

IMPUTE Example #2

SUDAAN Statements and Results Illustrated

- WSHD Multiple Imputations
- NOTSORTED option
- IMPBY
- IMPVAR (MULTIMP option)
- IMPNAME
- PRINT and OUTPUT

Input Data Set(s): WIC.SAS7bdat

Example

Using data from the Women, Infants and Children (WIC) study, perform multiple imputations (impute 4 times) for missing values of the length of the mother's hospital stay. Use information on the mother's race and smoking behavior as imputation class data.

Solution

The primary purpose of this example is to illustrate how to perform multiple imputations in a single call to IMPUTE. The SAS-callable version of SUDAAN was used to generate the output in this example.

The following variables from the WIC dataset are of interest in this example:

<u>Variable</u>	<u>Definition</u>
ID	Observation number
ANALWGT1	Final Weight
MOMRACE	Mother's Race (2 levels)
MOMSMK	Mother's Smoking Behavior (2 levels)
MOMHOSP	Length of Mother's Hospital Stay (in days)

Prior to performing WSHD imputation, PROC CROSSTAB is used to review the distribution of the imputation variable, MOMHOSP, and PROC RECORDS is used to print the nonrespondents. The CROSSTAB and RECORDS code and results are shown next.

Exhibit 1. CROSSTAB and RECORDS Code

```
LIBNAME in "\\rtints29\sudaan\data\wicwage";
libname out "c:\llwinbetatest\examples";

data one; set in.wicwage;

PROC CROSSTAB data=one design=srs;
  class momrace momsmk momhosp / include=missing;
  tables momrace momsmk momhosp;
  print nsum colper / nsumfmt=f6.0 colperfmt=f7.2 style=nchs;
  rtitle "Distributions";

PROC RECORDS DATA=one;
  subpopx momhosp=.;
  sortby momrace momsmk ;
  setenv colwidth=9 labwidth=10 decwidth=0;
  PRINT id momrace momsmk momhosp;
  rtitle "List the MOMHOSP Non-Respondents";
```

Since we are just interested in sample distributions here, we can specify the SRS design in CROSSTAB. And RECORDS does not use design information at all.

Exhibit 2. First Page of CROSSTAB Results

```

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DESIGN SUMMARY: Variances will be computed using the Taylor Linearization Method,
Assuming a
Simple Random Sample (SRS) Design

Number of observations read      :      953
Denominator degrees of freedom :      952
```

Exhibit 3. CROSSTAB Results: Sample Distribution of Mother's Race

Variance Estimation Method: Taylor Series (SRS)

Distributions

by: Mother Race.

Mother Race

	Sample Size	Col Percent
Total	953	100.00
1	480	50.37
2	473	49.63

Exhibit 4. CROSSTAB Results: Sample Distribution of Mother's Smoking Behavior

Variance Estimation Method: Taylor Series (SRS)

Distributions

by: Smoking behavior.

```
-----  
Smoking behavior  
                Sample   Col  
                Size     Percent  
-----  
Total                953    100.00  
Missing                21     2.20  
1                    230    24.13  
2                    702    73.66  
-----
```

There are 21 missing values for the imputation class variable mother's smoking.

Exhibit 5. CROSSTAB Results: Sample Distribution of Length of Hospital Stay (Days)

Variance Estimation Method: Taylor Series (SRS)

Distributions

by: Length of hospital stay -- mother.

```
-----  
Length of hospital  
  stay -- mother      Sample   Col  
                      Size     Percent  
-----  
Total                953    100.00  
Missing                21     2.20  
0                     25     2.62  
1                    320    33.58  
2                    350    36.73  
3                    142    14.90  
4                     48     5.04  
5                     22     2.31  
6                      7     0.73  
7                      4     0.42  
8                      2     0.21  
10                    1     0.10  
11                    1     0.10  
12                    3     0.31  
13                    7     0.73  
-----
```

There are 21 missing values for the imputation variable length of hospital stay.

Exhibit 6. RECORDS Results: Length of Hospital Stay Nonrespondents

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For Subpopulation: MOMHOSP = .

List the MOMHOSP Non-Respondents

by: Observation Number.

Observation Number	ID	Mother Race	Smoking behavior	Length of hospital stay -- mother
60	34129	1	1	.
63	21127	1	1	.
212	11013	1	1	.
234	41131	1	1	.
242	3127	1	2	.
464	3122	1	2	.
465	21126	1	2	.
468	30124	1	2	.
493	34131	1	2	.
496	22130	1	2	.
502	22136	1	2	.
518	23122	1	2	.
630	21130	1	2	.
631	30123	1	2	.
673	41129	1	2	.
674	41128	1	2	.
767	28125	2	1	.
769	28126	2	2	.
932	11125	2	2	.
933	10017	2	2	.
935	22127	2	2	.

In the next step of this example, we impute the missing values via PROC IMPUTE. The imputation will be performed 4 times. The code is displayed next.

Exhibit 7. IMPUTE Code to get Multiple Imputations

```
PROC IMPUTE data=one seed=6537181 method=wshd icnum notsorted;
  weight analwgt1;
  impby momrace momsmk ;
  impvar momhosp / multimp=4;
  impname momhosp="mhosp";
  impid id;
  idvar / all;
  setenv labwidth=12;
  print / donorstat=default means=default donorsumfmt=f5.0 rsumfmt=f5.0
        absdiffmeanfmt=f7.4 reldiffmeanfmt=f7.4;
  output / impute=default filename=out.multimp filetype=sas replace;
  rtitle "Hot-Deck Multiple Imputations for MOMHOSP";

PROC RECORDS data=out.multimp contents;
  setenv labwidth=10 colwidth=3 decwidth=0 colspce=0;
  print id momrace momsmk donoid1 donoid2 donoid3 donoid4
        momhosp mhosp_1 mhosp_2 mhosp_3 mhosp_4 /
        idfmt=f5.0 donoid1fmt=f5.0 donoid2fmt=f5.0 donoid3fmt=f5.0
        donoid4fmt=f5.0;
  rlabel donoid1="dnr1";
  rlabel donoid2="dnr2";
  rlabel donoid3="dnr3";
  rlabel donoid4="dnr4";
  rlabel momhosp="momhos";
  rlabel momrace="momrac";
  rlabel momsmk="momsmk";
  rlabel mhosp_1="hsp1";
  rlabel mhosp_2="hsp2";
  rlabel mhosp_3="hsp3";
  rlabel mhosp_4="hsp4";
  rtitle "WSDH Multiply Imputed Dataset";

PROC FREQ data=out.multimp;
  tables momhosp*(mhosp_1 mhosp_2 mhosp_3 mhosp_4) mhosp_1 mhosp_2 mhosp_3 mhosp_4 /
  list missing;
  title "WSDH Multiple Imputations: See How MOMHOSP is Imputation Revised";
```

The NOTSORTED option is used on the PROC statement because the file was not sorted by the IMPBY statement variables.

The IMPVAR statement includes the MULTIMP=4 option. This tells SUDAAN to perform the WSHD imputation procedure 4 times. The output dataset will include the results of all 4 of the multiple imputations.

The IMPBY statement specifies MOMRACE and MOMSMK as the imputation class variables.

The IMPNAME statement specifies that the imputed values of MOMHOSP will be called MHOSP (with the _1, _2, _3, and _4 suffixes added for the multiple imputations.

The PRINT statement requests the DONORSTAT and MEANS print groups.

The OUTPUT statement includes the option IMPUTE=DEFAULT. This, used in conjunction with the ALL option on the IDVAR statement, requests that all variables from the input dataset be included on the output dataset MULTIMP, plus the following variables created in IMPUTE:

<u>Variable</u>	<u>Definition</u>
ORIGVAR1	Original imputation variable (values from MOMHOSP)
MHOSP_1	Imputed variable from Imputation #1
MHOSP_2	Imputed variable from Imputation #2
MHOSP_3	Imputed variable from Imputation #3
MHOSP_4	Imputed variable from Imputation #4
DONORID1	Donor ID from Imputation #1
DONORID2	Donor ID from Imputation #2
DONORID3	Donor ID from Imputation #3
DONORID4	Donor ID from Imputation #4

Exhibit 8. First Page of IMPUTE Results

```

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The Impute Process has completed successfully.
Method: WSHD

Total Records Read from File: 953
Random Number Seed: 6537181

Total Imputation Classes: 6

Total Respondent Records: 932

Total Nonrespondent Records: 21
  Total Donor Records Imputation #1: 21
  Total Donor Records Imputation #2: 21
  Total Donor Records Imputation #3: 21
  Total Donor Records Imputation #4: 21
Total Records Imputed: 21

```

The default summary output displayed above provides useful information about the IMPUTE run. The method of imputation is WSHD, the number of valid records read from the file is 953, the number of item respondent records is 932, the number of item nonrespondent records is 21, the number of donor records is 21, and the number of records imputed is 21 for each round of imputations. There are 6 imputation classes, since mother's race has 2 levels and smoking behavior has 3 levels (levels 1, 2, and missing). The number of respondent records plus the number of nonrespondent records equals the total records read from the file (932+21=953).

Exhibit 9. IMPUTE Results: DONORSTAT PRINT Group (Imputation #1)

```

Hot-Deck Multiple Imputations for MOMHOSP

Imputation #1
Method = WSHD

by: Mother Race, Smoking behavior.
-----
Mother Race
Smoking
  behavi-
  or      Item      Item      Donor      Missing
          Resp      Non-Resp  Count      Data -
          Count     Count     Count     Post-imp
-----
1
Missing      6          0          0          0
1            186        4          4          0
2            272       12         12         0
2
Missing      15         0          0          0
1            39         1          1          0
2            414        4          4          0
-----

```

This table is the DONORSTAT print table for imputation #1. The rows in this table represent the complete cross-classification of the IMBY variables. In this example, there are 6 imputation classes (missing values for Smoking Behavior forms its own level). For each row, the table tells us the number of records considered to be respondents and nonrespondents, the number of donors contributing to imputation, and the number of records with missing data post-imputation. So for MOMRACE=1 and MOMSMK=1, there are 186 respondent records, 4 nonrespondent records (missing MOMHOSP), 4 donors contributing data, and no records with missing MOMHOSP after imputation.

The next 3 tables correspond to imputation rounds #2, 3, and 4.

Exhibit 10. IMPUTE Results: DONORSTAT PRINT Group (Imputation #2)

```

Hot-Deck Multiple Imputations for MOMHOSP

Imputation #2
Method = WSHD

by: Mother Race, Smoking behavior.
-----
Mother Race
Smoking
  behavi-
  or      Item      Item      Donor      Missing
          Resp      Non-Resp  Count      Data -
          Count     Count     Count     Post-imp
-----
1
Missing      6          0          0          0
1            186        4          4          0
2            272       12         12         0
2
Missing      15         0          0          0
1            39         1          1          0
2            414        4          4          0
-----

```

Exhibit 11. IMPUTE Results: DONORSTAT PRINT Group (Imputation #3)

Hot-Deck Multiple Imputations for MOMHOSP

Imputation #3
Method = WSHD

by: Mother Race, Smoking behavior.

Mother Race		Item	Item	Donor	Missing
Smoking	behavi-	Resp	Non-Resp	Count	Data -
or	or	Count	Count	Count	Post-imp

1	Missing	6	0	0	0
	1	186	4	4	0
	2	272	12	12	0
2	Missing	15	0	0	0
	1	39	1	1	0
	2	414	4	4	0

Exhibit 12. IMPUTE Results: DONORSTAT PRINT Group (Imputation #4)

Hot-Deck Multiple Imputations for MOMHOSP

Imputation #4
Method = WSHD

by: Mother Race, Smoking behavior.

Mother Race		Item	Item	Donor	Missing
Smoking	behavi-	Resp	Non-Resp	Count	Data -
or	or	Count	Count	Count	Post-imp

1	Missing	6	0	0	0
	1	186	4	4	0
	2	272	12	12	0
2	Missing	15	0	0	0
	1	39	1	1	0
	2	414	4	4	0

Exhibit 13. IMPUTE Results: MEANS PRINT Group (Imputation #1)

```

Hot-Deck Multiple Imputations for MOMHOSP

Imputation #1
Method = WSHD
Length of hospital stay -- mother

by: Mother Race, Smoking behavior.
-----
Mother Race
Smoking
behavi-   Mean -   Mean -   Abs.   Rel.
or        Pre-imp  Post-imp Diff.   Diff.
              (%)
-----
1
Missing   1.77     1.77     0.0000  0.0000
1         2.13     2.13     0.0013  -0.0622
2         2.02     1.99     0.0322  -1.5943
2
Missing   1.47     1.47     0.0000  0.0000
1         2.27     2.26     0.0164  -0.7204
2         2.35     2.35     0.0010  0.0429
-----

```

This table corresponds to Imputation Round #1. It tells us the weighted mean of MOMHOSP before vs. after hot-deck imputation, within each imputation class. If no imputation took place within a cell, the weighted mean is the same before vs. after imputation. The first and fourth rows show no change since there were no nonrespondents in these cells. The remainder of the cells only shows small changes after imputation.

Exhibit 14. IMPUTE Results: MEANS PRINT Group (Imputation #2)

```

Hot-Deck Multiple Imputations for MOMHOSP

Imputation #2
Method = WSHD
Length of hospital stay -- mother

by: Mother Race, Smoking behavior.
-----
Mother Race
Smoking
behavi-   Mean -   Mean -   Abs.   Rel.
or        Pre-imp  Post-imp Diff.   Diff.
              (%)
-----
1
Missing   1.77     1.77     0.0000  0.0000
1         2.13     2.12     0.0057  -0.2688
2         2.02     2.01     0.0101  -0.5020
2
Missing   1.47     1.47     0.0000  0.0000
1         2.27     2.32     0.0438  1.9287
2         2.35     2.33     0.0121  -0.5147
-----

```

Exhibit 15. IMPUTE Results: MEANS PRINT Group (Imputation #3)

Hot-Deck Multiple Imputations for MOMHOSP

Imputation #3
 Method = WSHD
 Length of hospital stay -- mother

by: Mother Race, Smoking behavior.

Mother Race					
Smoking		Mean -	Mean -	Abs.	Rel.
behavi-		Pre-imp	Post-imp	Diff.	Diff.
or					(%)
<hr/>					
1	Missing	1.77	1.77	0.0000	0.0000
	1	2.13	2.13	0.0049	0.2315
	2	2.02	2.01	0.0147	-0.7268
2	Missing	1.47	1.47	0.0000	0.0000
	1	2.27	2.44	0.1642	7.2269
	2	2.35	2.36	0.0131	0.5580
<hr/>					

Exhibit 16. IMPUTE Results: MEANS PRINT Group (Imputation #4)

Hot-Deck Multiple Imputations for MOMHOSP

Imputation #4
 Method = WSHD
 Length of hospital stay -- mother

by: Mother Race, Smoking behavior.

Mother Race					
Smoking		Mean -	Mean -	Abs.	Rel.
behavi-		Pre-imp	Post-imp	Diff.	Diff.
or					(%)
<hr/>					
1	Missing	1.77	1.77	0.0000	0.0000
	1	2.13	2.13	0.0067	0.3136
	2	2.02	2.01	0.0071	-0.3491
2	Missing	1.47	1.47	0.0000	0.0000
	1	2.27	2.26	0.0164	-0.7204
	2	2.35	2.34	0.0003	-0.0112
<hr/>					

Exhibit 17. RECORDS Results: Contents of WSHD Imputed Dataset

S U D A A N				
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SAS Record File OUT.MULTIMP				
Variables				
Name	Type	Output Format	SAS Format	Description

PROCNUM	Numeric	F2.		Procedure Number
DONORID1	Numeric	F8.		dnr1
DONORID2	Numeric	F8.		dnr2
DONORID3	Numeric	F8.		dnr3
DONORID4	Numeric	F8.		dnr4
MHOSP_1	Numeric			hsp1
MHOSP_2	Numeric			hsp2
MHOSP_3	Numeric			hsp3
MHOSP_4	Numeric			hsp4
ID	Numeric	F5.		ID
MARITAL2	Numeric			Marital Status
BABYDOB	Numeric		MMDDYY8.	Birth date--Baby
BABYSEX	Numeric	F12.		Sex of baby
BABHOSP	Numeric			Length of hospital stay -- baby
MOMHOSP	Numeric			momhos
BORN_US	Numeric			Place of birth -- US
COUNTRY	Alpha	A20		Place of birth -- outside of US
YRSINUS	Numeric			Years in US
HHSIZE	Numeric			HH size
MOMONWIC	Numeric			Mom on WIC while pregnant
DOBOMOM	Numeric		MMDDYY8.	Date of birth--Mom
MOINCOM	Numeric	F12.		Family income
MOMSMK	Numeric			momsmk
DURATION	Numeric	F3.		Breastfeeding duration -- computed
MOLASTBF	Numeric			Last breastfeeding month
MOSTOPBF	Numeric			Month reported breastfeeding stopped
BRFDINIT	Numeric	F12.		Breastfeeding Initiated
FRSTDAYS	Numeric	F3.		Age of baby when formula started
SITE	Numeric	F12.		Site
ANALWGT1	Numeric	F12.		Analysis
STRATUM	Numeric	F12.		STRATUM
PSU	Numeric	F12.		PSU
BABYWGT	Numeric	F12.		Baby Weight (ozs.)
EDUC	Numeric	F12.		Education
BFEED	Numeric	F12.		Breastfeeding Initiation
MOMRACE	Numeric	F12.		momrac
AGE	Numeric			AGE
MRTLSTAT	Numeric	F12.		Marital Status
BRFDEND	Numeric	F12.		Breastfeeding stopped (0-1-.)
ORIGVAL	Numeric			Original Variable

This table shows the contents of the WSHD imputed dataset.

Exhibit 18. RECORDS Results: First Page of WSHD Imputed Dataset

WSHD Multiply Imputed Dataset

by: Observation Number.

Observation Number	ID	mom rac	mom smk	dnr1	dnr2	dnr3	dnr4	mom hos	hsp 1	hsp 2	hsp 3	hsp 4
1	4001	1	2	2	2	2	2
2	19136	1	2	2	2	2	2
3	21017	1	1	1	1	1	1
4	32019	1	1	1	1	1	1
5	40016	1	1	1	1	1	1
6	42125	1	4	4	4	4	4
7	1001	1	1	2	2	2	2	2
8	1009	1	1	2	2	2	2	2
9	1018	1	1	2	2	2	2	2
10	1020	1	1	3	3	3	3	3
11	1021	1	1	3	3	3	3	3
12	1022	1	1	3	3	3	3	3
13	1023	1	1	13	13	13	13	13
14	3012	1	1	0	0	0	0	0
15	3128	1	1	1	1	1	1	1
16	4003	1	1	0	0	0	0	0
17	4008	1	1	1	1	1	1	1
18	4022	1	1	1	1	1	1	1
19	4026	1	1	4	4	4	4	4
20	4127	1	1	3	3	3	3	3
21	4130	1	1	1	1	1	1	1
22	4133	1	1	1	1	1	1	1
23	4135	1	1	5	5	5	5	5
24	6008	1	1	2	2	2	2	2
25	6009	1	1	2	2	2	2	2
26	7122	1	1	2	2	2	2	2
27	8131	1	1	1	1	1	1	1
28	10129	1	1	1	1	1	1	1
29	11013	1	1	12010	12014	8131	16016	.	3	1	1	2
30	11014	1	1	1	1	1	1	1
31	11123	1	1	2	2	2	2	2
32	12004	1	1	1	1	1	1	1
33	12005	1	1	2	2	2	2	2
34	12009	1	1	1	1	1	1	1
35	12010	1	1	3	3	3	3	3
36	12013	1	1	1	1	1	1	1
37	12014	1	1	1	1	1	1	1
38	12117	1	1	2	2	2	2	2
39	13005	1	1	2	2	2	2	2
40	13126	1	1	2	2	2	2	2
41	14005	1	1	1	1	1	1	1
42	14015	1	1	1	1	1	1	1

There are 953 records on the output dataset MULTIMP. This is the first page of the output dataset. It is sorted by MOMRACE and MOMSMK. DONORID is missing except when imputation takes place. On those records, DONORID is the ID of the donor record. So for ID=11013 (record 29), MOMHOSP is missing. The random donors (one for each round of imputation) from the same race and smoking behavior levels are DONORID=12010 (who contributed a value of MOMHOSP=3), 12014 (contributed MOMHOSP=1), 8131 (contributed MOMHOSP=1), and 16016 (contributed MOMHOSP=2).

Exhibit 19. SAS FREQ Results: See How MOMHOSP Is Imputed (Imputation #1)

MOMHOSP	mhosp_1	Frequency	Percent	Cumulative Frequency	Cumulative Percent
.	1	7	0.73	7	0.73
.	2	11	1.15	18	1.89
.	3	2	0.21	20	2.10
.	4	1	0.10	21	2.20
0	0	25	2.62	46	4.83
1	1	320	33.58	366	38.41
2	2	350	36.73	716	75.13
3	3	142	14.90	858	90.03
4	4	48	5.04	906	95.07
5	5	22	2.31	928	97.38
6	6	7	0.73	935	98.11
7	7	4	0.42	939	98.53
8	8	2	0.21	941	98.74
10	10	1	0.10	942	98.85
11	11	1	0.10	943	98.95
12	12	3	0.31	946	99.27
13	13	7	0.73	953	100.00

In Imputation Round #1, the 21 missing values for MOMHOSP were hot-deck imputed to either 1, 2, 3, or 4. All other values remained as is.

Exhibit 20. SAS FREQ Results: See How MOMHOSP Is Imputed (Imputation #2)

MOMHOSP	mhosp_2	Frequency	Percent	Cumulative Frequency	Cumulative Percent
.	1	8	0.84	8	0.84
.	2	8	0.84	16	1.68
.	3	3	0.31	19	1.99
.	5	1	0.10	20	2.10
.	7	1	0.10	21	2.20
0	0	25	2.62	46	4.83
1	1	320	33.58	366	38.41
2	2	350	36.73	716	75.13
3	3	142	14.90	858	90.03
4	4	48	5.04	906	95.07
5	5	22	2.31	928	97.38
6	6	7	0.73	935	98.11
7	7	4	0.42	939	98.53
8	8	2	0.21	941	98.74
10	10	1	0.10	942	98.85
11	11	1	0.10	943	98.95
12	12	3	0.31	946	99.27
13	13	7	0.73	953	100.00

In Imputation Round #2, the 21 missing values for MOMHOSP were hot-deck imputed to either 1, 2, 3, 5, or 7. All other values remained as is.

Exhibit 21. SAS FREQ Results: See How MOMHOSP Is Imputed (Imputation #3)

MOMHOSP	mhosp_3	Frequency	Percent	Cumulative Frequency	Cumulative Percent
.	1	5	0.52	5	0.52
.	2	8	0.84	13	1.36
.	3	4	0.42	17	1.78
.	4	3	0.31	20	2.10
.	5	1	0.10	21	2.20
0	0	25	2.62	46	4.83
1	1	320	33.58	366	38.41
2	2	350	36.73	716	75.13
3	3	142	14.90	858	90.03
4	4	48	5.04	906	95.07
5	5	22	2.31	928	97.38
6	6	7	0.73	935	98.11
7	7	4	0.42	939	98.53
8	8	2	0.21	941	98.74
10	10	1	0.10	942	98.85
11	11	1	0.10	943	98.95
12	12	3	0.31	946	99.27
13	13	7	0.73	953	100.00

In Imputation Round #3, the 21 missing values for MOMHOSP were hot-deck imputed to either 1, 2, 3, 4, or 5. All other values remained as is.

Exhibit 22. SAS FREQ Results: See How MOMHOSP Is Imputed (Imputation #4)

MOMHOSP	mhosp_4	Frequency	Percent	Cumulative Frequency	Cumulative Percent
.	0	1	0.10	1	0.10
.	1	5	0.52	6	0.63
.	2	10	1.05	16	1.68
.	3	2	0.21	18	1.89
.	4	2	0.21	20	2.10
.	7	1	0.10	21	2.20
0	0	25	2.62	46	4.83
1	1	320	33.58	366	38.41
2	2	350	36.73	716	75.13
3	3	142	14.90	858	90.03
4	4	48	5.04	906	95.07
5	5	22	2.31	928	97.38
6	6	7	0.73	935	98.11
7	7	4	0.42	939	98.53
8	8	2	0.21	941	98.74
10	10	1	0.10	942	98.85
11	11	1	0.10	943	98.95
12	12	3	0.31	946	99.27
13	13	7	0.73	953	100.00

In Imputation Round #4, the 21 missing values for MOMHOSP were hot-deck imputed to either 0, 1, 2, 3, 4, or 7. All other values remained as is.

One of the advantages of using WSHD imputation is that, in expectation, the distribution of item respondents is equivalent that generated from the imputation-completed data. The distribution of the imputed variables is printed using the FREQ procedure:

Exhibit 23. SAS FREQ Results: Distribution of MHOSP_1 (Imputation #1)

Imputation #1 for Imputed variable #1				
mhosp_1	Frequency	Percent	Cumulative Frequency	Cumulative Percent
0	25	2.62	25	2.62
1	327	34.31	352	36.94
2	361	37.88	713	74.82
3	144	15.11	857	89.93
4	49	5.14	906	95.07
5	22	2.31	928	97.38
6	7	0.73	935	98.11
7	4	0.42	939	98.53
8	2	0.21	941	98.74
10	1	0.10	942	98.85
11	1	0.10	943	98.95
12	3	0.31	946	99.27
13	7	0.73	953	100.00

Exhibit 24. SAS FREQ Results: Distribution of MHOSP_2 (Imputation #2)

Imputation #2 for Imputed variable #1				
mhosp_2	Frequency	Percent	Cumulative Frequency	Cumulative Percent
0	25	2.62	25	2.62
1	328	34.42	353	37.04
2	358	37.57	711	74.61
3	145	15.22	856	89.82
4	48	5.04	904	94.86
5	23	2.41	927	97.27
6	7	0.73	934	98.01
7	5	0.52	939	98.53
8	2	0.21	941	98.74
10	1	0.10	942	98.85
11	1	0.10	943	98.95
12	3	0.31	946	99.27
13	7	0.73	953	100.00

Exhibit 25. SAS FREQ Results: Distribution of MHOSP_3 (Imputation #3)

Imputation #3 for Imputed variable #1				
mhosp_3	Frequency	Percent	Cumulative Frequency	Cumulative Percent
0	25	2.62	25	2.62
1	325	34.10	350	36.73
2	358	37.57	708	74.29
3	146	15.32	854	89.61
4	51	5.35	905	94.96
5	23	2.41	928	97.38
6	7	0.73	935	98.11
7	4	0.42	939	98.53
8	2	0.21	941	98.74
10	1	0.10	942	98.85
11	1	0.10	943	98.95
12	3	0.31	946	99.27
13	7	0.73	953	100.00

Exhibit 26. SAS FREQ Results: Distribution of MHOSP_4 (Imputation #4)

Imputation #4 for Imputed variable #1				
mhosp_4	Frequency	Percent	Cumulative Frequency	Cumulative Percent
0	26	2.73	26	2.73
1	325	34.10	351	36.83
2	360	37.78	711	74.61
3	144	15.11	855	89.72
4	50	5.25	905	94.96
5	22	2.31	927	97.27
6	7	0.73	934	98.01
7	5	0.52	939	98.53
8	2	0.21	941	98.74
10	1	0.10	942	98.85
11	1	0.10	943	98.95
12	3	0.31	946	99.27
13	7	0.73	953	100.00

These results show that there are no missing values in the imputation variable for any of the 4 multiple imputations. The output also shows that the imputation-completed distributions for all imputations are similar to the item respondent distributions shown at the beginning of this example.