with imputations may lead to bias. We provide the new Stata command gmi which allows handling such bias-precision trade-off using either model reduction or model averaging techniques in the context of the generalized missing-indicator approach recently proposed by Dardanoni et al. (2011). If multiple imputations are available, our gmi command can be also combined with the built-in Stata prefix mi estimate to account for the extra variability due to the imputation process. The gmi command is illustrated with an empirical application which investigates the relationship between an objective health indicator and a set of socio-demographic and economic covariates affected by substantial item nonresponse.

Keywords: Missing covariates; Imputation; Bias-precision trade-off; Model reduction; Model averaging.
1 Introduction

When trying to run a regression of interest, researchers often face the problem of missing values on some of the variables. We focus on the case when only the covariates contain missing values and the data are Missing-At-Random (MAR). As argued by Little(1992), the related problem of missing values in the dependent variable is less interesting because if the covariates are complete and the missing values on the dependent variable are MAR, then the incomplete cases contribute no information about the regression parameters of interest.

One approach to this problem—complete-case analysis—is to drop all cases with missing values and run the regression using only the complete cases. Another approach, when imputations are available, is to fill-in the missing values with the imputations and run the regression using all the data, whether observed or imputed. This second approach—which we call the ‘naive’ approach—is actually becoming quite common, as public-use data files increasingly include imputations of key variables affected by missing data problems. Specialized software for carrying out imputations is also becoming increasingly available. One example is the mi suite of commands developed in Stata 11.

From the view point of inference about the regression parameter of interest, the availability of imputations generates a trade-off between bias and precision: the complete cases are often too few, so precision is lost, but filling-in the missing values with the imputations may lead to bias. Dardanoni et al. (2011), henceforth DMP, show that this trade-off is in fact equivalent to that arising in an extended or ‘grand’ regression model that includes two subsets of regressors: the focus regressors corresponding to the observed or imputed covariates, and a set of auxiliary regressors representing all possible interactions between the focus regressors and the missing-data indicators. In the ‘grand’ model, the trade-off is between bias and precision in estimating the coefficients on the focus regressors when we drop subsets of the auxiliary regressors. As discussed in DMP, this second trade-off is easier to deal with than the first, because a variety of methods are available. This paper presents the command gmi that implements several methods corresponding to two alternative strategies for handling such trade-off: model reduction and model averaging.

The remainder of this paper is organized as follows. Section 2 reviews the theoretical background in DMP. Section 3 describes the two alternative strategies for estimating the regression parameters of interest. Section 4 provides a detailed description of our gmi command. Section 5 illustrates the gmi command using data available on the Stata website. Finally, Section 6 use data from the first wave of SHARE (Survey of Health, Ageing and Retirement in Europe) to provide an empirical
application on the relationship between an objective health indicator and a set of socio-demographic and economic covariates affected by substantial item nonresponse.

2 Background

Consider modeling the relationship between an outcome $Y$ and a set of covariates $X$ using data where some covariate values are missing. We assume that, in the absence of missing values, the data would satisfy the classical linear model

$$y = X\beta + u,$$

where $y$ is the $N \times 1$ vector of observations on the outcome of interest, $X$ is an $N \times K$ matrix of observations on the covariates, $\beta$ is the $K \times 1$ vector of regression parameters, and $u$ is an $N \times 1$ vector of regression errors that are homoskedastic, serially uncorrelated and have zero mean conditional on $X$. This means that the full-information OLS estimator from a regression of $y$ on $X$ would be unbiased for $\beta$ and efficient in the Gauss-Markov sense.

We also assume that all missing covariate values can be replaced by imputations. These imputations may be provided by the data-producing agency or may be constructed by the researcher, for example by using the Stata command `mi impute`.

Because the first element of $X$ is taken to be the constant term, which is always observed, the number of possible missing-data patterns is equal to $2^{K-1}$ (no missing data, only the first covariate missing, only the first and the second missing, etc.). A particular data set need not contain all the possible patterns, so we simply index by $j = 0, \ldots, J$ the patterns that are present in the data, with $j = 0$ corresponding to the subsample with complete data, which is assumed to be always available, and $J \leq 2^{K-1} - 1$. To keep track of exactly which covariate values are missing, we introduce the $N \times K$ missing-data indicator matrix $M$, whose $(n,k)$ element is equal to one if the $n$th case has a missing value on the $k$th covariate and is equal to zero otherwise.

We are concerned with the problem of how to combine the observed and the imputed values in order to estimate the regression parameter $\beta$.

2.1 Complete-case analysis

This approach amounts to ignore the imputed values and use only the subsample with complete data. Complete-case analysis is our benchmark because, under two key assumptions, it delivers an unbiased estimator of the regression parameter $\beta$. 
Denoting the subsample with complete data by \([X^0, y^0]\), where \(X^0\) is an \(N_0 \times K\) matrix and \(y^0\) is an \(N_0 \times 1\) vector, the two key assumptions are full rank of \(X^0\) and ignorability of the missing-data process.

**Assumption 1 (Complete-case full rank)** \(X^0\) has full column rank.

**Assumption 2 (Ignorability)** \(M\) and \(y\) are conditionally independent given \(X\).

For Assumption 1 to hold, there must be enough cases (at least \(K\)) with non-missing covariate values. Assumption 2 is weaker than the standard MAR assumption because it only requires mean independence and not independence. Thus, it admits patterns where cases with low or high levels of some covariates systematically have a greater percentage of missing values. This assumption fails if, for example, observations with missing covariate values have a different regression function than observations with no missing values. In this case, an alternative is some type of sample selection model (see for example the Stata command `heckman`). Under these two assumptions, we have the following result, which represents the main justification for complete-case analysis:

**Result 1** If Assumptions 1 and 2 hold, then the complete-case OLS estimator from a regression of \(y^0\) on \(X^0\) is unbiased for \(\beta\).

Although unbiased, the complete-case OLS estimator has the drawback of being much less precise than the full-information OLS estimator, except when the fraction of complete cases is large.

### 2.2 The ‘naive’ and the simple missing-indicator approaches

A common alternative to complete-case analysis is to use all cases and regress \(y\) on the completed design matrix \(W\), whose \((n, k)\) element is equal to the corresponding element of \(X\) if a covariate value is not missing and is equal to the imputed value otherwise. This ‘naive’ approach ignores the fact that the imputations are not the same as the missing covariate values, so it gives an estimator of \(\beta\) that is more precise than the complete-case OLS estimator but is also biased.

Another alternative, the so-called simple missing-indicator approach, consists of regressing \(y\) on the completed design matrix \(W\) and a set of \(J\) dummies \(d_1, \ldots, d_J\), where the elements of \(d_j\) are equal to one for cases that belong to the \(j\)th missing-data pattern and are equal to zero otherwise (the subsample with complete data represents the baseline). Adding dummies for the missing-data patterns increases the flexibility of the model by allowing the intercepts to differ across patterns but, again, unbiasedness is lost (Horton and Kleinman 2007, Jones 1996, Little 1992).
2.3 The generalized missing-indicator approach

The bias arising from the use of imputations may be eliminated by fully interacting the columns of the completed design matrix $W$ with the dummies for the missing-data patterns. DMP call this a generalized missing-indicator approach.

They show that, if $y^j$ and $W^j$ respectively denote the $N_j \times 1$ subvector of $y$ and the $N_j \times K$ submatrix of $W$ corresponding to the $j$th missing-data pattern, then the generalized missing-indicator approach corresponds to using the following ‘grand’ model

$$
\begin{bmatrix}
  y_0 \\
  y_1 \\
  \vdots \\
  y^J
\end{bmatrix}
= \begin{bmatrix}
  X^0 \\
  W^1 \\
  \vdots \\
  W^J
\end{bmatrix} \beta + \begin{bmatrix}
  0 & \cdots & 0 \\
  W^1 & \cdots & W^J \\
  \vdots & \ddots & \vdots \\
  \vdots & & \vdots
\end{bmatrix} \delta + \begin{bmatrix}
  u_0 \\
  v^1 \\
  \vdots \\
  v^J
\end{bmatrix},
$$

where $\beta$ is the regression parameter of interest, the $\delta^j$ are $K \times 1$ vectors of nuisance parameters that may be interpreted as the asymptotic bias in the regression of $y^j$ on $W^j$, and the $v^j$ are $N_j \times 1$ vectors of projection errors that have mean zero and are orthogonal to the columns of $W^j$. A more compact representation of the ‘grand’ model is

$$
y = W \beta + Z \delta + v, \quad (2)
$$

where

$$
W = \begin{bmatrix}
  X^0 \\
  W^1 \\
  \vdots \\
  W^J
\end{bmatrix}, \quad Z = \begin{bmatrix}
  0 & \cdots & 0 \\
  W^1 & \cdots & W^J \\
  \vdots & \ddots & \vdots \\
  \vdots & & \vdots
\end{bmatrix}, \quad \delta = \begin{bmatrix}
  \delta^1 \\
  \vdots \\
  \delta^J
\end{bmatrix}, \quad v = \begin{bmatrix}
  u_0 \\
  v^1 \\
  \vdots \\
  v^J
\end{bmatrix},
$$

respectively an $N \times K$ matrix of observed or imputed covariates, an $N \times JK$ matrix of auxiliary variables, a $JK \times 1$ vector of nuisance parameters, and an $N \times 1$ error vector. The matrix of auxiliary variables $Z$ consists of $JK$ interactions between the set of $J$ dummies $d_1, \ldots, d_J$ for the missing-data patterns and the $K$ columns of the completed design matrix $W$. Notice that this matrix is not required to have full column rank. This occurs when some of the $W^j$ does not have full column rank, either because $N_j < K$ or because $N_j \geq K$ but the columns of $W^j$ are linearly dependent, as when mean imputation or deterministic regression imputation is used. Incidentally, such imputation methods are known to produce data sets with undesirable properties, see e.g. Lundstrom and Sarnndal (2002). When some of the $W^j$ does not have full column rank, only a subset of the coefficients in $\delta^j$ is identifiable but this does not affect the estimates of $\beta$. Also notice that the regression errors in model (2) need not have constant variance because the projections errors $v^1, \ldots, v^J$ may be heteroskedastic.
The main result in DMP is the following:

**Result 2** If Assumption 1 holds then, for any choice of imputations, the OLS estimate of $\beta$ in model (2) is numerically the same as the complete-case OLS estimate of $\beta$.

Thus, if the ignorability assumption holds, regressing $y$ on $W$ and $Z$ allows one to fully exploit the available information and to obtain an unbiased estimator of the regression parameter $\beta$.

3 Alternative strategies for estimating $\beta$

Both the ‘naive’ and the simple missing-indicator approach correspond to using restricted versions of model (2) obtained by placing restrictions on the vector $\delta$. The ‘naive’ approach restricts $\delta$ to be equal to zero, while the simple missing-indicator approach restricts all the $\delta^j$ to be equal to zero except for their first element. When these restrictions are at odds with the data, imposing them leads to an estimator of $\beta$ that is biased but more precise (less variable) than the OLS estimator of $\beta$ in model (2) which, in turn, is numerically the same as the complete-case estimator of $\beta$. This suggest that, by placing restrictions on $\delta$, or equivalently by excluding some of the auxiliary variables in $Z$, one may obtain an estimator of $\beta$ that is better in the mean squared error (MSE) sense than the complete-case estimator. The Stata command in this paper implements two alternative strategies for obtaining such an estimator of $\beta$: model reduction and model averaging.

3.1 Model reduction

Model reduction involves selecting first an intermediate model between the ‘grand’ model (2) and the ‘naive’ model corresponding to $\delta = 0$, and then estimating the parameter of interest $\beta$ conditional on the selected model. Because the variables in the completed design matrix $W$ are treated as focus regressors and are always included, an intermediate model corresponds to one of the $2^{JK}$ possible subsets of auxiliary regressors in $Z$.

Model reduction may be carried out through a number of variable selection methods, such as those implemented by the built-in Stata command `stepwise` or by the `vselect` command discussed in Lindsey and Sheather (2010). Dropping one of the variables in $Z$ amounts to restricting one element of $\delta^j$ to zero. This in turn corresponds to selecting one of the $J$ missing-data patterns and forcing the coefficient on a particular covariate for that data pattern to be the same as for the subsample with complete data. The various methods differ depending on how one explores the set of all the possible models (e.g. via a general-to-specific or via a specific-to-general approach) and
the decision rule used to judge validity of each model considered (e.g. a fixed significance level or an information criterion such as AIC or BIC).

One well known problem with this strategy is pretesting.\footnote{See Magnus (1999) and the FAQ http://www.stata.com/support/faqs/stat/stepwise.html.} Another is the fact that model reduction and estimation are completely separated. As a result, the reported conditional estimates tend to be interpreted as if they were unconditional. A third problem is that, since there are $J$ subsamples with incomplete data and $K$ covariates (including the constant term), the model space may contain up to $2^{JK}$ models. Thus, the model space is huge, unless both $J$ and $K$ are small. Simple model reduction techniques, such as backward and forward selection, analyze at most $JK(JK + 1)/2$ models. More complicated model reduction techniques, such as the leaps and bounds technique implemented in \texttt{vselect}, usually analyze a larger number of models.

### 3.2 Model averaging

Model averaging takes a different route. Instead of selecting a model out of the available set of models, one first estimates the parameter of interest $\beta$ conditional on each model in the model space, and then computes the estimate of $\beta$ as a weighted average of these conditional estimates. When the model space contains $I$ models, a model averaging estimate of $\beta$ is of the form

$$\hat{\beta} = \sum_{i=1}^{I} \lambda_i \hat{\beta}_i,$$

where the $\lambda_i$ are non-negative random weights that add up to one and $\hat{\beta}_i$ is the estimate of $\beta$ obtained by conditioning on the $i$th model. In Bayesian model averaging (BMA), each $\hat{\beta}_i$ is weighted by the posterior probability of the corresponding model. If equal prior probabilities are assigned to each model, then $\lambda_i$ is proportional to the marginal likelihood of $y$ under model $i$. The BMA literature is vast and we refer the reader to Raftery et al. (1997) for a starting point.

Our Stata implementation of standard BMA is based on the \texttt{bma} command provided by De Luca and Magnus (2011). This approach assumes a classical Gaussian linear model for (2), noninformative priors for $\beta$ and the error variance, and a multivariate Gaussian prior for $\delta$. Notice that the computational burden required to obtain a standard BMA estimate is proportional to the dimension $I$ of the model space. In our case $I = 2^{JK}$, so this computational burden is substantial unless both $J$ and $K$ are small.

Another type of BMA is Weighted-Average Least Squares (WALS), introduced by Magnus et al. (2010). WALS also assumes a classical Gaussian linear model for (2) and noninformative priors for $\beta$
and the error variance but, instead of a multivariate Gaussian prior for $\delta$, it uses a distribution with zero mean for the independently and identically distributed elements of the transformed parameter vector $\eta = \eta(\delta)$, whose $h$th element $\eta_h$ is the population $t$-statistic for testing the significance of the $h$th element of $\delta$. Magnus et al. (2010) use the Laplace distribution, while Einmahl et al. (2011) use the Subbotin family which leads to estimators with better asymptotic properties. The assumption that the regression errors in (2) are homoskedastic and serially uncorrelated is not crucial for WALS, and the method can be generalized to non-spherical errors (Magnus et al. 2011).

WALS has three main advantages over standard BMA. First, its computational burden is only proportional to $JK$. Second, its choice of priors corresponds to a more intuitive concept of uncertainty about the role of the auxiliary variables. Third, WALS estimates have bounded risk and are near-optimal in terms of a well-defined regret criterion (Magnus et al. 2010). Our Stata implementation of WALS, for both Laplace and Subbotin priors, is based on the \texttt{wals} command provided by De Luca and Magnus (2011).

### 3.3 Standard errors of the estimators

Like standard Stata estimation commands, we provide estimated coefficients, standard errors and $t$-ratios. We do not provide $p$-values and confidence intervals because our estimators are generally biased and their distribution need not be Gaussian, not even asymptotically. On the other hand, the $h$th regressor may be considered to be robustly correlated with the outcome if the $t$-ratio $\eta_h$ on its coefficient is greater than one in absolute value, in which case the MSE of the unrestricted OLS estimator of the coefficient is lower than that of the restricted OLS estimator (see e.g. Magnus 2002). On the basis of this criterion, we also provide two-standard error bands for the estimated coefficients.

Computation and interpretation of the standard errors differ depending on the estimation strategy (model reduction vs. model averaging) and the general approach to estimation (frequentist vs. Bayesian).

For model reduction, the default is ‘classical’ standard errors of the OLS estimator of the selected model. These standard errors do not take into account heteroskedasticity or serial correlation in the data and, most importantly, ignore the additional sampling variability induced by the model selection step. The option \texttt{bootstrap} gives standard errors based on the wild bootstrap which are valid under conditional heteroskedasticity and also take into account the additional variability due to model selection.
For BMA, the default standard errors have the usual Bayesian interpretation of measuring the spread of the posterior distribution of the parameters of interest given the data and take model uncertainty explicitly into account. In this case, the option `bootstrap` provides a frequentist measure of the variability due to sampling, including the variability due to model selection.

Notice that neither model reduction nor model averaging take into account the additional sampling variability due to imputation. This problem could be addressed by multiple imputation methods (Rubin 1987). As illustrated in Sections 5 and 6, our `gmi` command can be combined with the build-in Stata prefix `mi estimate` (see `mi estimate`).

### 4 Stata command

The new Stata command `gmi` allows handling the trade-off between bias and precision when estimating a classical linear regression model with imputed covariates. The earliest version of Stata required to run this command is version 11.1. The syntax is as follows:

```
gmi depvar [varlist] [if] [in] , imputed(varlist) missing(varlist) [ summarize cc naive smi sw vs bma wals stepwise_options vselect_bma_options wals_options full vce(bootstrap [, bootstrap_options]) auxiliary(string) keep nowarn ]
```

where `depvar` is the dependent variable, `varlist` is an optional list of observed covariates (i.e. covariates whose values are fully observed), `imputed` is the list of imputed covariates (i.e. covariates whose missing values are replaced by imputed values) and `missing` is the relevant list of missing-data indicators (i.e. the non-zero columns of the matrix \(M\) corresponding to the set of imputed covariates). Missing-data indicators take value 0 for observed cases and value 1 for imputed cases. The number of imputed covariates must coincide with the number of missing-data indicators. The first variable in `missing` is paired with the first indicator in `imputed`, the second variable in `missing` is paired with the second indicator in `imputed`, and so on. The constant term (which is always included) plus the set of observed and imputed covariates correspond to the \(K\) columns of the completed design matrix \(W\). The auxiliary regressors in \(Z\) (i.e. the \(JK\) interactions between the \(J\) dummies for the missing-data patterns and the \(K\) columns of \(W\)) are instead automatically generated by the command using the information from `missing`. The `gmi` command shares the same features of all Stata estimation commands, including access to the estimation results. Factor variables, time-series operators and weights are not allowed. A description of the options that are specific to this command is provided in the following sections.
4.1 Options of the gmi command

`summarize`, the default, provides a description of the ‘grand’ model (number of observations, number of observed and imputed covariates, number of focus and auxiliary regressors, number of missing-data patterns, and dimension of the model space), plus summaries of the distribution of `depvar` (number of observations, mean and standard deviation) for the complete-case and each missing-data pattern.

`cc` provides the complete-case estimate of \( \beta \), namely the OLS estimate from a regression of `depvar` on the \( K \) focus regressors in \( W \) using only the complete cases. This is numerically the same as the OLS estimate of \( \beta \) in the grand model (2).

`naive` provides the ‘naive’ estimate of \( \beta \), namely the OLS estimate from a regression of `depvar` on the \( K \) focus regressors in \( W \) using all cases.

`smi` provides the simple missing-indicator estimate of \( \beta \), namely the OLS estimate of \( \beta \) from a regression of `depvar` on the \( K \) focus regressors in \( W \) and the \( J \) dummies for the missing-data patterns using all cases.

`sw` provides the OLS estimate of \( \beta \) from a regression of `depvar` on the \( K \) focus regressors in \( W \) and the subset of auxiliary regressors in \( Z \) selected through the build-in Stata command `stepwise`. This estimate of \( \beta \) is conditional on the selected model. A brief description of the options for the `stepwise` command is given in Section 4.2.

`vs` provides the OLS estimate of \( \beta \) from a regression of `depvar` on the \( K \) focus regressors in \( W \) and the subset of auxiliary regressors in \( Z \) selected through the `vselect` command by Lindsey and Sheather (2010). As for the `sw` option, this estimate of \( \beta \) is conditional on the selected model. A brief description of the options for the `vselect` command is given in Section 4.3.

`bma` provides the BMA estimate of \( \beta \) in the ‘grand’ model (2) using the `bma` command implemented by De Luca and Magnus (2011). This option assumes a classical Gaussian linear model for (2), noninformative priors for the regression parameter \( \beta \) and the error variance, and a multivariate Gaussian prior for the auxiliary parameter \( \delta \). This estimate is obtained as a weighted average of the estimates of \( \beta \) from each of the \( 2^{JK} \) possible models in the model space with weights proportional to the marginal likelihood of `depvar` in each model. A brief description of the options for the `bma` command is given in Section 4.4.

`wals` provides the WALS estimate of \( \beta \) in the ‘grand’ model (2) using the `wals` command implemented by De Luca and Magnus (2011). Like BMA, this option assumes a classical Gaussian linear model for (2) and noninformative priors for the regression parameter \( \beta \) and the error
variance. Unlike BMA, WALS uses orthogonal transformations of the auxiliary regressors and their parameters, which reduces to $JK$ the order of magnitude of the required calculations. Further, the transformed auxiliary parameters are assumed to be identically and independently distributed according to either a Laplace or a Subbotin prior. A brief description of the options for the \texttt{wals} command is given in Section 4.5.

\texttt{full} requires to display the estimation results for all model parameters (i.e. focus and auxiliary parameters) and to return the associated estimates and their variance-covariance matrix in the vector $\texttt{e(b)}$ and the matrix $\texttt{e(V)}$ respectively. By default, display of the estimation results is restricted to the focus parameters of interest, the associated estimates and their variance-covariance matrix are returned in the vector $\texttt{e(b)}$ and the matrix $\texttt{e(V)}$ respectively, while estimates of the auxiliary parameters and their variance-covariance matrix are returned in the vector $\texttt{e(b\_aux)}$ and the matrix $\texttt{e(V\_aux)}$ respectively.

\texttt{vce(bootstrap [, bootstrap_options])} uses wild bootstrap to estimate the variance-covariance matrix of the parameter estimates (see \texttt{[R] bootstrap}). By default, bootstrap estimates of the variance-covariance matrix are computed only for the focus parameters. To obtain bootstrap estimates of the variance-covariance matrix the focus and the auxiliary parameters, the option \texttt{vce(bootstrap)} must be combined with the option \texttt{full}. In any case, \texttt{vce(bootstrap)} and \texttt{full} cannot be jointly specified when applying model reduction techniques (i.e. the options \texttt{sw} and \texttt{vs}) because the subset of selected auxiliary regressors can vary across bootstrap replicates. Standard options for bootstrap estimation can be specified as sub-options within \texttt{vce(bootstrap)} (see \texttt{[R] vce option}).

\texttt{auxiliary(string)} specifies the prefix for the name of the auxiliary regressors. The default is $D$.

Thus, auxiliary regressors are named as $Dj$ and $Dj\_varname$ where $j = 1, \ldots, J$ is an index for the sub-samples of missing data and \texttt{varname} is the name of each variable listed in \texttt{varlist}\ and \texttt{imputed}.

\texttt{keep} specifies whether auxiliary regressors have to be kept in the data after estimation. By default, they are dropped.

\texttt{nowarn} suppresses the display of a warning message on dropped collinear regressors.

### 4.2 Options for stepwise

By specifying the \texttt{sw} option, \texttt{gmi} carries out model reduction through the build-in Stata command \texttt{stepwise} (see \texttt{[R] stepwise} for details). The relevant options of the \texttt{stepwise} command are $\texttt{pr(\#)}$
(significance level for backward selection), \texttt{pe(\#)} (significance level for forward selection), \texttt{forward} (backward stepwise) and \texttt{lr} (likelihood-ratio test of term significance). Since the auxiliary regressors in $Z$ have no hierarchical ordering, backward hierarchical selection and forward hierarchical selection are not allowed.

### 4.3 Options for \texttt{vselect}

By specifying the \texttt{vs} option, \texttt{gmi} carries out model reduction through the \texttt{vselect} command implemented by Lindsey and Sheather (2010). This command offers three model reduction techniques: backward selection (the default), forward selection (\texttt{forward}), and leaps-and-bounds selection (\texttt{best}). An information criterion is used to judge the validity of each model through the options \texttt{r2adj} (adjusted $R^2$), \texttt{aic} (AIC), \texttt{aicc} (corrected AIC), \texttt{bic} (BIC), \texttt{cp1} or \texttt{cp2} (Mallow’s $C_p$). Mallow’s $C_p$ criterion can only be used with leaps-and-bounds selection and the decision rule can be either a value of $C_p$ close to zero (\texttt{cp1}) or a value close to the number of covariates (\texttt{cp2}). For additional information see Lindsey and Sheather (2010).

### 4.4 Options for BMA

By specifying the \texttt{bma} option, \texttt{gmi} carries out BMA through the \texttt{bma} command implemented by De Luca and Magnus (2011). In this case, one can use two additional options. The option \texttt{scaling} provides an alternative way of scaling the model weights $\lambda_i$ when the default scaling procedure suffers from numerical problems. Although scaling of the model weights does not affect BMA estimates, the computing time required by this alternative procedure is almost double. The option \texttt{nodots} suppresses the display of the dots to track the progress of \texttt{bma} estimation. By default, dots are displayed only if the model space consists of more than 128 models. One dot means that 1% of the models in the model space have been estimated.

### 4.5 Options for WALS

By specifying the \texttt{wals} option, \texttt{gmi} carries out model averaging through the \texttt{wals} command implemented by De Luca and Magnus (2011). As for the prior on the transformed auxiliary parameters, one can choose between Laplace or Subbotin priors through the option \texttt{q(\#)} which defines the free parameter $0 < q \leq 1$ of a Subbotin density with the prior median of $\eta_h$ equal to zero and the prior median of $\eta_h^2$ equal to one. The default is $q = 1$ corresponding to a Laplace prior. Values of $q$ in the interval $(0, 1)$ give instead a class of Subbotin priors. Einmahl et al. (2011) argue that values of $q$ close to zero are unappealing from the point of view of ignorance. For empirical applications,
they recommend $q = 0.5$. For a Subbotin prior with $q \neq 1$ and $q \neq 0.5$, one can also specify a set of additional options (i.e. `intpoints(#)`, `eps(#)` and `iterate(#)) to control the accuracy of the numerical process for approximating the constrained parameter of a Subbotin density. Additional information can be found in De Luca and Magnus (2011).

5 Example

This section illustrates the `gmi` command using data available on the Stata website.

```
. describe
obs: 1,647  Albuquerque Home Prices Feb15-Apr 30, 1993
vars: 13  19 Jun 2009 10:50
size: 54,351 (99.9% of memory free)  (_dta has notes)

variable name storage display value label variable label
variable name storage display value label variable label
price  int  %8.0g  Sale price (hundreds)
sqft  int  %8.0g  Square footage of living space
age  float  %10.0g  Home age (years)
ne  byte  %8.0g  Number of certain features
ne  byte  %8.0g  Located in northeast (largest
custom  byte  %8.0g  Custom build
corner  byte  %8.0g  Corner location
tax  float  %10.0g  Tax amount (dollars)
lnage  float  %9.0g  
lnage  float  %9.0g  
_mi_miss  byte  %8.0g  
_mi_m  int  %8.0g  
_mi_id  int  %12.0g  
```

We want to estimate a classical linear regression model for the relationship between home sale price (`price`) and home characteristics (`sqft`, `nfeatures`, `ne`, `custom`, `corner`, `lnage` and `lnage`). Since there are cases with `age` and `tax` missing, `lnage` and `lnage` are affected by a missing-data problem and their missing values have been imputed using a multivariate normal regression model (see `mi impute mvn`).

```
. mi describe
Style: mlong
last mi update 19jun2009 10:50:22, 243 days ago
Obs.: complete  66
incomplete  51  (M = 30 imputations)
total  117
Vars.: imputed:  2; lnage(49) lnage(10)
passive:  2; age(49) tax(10)
regular:  6; price sqft nfeatures ne custom corner
```
system: 3; _mi_n _mi_id _mi_miss
(there are no unregistered variables)

. mi misstable summarize lnage lntax

<table>
<thead>
<tr>
<th>Variable</th>
<th>Obs&lt;.</th>
<th>Obs&gt;</th>
<th>Obs&lt;.</th>
</tr>
</thead>
<tbody>
<tr>
<td>lnage</td>
<td>49</td>
<td>68</td>
<td>30</td>
</tr>
<tr>
<td>lntax</td>
<td>10</td>
<td>107</td>
<td>95</td>
</tr>
</tbody>
</table>

Thus, the data contain 117 observations plus 30 multiple imputations stored in the mlong style (see styles) for each of the 51 incomplete cases. Below, we generate the missing-data indicators for lnage and lntax and the local first_imp which is used to restrict the estimation sample to the first imputation. Continuous covariates are centered to their median values to obtain meaningful estimates of the constant term.

. generate mis_lnage=(lnage==.)
. generate mis_lntax=(lntax==.)
. bys _mi_id: egen M_lnage=max(mis_lnage)
. bys _mi_id: egen M_lntax=max(mis_lntax)
. local first_imp "(_mi_miss==0|_mi_m==1)"
. foreach x of varlist sqft nfeatures lnage lntax {
  2. quietly summarize x if first_imp, d
  3. quietly replace `x'=`x'-(p50)
  4. }

The gmi command with its default option summarize produces the following output:

```
. gmi price sqft nfeatures ne custom corner if `first_imp', ///
> imp(lnage lntax) mis(M_lnage M_lntax)
note: D1_nfeatures D1_ne D1_custom D1_corner D1_lnage D1_lntax D3_corner omitted
> d because of collinearity

Grand model
```

| Number of obs | 117 |
| Number of observed covariates | 6 |
| Number of imputed covariates | 2 |
| Number of focus covariates | 8 |
| Number of missing data patterns | 3 |
| Number of auxiliary covariates | 17 |
| Dimension of model space | 131072 |

```
<table>
<thead>
<tr>
<th>Summary of price by missing data pattern</th>
<th>Missing data patterns</th>
</tr>
</thead>
<tbody>
<tr>
<td>Group</td>
<td>Freq.</td>
</tr>
<tr>
<td>-------</td>
<td>-------</td>
</tr>
<tr>
<td>0</td>
<td>66</td>
</tr>
<tr>
<td>1</td>
<td>2</td>
</tr>
<tr>
<td>2</td>
<td>41</td>
</tr>
<tr>
<td>3</td>
<td>8</td>
</tr>
</tbody>
</table>
```

Our model includes 8 focus regressors, of which 6 (including the constant term) are observed and 2 are imputed. Excluding the subset of complete cases (66 observations), there are \(2^2 - 1 = 3\)
missing-data patterns: (i) \texttt{lnage} observed and \texttt{lntax} missing (2 observations), (ii) \texttt{lnage} missing and \texttt{lntax} observed (41 observations), and (iii) \texttt{lnage} and \texttt{lntax} both missing (8 observations). The grand model therefore includes $3 \cdot 8 = 24$ auxiliary variables, but 7 of them are dropped because of perfect collinearity. In particular, because the variable \texttt{corner} is constant for the third missing-data pattern, the auxiliary variables \texttt{D3} and \texttt{D3\_corner} are perfectly collinear, so the latter is dropped.

Other 6 auxiliary variables are dropped because the first missing-data pattern includes only 2 observations, so we can identify at most 2 of the 8 associated auxiliary parameters. After dropping from \texttt{Z} all collinear variables, the dimension of the model space reduces to $2^{17} = 131072$. The summary statistics for the dependent variable across missing-data patterns reveal that both the mean and the variance of \texttt{price} are considerably higher for the subsample with complete cases.

We obtain the complete-case OLS estimator of the focus parameters $\beta$ by specifying the \texttt{cc} option.

These estimates could also be obtained through the built-in Stata command \texttt{regress} after restricting the estimation sample to the subset of complete data. They are also numerically the same as the OLS estimate of $\beta$ in the grand model (2). Result 1 implies that, under the ignorability assumption, the complete-case OLS estimator is unbiased for $\beta$. Our findings suggest that home sale price is positively related to square footage of living space, log of taxes paid and whether the home is located in a custom building. On the other side, there is negative association with log of home
age and whether the home has a corner location. The effects of the other covariates are not robust because the corresponding $t$-ratios are smaller than one in absolute value. It is also worth noticing that the complete-case estimator is likely to be highly inefficient as it discards about 44 percent of the sample observations.

To explore the trade-off between bias and precision, consider now the ‘naive’ and the simple missing-indicator approaches. The former ignores the fact that missing values have been imputed by restricting all auxiliary parameters to zero, while the latter restrict all auxiliary parameters to zero except the coefficients on the dummies for the missing-data patterns.

```
. gmi price sqft nfeatures ne custom corner if `first_imp', ///
    imp(lnage lntax) mis(M_lnage M_lntax) naive nowarn
Naive OLS estimates
Number of obs = 117
df_m = 7

_cons     984.3707    35.50699  27.72 [948.8638 1019.878]
sqft      0.382786    0.0729738  5.25  [0.3098122 0.4557598]
nfeatures 3.622533    13.89274  0.26 [-10.27021 17.51527]
ne        28.93578    37.16146  0.78 [-8.225879 66.09725]
custom    145.1389    46.45179  3.12  [98.68716 191.5907]
corner   -85.8675    42.73586 -2.01 [-128.6034 -43.13164]
lnlntax  262.9705    106.5927  2.47  [156.3778 369.5632]
```

```
. gmi price sqft nfeatures ne custom corner if `first_imp', ///
    imp(lnage lntax) mis(M_lnage M_lntax) smi nowarn
SMI OLS estimates
Number of obs = 117
df_m = 10

_cons    1007.3570    35.88174  28.07 [971.4752 1043.239]
sqft     0.3993985    0.0718978  5.56  [0.3275006 0.4712963]
nfeatures -5.977141   14.29397 -0.42 [-20.27111 8.316833]
ne       49.92553    37.20047  1.34  [12.72506 87.12601]
custom   157.4772    47.33692  3.33  [110.1403 204.8141]
corner  -103.4662   42.61305 -2.43 [-146.0793 -60.85319]
lnage   -30.55087   21.47985 -1.42 [-52.03073 -9.071018]
lnlntax 204.6133    108.1598  1.89  [96.45353 312.7731]
```

```
. matrix list e(b_aux)
e(b_aux)[1,3]  
    D1       D2       D3  
y1    -119.71306  -82.584248 -164.86743

. matrix list e(V_aux)
  symmetric e(V_aux)[3,3]  
    D1       D2       D3  
D1 18343.655  
D2 826.58661 1662.3468  
D3 527.1418  680.08892 4520.6579
```

Both approaches impose arbitrary restrictions on the auxiliary parameter $\delta$, so they are likely to result in biased estimates of the focus parameter $\beta$. However, as suggested by their considerably
lower standard errors, these estimators are more precise than the complete-case OLS estimator. The most striking differences are in the estimated coefficients of \text{corner} and \text{lntax}. Notice that, to force users to treat the auxiliary parameters as nuisance parameters, their estimates and the associated variance-covariance matrix are returned in the vector \text{e(b\_aux)} and the matrix \text{e(V\_aux)} respectively.

The \text{gmi} command provides two alternative strategies for finding a better estimator of $\beta$ in the MSE sense: model reduction and model averaging. Although the choice between these two strategies is left to the users, we strongly encourage choosing model averaging in order to avoid the problems caused by pretesting.

Model reduction can be carried out through the built-in Stata command \text{stepwise} or the \text{vselect} command by Lindsey and Sheather (2010). There are reasons to prefer the latter, as model reduction is based on an information criterion instead of an arbitrary significance level, and the leaps-and-bounds algorithm is expected to select the best model. To save space, we only present the OLS estimates of the model selected by \text{vselect} with the \text{best} and the \text{bic} options.

\begin{verbatim}
. gmi price sqft nfeatures ne custom corner if first_imp, ///
> imp(lnage lntax) mis(M_lnage M_lntax) vs best bic full nowarn
Model reduction: leaps and bounds with bic Number of obs = 117
   df_m = 9

<table>
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<tr>
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<tbody>
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<td>32.5224</td>
<td>30.24</td>
</tr>
<tr>
<td>sqft</td>
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<td>0.0722097</td>
<td>6.80</td>
</tr>
<tr>
<td>nfeatures</td>
<td>1.022459</td>
<td>12.73723</td>
<td>0.08</td>
</tr>
<tr>
<td>ne</td>
<td>6.726864</td>
<td>34.54129</td>
<td>0.19</td>
</tr>
<tr>
<td>custom</td>
<td>163.2298</td>
<td>43.00966</td>
<td>3.80</td>
</tr>
<tr>
<td>corner</td>
<td>-80.96139</td>
<td>39.22133</td>
<td>-2.06</td>
</tr>
<tr>
<td>lnage</td>
<td>-25.25726</td>
<td>19.84414</td>
<td>-1.27</td>
</tr>
<tr>
<td>lntax</td>
<td>257.7811</td>
<td>98.19124</td>
<td>2.63</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>D2_sqft</td>
<td>-0.2688726</td>
<td>0.0622148</td>
<td>-4.32</td>
</tr>
<tr>
<td>D3_custom</td>
<td>-400.7815</td>
<td>168.7942</td>
<td>-2.37</td>
</tr>
</tbody>
</table>
\end{verbatim}

In this case, we specified the \text{full} option to display estimates of the focus and the auxiliary parameters. The selected model includes two auxiliary variables: the interaction between \text{sqft} and the dummy \text{D2} for the second missing-data pattern, and the interaction between \text{custom} and the dummy \text{D3} for the third missing-data pattern. Notice that, the standard errors are conditional on the model selected by \text{vselect} and therefore should be treated with caution.

Next, we focus on model averaging using BMA and WALS respectively.

\begin{verbatim}
. gmi price sqft nfeatures ne custom corner if first_imp, ///
> imp(lnage lntax) mis(M_lnage M_lntax) bma nowarn
Model space: 131072 models
\end{verbatim}
Magnus et al. (2010) argue that WALS is theoretically superior to BMA in the choice of priors for the auxiliary parameters, and is practically superior because of the substantially lower computational burden. Although the Stata command `bma` is much faster than the Magnus’ original Matlab routine, we recognize that BMA can be very time consuming when the covariates or missing-data patterns is moderate or large. In such circumstances, users are encouraged to rely on WALS, at least when performing preliminary model specification search. In this example, we find that most of the estimated coefficients from BMA and WALS are similar, suggesting that differences in the priors for the auxiliary parameters play a minor role. Similar findings are also supported by the estimates from WALS with a Subbotin prior for the auxiliary parameters.
In the above example, standard errors are estimated by the wild bootstrap with 100 replications. Bootstrapped standard errors are usually larger than traditional ones because they account for heteroskedasticity of unknown form. As argued in Section 3.3, the wild bootstrap also provides an easy way to ensure comparability of the standard errors across the different estimation methods.

Finally, we can use the 30 multiple imputations on lnage and lntax to account for the sampling variability induced by imputation of missing values. This can be done by combining our `gmi` command with the built-in Stata prefix `mi estimate`.

```plaintext
.mi estimate: gmi price sqft nfeatures ne custom corner, ///
> imp(lnage lntax) mis(M_lnage M_lntax) wals q(.5) nowarn full
```

```
Multiple-imputation estimates
Imputations = 30
WALS estimates - Subbotin(q=0.5) priors
Number of obs = 117
Average RVI = 0.1202
Complete DF = 92
DF adjustment: Small sample
DF: min = 53.04
          = 81.87
max = 89.46
Model F test: Equal FMI
F( 24, 89.4) = 18.93
Prob > F = 0.0000
```

```
price | Coef.   Std. Err.      t    P>|t|     [95% Conf. Interval]
------|--------|----------------------|---------|--------|----------------------------------------
    _cons | 991.3835   37.04141    26.76    0.000  917.7875   1064.979
        sqft | 0.4317465   0.0867584    4.98    0.000    0.2592937    0.6041993
        nfeatures | -0.2094938  16.38439   -0.01    0.990  -32.76653    32.34754
        ne | 21.21545   42.39104    0.50    0.618  -63.00865    105.4396
        custom | 169.9147    50.06324    3.39    0.001  109.44587   229.38336
        corner | -78.7322   46.13668   -1.71    0.091  -174.9073    16.4451
        lnage | -42.43074   26.08771   -1.63    0.108  -94.33088    0.469404
        lntax | 280.3926   127.9489    2.19    0.031   26.02861    534.7565
        D1 | -113.878   110.019    -1.04    0.304  -332.663    104.9071
        D1_sqft | 0.3657415  0.4004362    0.91    0.364  -0.430267    1.16175
        D2 | -54.71213   69.21678   -0.79    0.431  -192.3361    82.91185
        D2_sqft | -0.089371   0.1178286   -0.75    0.452  -0.3038581    0.024081
        D2_nfeatures | -3.48209    25.28178   -0.14    0.891  -53.76192    46.79774
        D2_ne | 17.66156    65.09508    0.27    0.787  -111.7321   147.0552
        D2_custom | -33.78444   80.12972   -0.42    0.674  -193.1156   125.5476
        D2_corner | -42.7017    74.38713   -0.57    0.567  -190.6397   105.2363
        D2_lnage | 39.95991    41.30198   -0.97    0.335  -46.79896    86.78792
        D2_lntax | -151.7439   171.2648   -0.89    0.378  -492.8674    189.3797
        D3 | -167.1674   275.5164   -0.61    0.546  -716.1743   381.8394
        D3_sqft | 0.176931    0.8775508   0.20    0.841  -1.569222    1.922806
        D3_nfeatures | -25.63685   77.73594   -0.33    0.742  -180.3954   129.1217
        D3_ne | 146.6077    276.0254    0.53    0.599  -404.0849    695.3002
```

18
The prefix `mi estimate` runs the specified `gmi` command on each imputed dataset to obtain a set of alternative estimates of the model parameters and their variance-covariance matrix. Multiple imputation estimates are then obtained by applying the combination rules of Rubin (1987) on the resulting set of alternative estimates (see `mi estimate`). Here, it is worth noticing that the prefix `mi estimate` has its own reporting output and does not respect the reporting output of the `gmi` command. As discussed in Section 3.3, p-values and confidence intervals must then be treated with caution especially when the number multiple imputations is small. Also notice that the option `full` of the `gmi` command is always needed to obtain valid information on the average relative variance increase (RVI) due to nonresponse and the summaries about parameter-specific degrees of freedom (DF). In any case, the prefix `mi estimate` and the option `full` of the `gmi` command cannot be jointly combined when using model reduction techniques (i.e. options `sw` and `vs`) because the subset of selected auxiliary regressors can vary across imputations.

### 6 Empirical application

This application investigates the relationship between hand grip strength (GS) and a set of socio-demographic and economic covariates using data on the elderly European population. As argued by Andersen et al. (2009), GS is an important measure of health because it is objectively measured, it directly affects every day activity functions, it is known to decline linearly with age, and it is a strong predictor of disability, morbidity, frailty and mortality. Furthermore, measuring GS is cheap and can be carried out by trained survey interviewers in non-clinical studies.

Our data are from release 2.4.0 of the first wave of the Survey of Health, Ageing and Retirement in Europe (SHARE), a multidisciplinary and cross-national household panel survey coordinated by the Mannheim Research Institute for the Economics of Aging (MEA).² SHARE collects data on self-reported and objective measures of health, socio-economic status, and social and family networks for nationally representative samples of elderly people in the participating countries. The first wave, conducted in 2004, covers about 28,500 individuals in 11 European countries (Austria, Belgium, Denmark, France, Germany, Greece, Italy, the Netherlands, Spain, Sweden and Switzerland). In each country, the target population consists of people aged 50 and older, plus their possibly younger

---

² Data can be downloaded free of charge from the SHARE Research Data Center: http://www.share-project.org. To get access to the dataset, researchers have to complete a statement concerning the use of the microdata.
partners. Although all national samples are selected through probability sampling, the sampling procedures are not completely standardized across countries because of the lack of a common sampling frame. The unweighted household nonresponse rate in the first wave ranges between a maximum of 62 percent in Switzerland and a minimum of 26 percent in France, and is equal to 45 percent on average. The data collection mode is computer-assisted personal interview (CAPI), supplemented by show-cards and a self-administered paper-and-pencil questionnaire.³

The GS test in SHARE involves two measurements on each hand (alternating between the two hands) using a hand-grip dynamometer. Before the fieldwork period, interviewers participated in centrally designed training sessions to learn a common protocol for measuring GS. They are instructed to demonstrate the use of the hand-grip dynamometer, help the respondent to assume a suitable position before performing the test, and verbally encourage the respondent to squeeze the handles of the hand-grip dynamometer as hard as possible. Respondents are excluded from the GS test only for swelling, inflammation, severe pain, recent injury or surgery to both hands in the last 6 months. For respondents with problems in one hand, the GS test is performed on the other hand only. The measurement of GS on each hand is considered valid if the two assessments on the same hand were greater than 0 Kg, lower than 100 Kg and they do not differ from each other by more than 20 kg. The overall GS test is considered valid if there is at least one valid measurement on one hand. Following Andersen et al. (2009), our dependent variable is the maximum GS (‘maxgrip’) measurement resulting from valid test.

Our set of socio-demographic and economic covariates includes age, gender, macro-region of residence (Northern, Central or Southern countries), self-reported weight and height, an indicator for educational attainment, per-capita household income and household net worth. To ensure cross-country comparability of the information on educational attainment, the original values have been recoded using the 1997 International Standard Classification of Education (ISCED-97). For similar reasons, per-capita household income and household net worth have been adjusted for the differences in purchasing power across countries. Thus, nominal amounts have been divided by the national purchasing power parity to obtain real amounts denominated in German prices for the year 2005.⁴

Unlike Andersen et al. (2009), who use imputed values of household income and household net worth by relying on the estimates from the ‘naïve’ approach, we are interested in investigating

³ For additional information on survey design and response rates see Börsch-Supan et al. (2005).
⁴ Data for these calculations are obtained from the purchasing power parity survey carried out by OECD in 2005. Further information can be found in the SHARE documentation of release 2.4.0.
the trade-off between bias and precision when replacing the missing values on these two variables with imputations. This is an important issue to consider because these covariates are affected by substantial item nonresponse. The item nonresponse rates for household income and household net worth range, respectively, between a maximum of 76 and 77 percent in Belgium and a minimum of 49 and 52 percent in Greece, and are equal to 62 and 64 percent on average. The substantial amount of item nonresponse reflects three problems. First, these variables are not asked directly to respondents but are obtained by aggregating a large number of income and wealth components (27 and 13 respectively), collected at both the individual and the household level. Second, information about incomes, real and financial assets, mortgage and other debts are asked through open-ended and retrospective questions that are sensitive and difficult to answer. Third, according to SHARE fieldwork rules, a household with two spouses is considered as interviewed if at least one of them agrees to participate. If the other does not, then household income and household net worth must be imputed because the individual components are missing for the nonresponding spouse.

To deal with the potential selectivity effects generated by item nonresponse, the public-use SHARE data include 5 multiple imputations of the key survey variables. As discussed at length in Christelis (2011), these imputations are constructed by the multivariate iterative procedure of Buuren et al (2006) which attempts to preserve the correlation structure of the imputed data. In what follows, we account for the additional sampling variability induced by the imputation process using the combination rules proposed by Rubin (1987) on the 5 multiple imputations of household income and household net worth.

Another important difference with respect to Andersen et al. (2009) is that we focus on respondents aged between 50 and 80 years who do not report serious health problems. This choice is primarily motivated by the need of compensating for cross country differences in coverage of the institutionalized target population. While the sampling frames of Denmark, Netherlands and Sweden cover people living in institutions for the elderly, this segment of the 50+ population is excluded by the national sampling frames of the other SHARE countries. Moreover, Southern European countries are known to have fewer nursing home than Northern and Continental European countries and a cultural tradition of old parents living with a child. To limit the impact of these cross country differences, we select respondents who have at most one limitation with activities of daily living, at most one chronic disease, and whose self-reported health status is at least fair. After applying this sample selection criterion, dropping the invalid measurements of maxgrip (about 5 percent of the cases) and the few missing data on weight, height and education (about 1 percent of
the cases), our working sample consists of 13,724 observations. Summary statistics for the outcome and the covariates are presented in Table 1, separately by gender and macro-region.

Given the high level of comparability of the SHARE data, we pool data from countries in the same macro-region, and estimate our linear regression model of interest separately by gender and macro-region. For simplicity, we assume that the errors in the grand model are independent and spherically distributed. The model specification in each subgroup includes 7 focus regressors, of which 5 (age, weight, height, education and the constant term) are observed and 2 (per-capita household income and household net worth) are imputed, 3 subsamples with incomplete data, and 21 non-collinear auxiliary variables. The resulting dimension of the model space is 2,097,152. After centering the focus covariates on the corresponding medians of each subgroup, we compare the estimates from five alternative approaches: complete case, ‘naive’, model reduction, BMA and WALS. Model reduction estimation is carried out using the vs estimation option of the gmi command with leaps-and-bounds selection and AIC as model information criteria, while WALS estimation is carried out using a Subbotin prior with parameter $q = 0.5$.\footnote{Estimates from the simple missing-data indicator approach are omitted because similar to those obtained from the ‘naive’ approach. Estimates from WALS with a Laplace prior are instead omitted because very similar to those obtained with a Subbotin prior.}

The estimated coefficients and their standard errors are presented in Tables 2 and 3, separately by gender and macro-region. Qualitatively, our results are consistent with the empirical findings in Andersen et al. (2009). In all specifications, maxgrip is negatively related to age and positively related to self-reported weight and height. Women have a lower level of maxgrip than men, but they also present a considerably flatter decline with advancing age. The positive gradient between Northern-Continental and Southern countries persists even after focusing on the healthier segment of the elderly European population. For men, the age-related decline in maxgrip is steeper for those living in Southern countries. For women, it is instead steeper for those living in Northern and Continental countries. Education, per-capita household income and household net worth do not seem to be robustly correlated with maxgrip. The only exceptions are the positive correlations between maxgrip and education for men and women living in Continental countries, between maxgrip and per-capita household income for women living in Southern countries, and between maxgrip and household net worth for men and women living in Southern countries.

Notice that, although there is broad agreement with previous studies on the sign of the estimated associations, their magnitude and the size of the standard errors are subject to non-negligible differences. For example, the point estimate of the coefficient on weight in the specification Male-
North ranges between a minimum 0.106 with a standard error of 0.022 using the ‘naive’ approach to a maximum of 0.214 with a standard error of 0.057 using complete-case analysis. Similar differences are observed for the estimated coefficients on education in the specifications Male-North and Male-Center, household net worth in the specification Male-South, weight in the specification Female-Center, and per-capita household income in the specification Female-South. The estimates from model reduction and model averaging are somewhat in-between the estimates from the complete-case and the ‘naive’ approach. In particular, the conditional estimates from model reduction are quite close to the unconditional estimates from BMA. This suggests that, in this example, the effects of pretesting are not very important. The differences in the unconditional estimates from BMA and WALS suggest instead that alternative assumptions on the prior distributions for the auxiliary parameters may matter. From this view point, WALS has the advantage of using priors that ensure bounded risk and a coherent treatment of ignorance about the auxiliary parameters.
References


Table 1: Descriptive statistics for the dependent and the independent variables (weight is in kilograms, height is in centimeters, PPP-adjusted per-capita household income is in 10,000 Euro and household net worth is 100,000 Euro).

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<th>Region</th>
<th>Variable</th>
<th>Male P25</th>
<th>Med</th>
<th>P75</th>
<th>Female P25</th>
<th>Med</th>
<th>P75</th>
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<td>North</td>
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</tr>
<tr>
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Table 2: Estimated coefficients and standard errors (in parentheses) for males by macro-region. Estimation is based on $M = 5$ multiple imputations for income and net worth. Results for the auxiliary regressors are omitted to save space.

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Table 3: Estimated coefficients and standard errors (in parentheses) for females by macro-region. Estimation is based on $M = 5$ multiple imputations for income and net worth. Results for the auxiliary regressors are omitted to save space.

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