Imputation for Questionnaires and Scales

Proration (Averaging the Available Items)

Scale scores sum or average a set of questionnaire items measuring the same construct

When component items are missing, researchers often compute prorated scale scores scales by averaging the available item responses

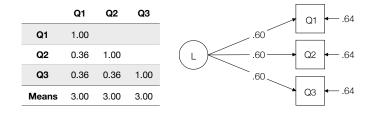
Equivalent to imputing values with a person's mean

Proration = Person-Mean Imputation

	Р	roratio	n		Р	erson-N	/lean Ir	nputa	tion
ID	Q1	Q2	Q3	Scale	ID	Q1	Q2	Q3	Scale
1	1	2	1	1.3	1	1	2	1	1.3
2	5	NA	4	4.5	2	5	4.5	4	4.5
3	3	2	4	3.0	3	3	2	4	3.0
4	NA	3	NA	3.0	4	3.0	3	3.0	3.0

Proration Assumptions

Proration requires two very strict assumptions: identical item means and inter-item correlations (parallel factor structure) and an MCAR mechanism



Scale-Level and Item-Level Imputation

Scale-level imputation: (a) compute scale scores, treating the scale as missing when one or more items is missing, then (b) impute the scale scores directly without the items

Item-level imputation: (a) impute the items, then (b) compute scale scores from the filled-in items

Item-level imputation offers a dramatic power gain

	Oriç	ginal	varial	Imputation variables				
ID	\mathbf{X}_1	X ₂	X 3	Y ₁	Y ₂	ID	ScaleX	ScaleY
1	1	2	1	NA	3	1	4	NA
2	5	NA	4	NA	NA	2	NA	NA
3	3	2	4	3	4	3	9	7
4	NA	3	NA	5	5	4	NA	10
200	4	5	4	3	4	200	13	7

Item-Level Imputation

Original variables					
ID	X ₁	X_2	X ₃	Y ₁	Y ₂
1	1	2	1	NA	3
2	5	NA	4	NA	NA
3	3	2	4	3	4
4	NA	3	NA	5	5
200	4	5	4	3	4

Motivating Example

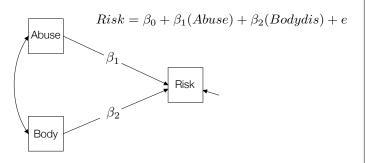
Questionnaire data from a study of eating disorder risk in a sample of 500 college-aged women

Variables include body mass index (BMI), questionnaire items measuring body dissatisfaction and eating disorder risk, past sexual abuse history (0 = no abuse history, 1 = abuse history)

All questionnaire items measured on a 7-point scale

Analysis Model

Body dissatisfaction and eating disorder risk are scale scores computed as the sum of the item responses



Analysis Summary

,												
	Ite	m-Level In	nputation									
Est. SE z p RIV												
B_0	10.04	0.88	11.36	< .001	0.04							
B ₁ (Abuse)	1.36	0.51	2.69	0.01	0.25							
B ₂ (Diss)	0.50	0.03	16.27	< .001	0.04							
Scale-Level Imputation												
Est. SE z p RIV												
B ₀	9.45	1.14	8.27	< .001	0.80							
B ₁ (Abuse)	1.70	0.60	2.81	< .01	0.98							
B ₂ (Diss)	0.52	0.04	12.66	< .001	0.86							

Important Takeaways

Item-level imputation offers a dramatic gain in precision

RIV = .86 (relative increase in variance) means that missing data error was 86% as large as complete-data sampling error, versus only 4% for item-level imputation

In order to obtain the same standard errors as item-level imputation, scale-level imputation would require a 40% to 90% increase in the sample size

Ex6.1.imp Blimp Diagnostic Script

DATA: ~/desktop/examples/eatingrisk.csv;

VARNAMES: id abuse bmi bds1-bds7 edr1-edr6;

NOMINAL: abuse;

ORDINAL: bds1-bds7 edr1-edr6;

MISSING: -99;

MODEL: ~ abuse bmi bds1-bds7 edr1-edr6;

SEED: 90291; BURN: 20000; THIN: 1; NIMPS: 2:

OUTFILE: ~/desktop/examples/imp*.csv;

OPTIONS: separate psr; CHAINS: 2 processors 2;

Diagnostic Output

Fix Eff	Ran Eff Var	Err Var	Threshold	
Max PSR	1.007	nan	1.000	1.055
Missing Variable	edr2		bmi	bds1
Comparing iterations 9301 to 18600 for 2 chains.				
Fix Eff	Ran Eff Var	Err Var	Threshold	
Max PSR	1.007	nan	1.000	1.046
Missing Variable	edr2		bmi	bds1

Ex6.2.imp Blimp Imputation Script (Mplus Format)

```
DATA: ~/desktop/examples/eatingrisk.csv;

VARNAMES: id abuse bmi bds1-bds7 edr1-edr6;

NOMINAL: abuse;

ORDINAL: bds1-bds7 edr1-edr6;

MISSING: -99;

MODEL: ~ abuse bmi bds1-bds7 edr1-edr6;

SEED: 90291;

BURN: 10000;

THIN: 10000;

NIMPS: 20;

OUTFILE: ~/desktop/examples/imp*.csv;

OPTIONS: separate;

CHAINS: 2 processors 2;
```

Blimp Output

```
VARIABLE ORDER IN SAVED DATA:

id abuse bmi bds1 bds2 bds3 bds4 bds5 bds6 bds7 edr1 edr2
edr3 edr4 edr5 edr6
```

Ex6.3.inp Mplus Analysis Script

```
DATA:

file = implist.csv;

type = imputation;

VARIABLE:

names = id abuse bmi bds1-bds7 edr1-edr6;

usevariables = abuse bodydis eatrisk;

DEFINE:

bodydis = sum(bds1-bds7);

eatrisk = sum(edr1-edr6);

MODEL:

eatrisk on abuse bodydis;

OUTPUT:

standardized(stdyx);
```

Mplus Analysis Output

MODEL RESULTS Two-Tailed Rate of S.E. Est./S.E. P-Value Estimate Missing EATRISK ON ABUSE 1.355 0.512 2.646 0.008 0.206 BODYDIS 16.188 0.500 0.031 0.000 0.047 Intercepts EATRISK 0.879 11.479 0.000 0.032 10.092 Residual Variances EATRISK 12.142 0.800 15.170 0.000 0.080

Ex6.4.imp Blimp Imputation Script (R, SAS, SPSS, and Stata Format)

```
DATA: ~/desktop/examples/eatingrisk.csv;
VARNAMES: id abuse bmi bds1-bds7 edr1-edr6;
NOMINAL: abuse;
ORDINAL: bds1-bds7 edr1-edr6;
MISSING: -99;
MODEL: ~ abuse bmi bds1-bds7 edr1-edr6;
SEED: 90291;
BURN: 10000;
THIN: 10000;
NIMPS: 20;
OUTFILE: ~/desktop/examples/imps.csv;
OPTIONS: stacked;
CHAINS: 2 processors 2;
```

Blimp Output

```
VARIABLE ORDER IN SAVED DATA:

imp# id abuse bmi bds1 bds2 bds3 bds4 bds5 bds6 bds7 edr1
edr2 edr3 edr4 edr5 edr6
```

Ex6.5.r R Analysis Script

```
# Required packages
library(mitml)

# Read data
impdata <- read.csv(file = "~/desktop/examples/
imps.csv",head = FALSE, sep = ",")
names(impdata) = c("imp","id","abuse","bmi",
    "bds1","bds2","bds3","bds4","bds5","bds6","bds7",
    "edr1","edr2","edr3","edr4","edr5","edr6")

# Compute scale scores
impdata$bodydis <- rowSums(subset(impdata,
    select = bds1:bds7))
impdata$eatrisk <- rowSums(subset(impdata,
    select = edr1:edr6))</pre>
```

Ex6.5.r R Analysis Script

```
# Analyze data and pool estimates
implist <- split(impdata, impdata$imp)
implist <- as.mitml.list(implist)
analysis <- with(implist, lm(eatrisk ~ abuse + bodydis))
testEstimates(analysis, df.com = 497)

# Test full model with Wald test
emptymodel <- with(implist, lm(eatrisk ~ 1))
testModels(analysis, emptymodel, method = "D1")</pre>
```

R Analysis Output

Final parameter estimates and inferences obtained from 20 imputed data sets.

	Estimate	Std.Error	t.value	df	P(> t)	RIV	FMI
(Intercept)	10.092	0.882	11.445	467.989	0.000	0.032	0.031
abuse	1.355	0.513	2.640	213.738	0.009	0.253	0.205
bodydis	0.500	0.031	16.141	447.769	0.000	0.049	0.047

Model comparison calculated from 20 imputed data sets. Combination method: D1

F.value df1 df2 P(>F) RIV 155.593 2 2010.888 0.000 0.142

Unadjusted hypothesis test as appropriate in larger samples.

Ex6.6.sps SPSS Analysis Script

data list free file = '/users/craig/desktop/examples/imps.csv' /imputation_ id abuse bmi bds1 to bds7 edr1 to edr6. exe.

- * Compute scale scores compute bodydis = sum(bds1 to bds7). compute eatrisk = sum(edr1 to edr6).
- * Initiate pooling routines. sort cases by imputation_. split file layered by imputation_.
- * Analysis and pooling.
 regression
 /descriptives mean stddev corr sig n
 /dependent eatrisk
 /method=enter abuse bodydis.

SPSS Analysis Output

						Coefficients ^a						
				Unstandardize	d Coefficients	Standardized Coefficients						
	imputation_	Model		В	Std. Error	Beta	t	Sig.				
ı	1.00	1	(Constant)	10.041	.855		11.738	.000				
			abuse	1.303	.452	.104	2.883	.004				
			bodydis	.503	.030	.609	16.885	.000				
ì	20.00	1	(6	0.040	052		11.672	000				

20.00	1	(Constant)	9.940	.852		11.673	.000
		abuse	1.115	.453	.088	2.461	.014
		bodydis	.507	.030	.615	17.108	.000
Pooled	1	(Constant)	10.092	.882		11.445	.000
		abuse	1.355	.513		2.640	.009
		bodydis	.500	.031		16.141	.000

Ex6.7.do Stata Analysis Script

```
// Import and save original data
import delimited "~/desktop/examples/eatingrisk.csv"
rename (v1 - v16)(id abuse bmi bds1 bds2 bds3 bds4 bds5 bds6 bds7 edr1
edr2 edr3 edr4 edr5 edr6)
generate imp = 0

// Recode missing values
foreach var of varlist id - edr6 {
    replace 'var' = . if 'var'== -99
}
save original, replace

// Import and save imputed data
clear
import delimited "~/desktop/examples/imps.csv"
rename (v1 - v17)(imp id abuse bmi bds1 bds2 bds3 bds4 bds5 bds6 bds7
edr1 edr2 edr3 edr4 edr5 edr6)
save imputed, replace
```

Stata Analysis Output

Ex6.7.do Stata Analysis Script

```
// Append original and imputed data
use original, clear
append using imputed

// Compute scale scores
generate bodydis = bds1 + bds2 + bds3 + bds4 + bds5 + bds6 + bds7
generate eatrisk = edr1 + edr2 + edr3 + edr4 + edr5 + edr6

// Convert to mi data
mi import flong, m(imp) id(id) imputed(abuse bmi bds1 - bds7
edr1 - edr6 bodydis eatrisk) clear

// Analyze data and pool results
mi estimate, cmdok: regress eatrisk abuse bodydis
```