

Project: Diagnostics for Informative Censoring (2015-0617)
Authors: Jackson JW, Schnellinger E, Valeri L, Henderson D.

Objective Obtaining intention-to-treat (ITT) effect estimates on symptom reduction and tolerability requires that investigators impute the missing outcomes^{1,2}. We applied recently developed data visualizations from the causal inference literature¹⁷ that describe, over time, how dynamic characteristics of patients who dropout differ from those who remain to a placebo-controlled efficacy trial of paliperidone extended release (NCT00397033)²⁰. Our objectives were to: (a) describe the dropout mechanism; (b) construct weights that would enable either an ITT comparison of final PANSS scores; (c) use data visualization tools to assess the quality of those weights; (d) report the results before and after applying the weights.

Methods We used visit-specific data to construct a person-time file (target days 0, 4, 8, 15, 22, 29, and 43) which included the assigned treatment arm and data on baseline and time-varying characteristics. Study discontinuation was indexed at the last visit where data was provided; thus, patients providing data at all visits were considered as having completed the study. We excluded the 0-4% those who did not initiate their first study medication and censored patients when they ended double-blind treatment. From a list of potential covariates, we chose those that were independently associated with the outcome or study discontinuation using linear and logistic regression (see Table 1). Fully conditional specification^{31,32} was used to impute the relatively minor missing data due to non-response (up to 2% for each covariate in both datasets) using PROC MI in SAS. Generalized additive models were fit on the imputed data to obtain the predicted probability of assigned treatment arm, with and without baseline covariates as predictors, with cubic splines to smooth predictions from continuous covariates. Generalized additive models were also fit on the imputed data to obtain the predicted probability of study discontinuation at visit *m*. We used these models to construct inverse probability of censoring weights (IPCW) as described in Cole and Hernan 2008.²⁴

We plotted the standardized mean difference of each time-varying covariate comparing the censored to uncensored at each visit, both before and after applying the weights to the data. We used GEE to fit unweighted and weighted linear regression models for change in PANSS score from baseline through day 43, both before and after weighting, again assuming an independence covariance matrix. In both applications, a robust-variance sandwich estimator was used to obtain the standard errors of model parameters.²⁶ P-values from multiple comparisons were adjusted by the Hochberg procedure. All data management and analysis was carried out SAS version 9.3 via the SAS Drug Development Clinical Trial Drug Transparency Platform. All plots were produced using ggplot2 in R. This study was conducted under the approval of the Institutional Review Board at the Harvard T.H. Chan School of Public Health, following all relevant guidelines and protocols.

Results

We obtained 307 patients who were randomized and initiated their assigned treatment. With the exception of race, the distribution of characteristics at baseline was similar (Table 1). There were 99 patients who ended double blind treatment early (of whom 9 provided data at the end of the study but we censored). As expected, dropout was lowest in the high-dose arm (22%), highest in the placebo arm (41%), and was most severe during the first two weeks of follow-up, peaking at target day 15 (Figure 1). The most common reported reason for dropout was lack of efficacy (26% to 50%). These numbers are largely consistent with the lead publication from this study.²⁰

The associations between dropout and time-varying symptoms were quite strong (Figure 1). For example, on target day 15 where most of the dropout occurred, many mean differences in PANSS score decline and CGI-Severity were more than a standard deviation apart for the placebo and low-dose arms, and ranged from one to three fifths of a standard deviation for the high-dose arm. Depression was also strongly

associated with dropout. These patterns of higher illness among the censored were consistent over the course of the trial. We found that applying the weights did not resolve these differences and in some instances exacerbated them. This is consistent with the weights' distribution having means farther from one among the placebo and low-dose arms (Figure 2). Concerningly, the treatment effects for the high-dose vs. placebo arms became more pronounced after weighting, from -9.1 (95%CI -14.2, -4.1) before weighting to -11.9 (95%CI -17.8, -6.0) after weighting; likewise, the treatment effect in the low-dose arm increased from -4.0 (95%CI -9.3, -1.4) to -9.2 (95%CI -15.1, -3.2). Examining the covariate distribution post-weighting informs us that the changes in effect size are likely not due to bias reduction and should not be taken seriously. For these reasons, we only present the unweighted change in PANSS score in Figure 4. The weights' poor performance is likely due to finite sample and related issues where patients who dropout lack similar counterparts among the uncensored data who can stand in for them.

Conclusions

We found the diagnostics to be instrumental in carrying out an objective analysis and were extremely useful in building and refining models for the dropout mechanism. As we expected to find, dropout was indeed informative of markers of treatment effect. As part of this project we developed freely available SAS macros that can be used to implement the visualizations reported here, available on the lead author's GitHub page. Example code for using these tools is provided in the Appendix.

References

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Table 1. Baseline characteristics and disposition in the paliperidone ER study			
	Placebo	Low-dose	High-dose
	N=107 (100)	N=104 (100)	N=96 (100)
Age	37.1 (10.8)	38 (9.6)	36.5 (10.3)
Gender			
Male	67 (63)	67 (64)	63 (66)
Female	40 (37)	37 (36)	33 (34)
Race/Ethnicity			
White	53 (50)	45 (43)	42 (44)
Asian	34 (32)	34 (33)	33 (34)
Black	20 (19)	23 (22)	20 (21)
Other	0 (0)	2 (2)	1 (1)
Randomization Strata			
Receiving mood stabilizer	41 (38)	38 (37)	38 (40)
Country			
Asia	33 (31)	34 (33)	30 (32)
Europe and other	35 (33)	31 (30)	31 (32)
North America	39 (36)	39 (38)	35 (36)
Baseline measures			
Total PANSS	91.6 (12.5)	96.0 (13.1)	93.3 (13.0)
CGI-Severity	4.6 (0.6)	4.6 (0.6)	4.6 (0.6)
YRMS	18.1 (6.8)	17.8 (7.4)	18.5 (8.2)
HAMD	15.7 (7.9)	17.6 (8.5)	17.3 (8.9)
Simpson-Agnes EPS	0.1 (0.2)	0.1 (0.3)	0.1 (0.3)
Weight	174.8 (44.7)	162.8 (39.7)	171.5 (49.1)
Disposition			
Remained in study	62 (63)	74 (71)	79 (76)
Ended double-blind treatment	44 (41)	34 (33)	34 (22)
... due to adverse event	7 (16)	9 (26)	4 (12)
... due to lack of efficacy	22 (50)	12 (35)	9 (26)
... due to loss to follow-up	3 (7)	3 (9)	2 (6)
... due to other reasons	0 (0)	4 (12)	1 (3)
... due to withdrawal of consent	12 (27)	6 (18)	5 (15)
<p>Mean (sd) for continuous, N (%) for categorical variables Acronyms: Positive and Negative Symptom Syndrome Scale (PANSS); Clinical Global Impressions Severity (CGI-Severity); Youth Mania Rating Scale (YRMS); Hamilton Depression Rating Scale (HAMD); Extrapyramidal Symptoms (EPS) PANSS, CGI-Severity, YRMS, and HAMD were measured at target days 0, 4, 8, 15, 22, 29, 43 and when double blind treatment ended early Simpson-Agnes EPS score was only measured at baseline and target day 15, therefore only baseline values were used. The score was taken as the mean of the tremor, fixation, elbow, shoulder, arm, and gait items to match the CATIE analysis Percentages for reasons for ending double blind treatment sum to 100 among those who did so</p>			

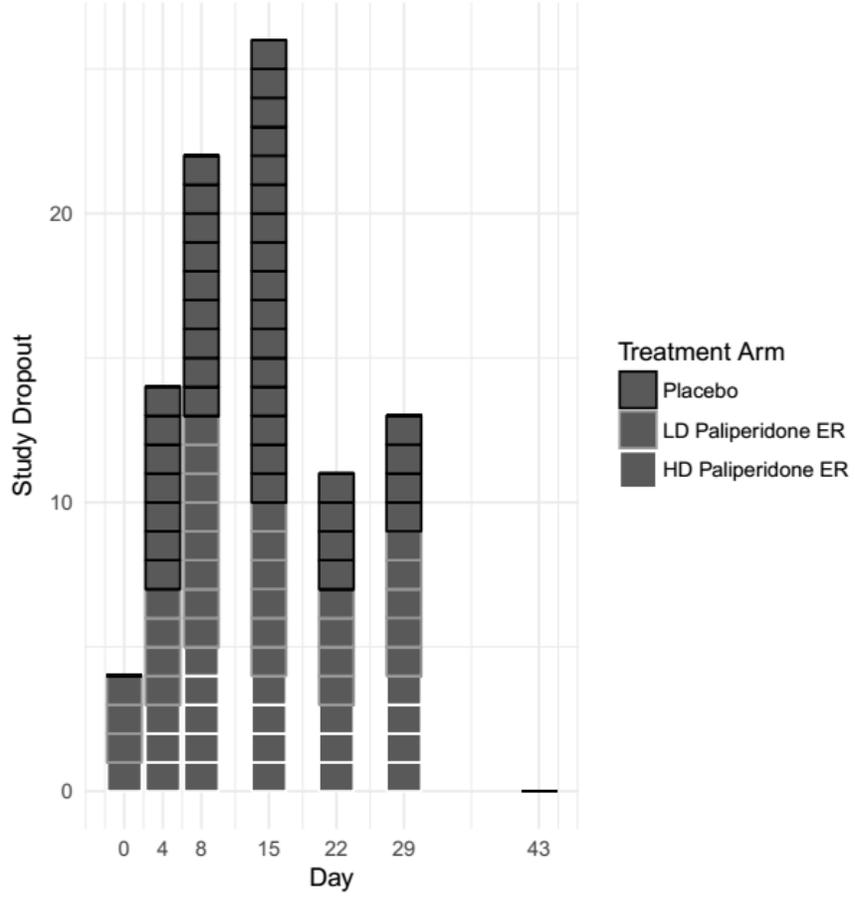


Figure 1. Study Dropout over the course of the Paliperidone ER study, by treatment arm

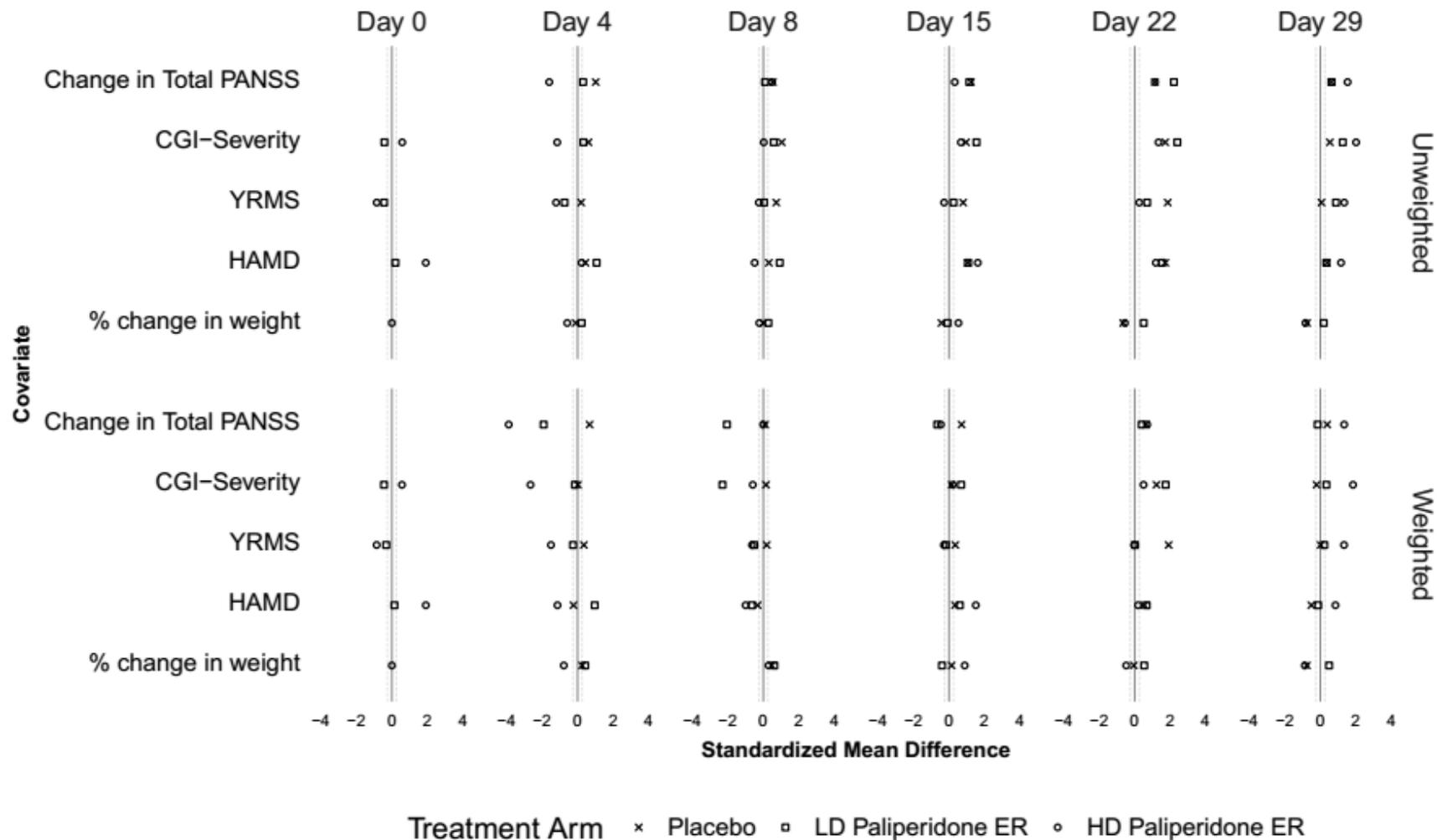


Figure 2. Mean difference of time-varying covariates comparing the censored vs. the uncensored over the course of the Paliperidone ER study, divided by the standard deviation among the uncensored, before and after application of IPCW weights. Colors distinguish the standardized mean differences for specific treatment arms. The dashed grey reference lines are placed at a positive and negative quarter of a standardized mean difference.

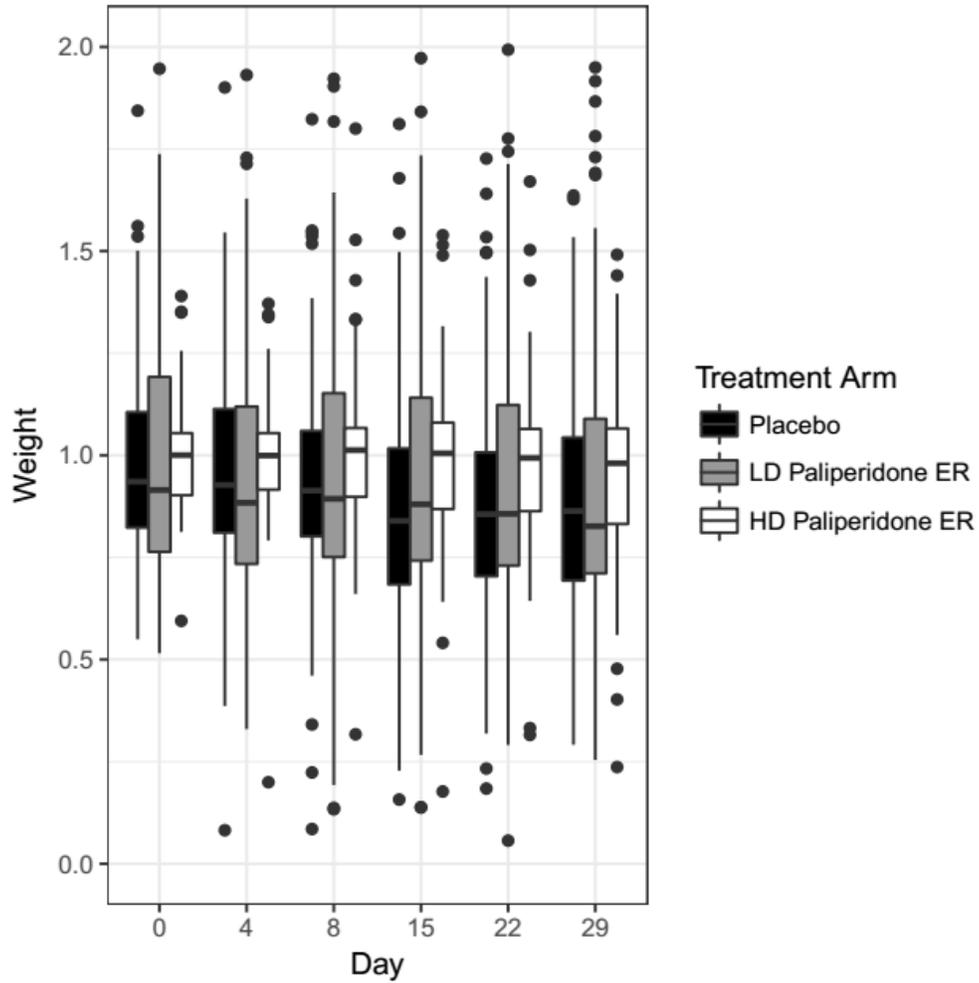


Figure 3. Boxplots depicting the distribution of the stabilized IPCW weights over the course of the study, by treatment arm. Though not shown, the full range of the weights was [0.06, 32.4] indicating very influential observations. The weights for the low-dose and placebo arms substantially diverge from one over time.

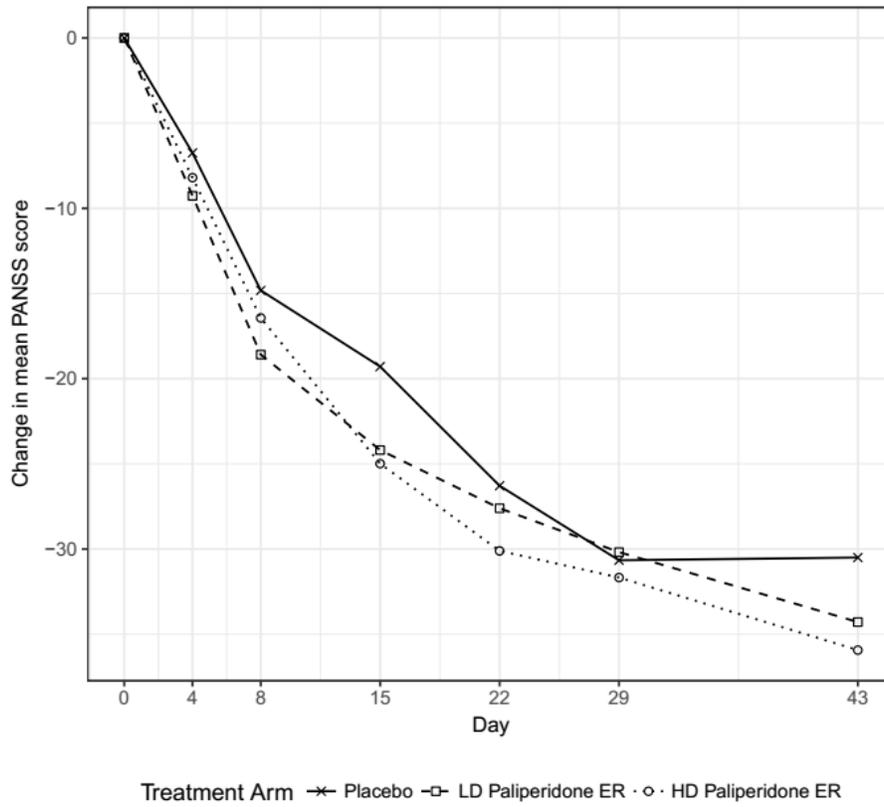


Figure 4. Change in Total PANSS score over the course of the Paliperidone ER study, by treatment arm.

Appendix

Weight specification

$$W(t) = \frac{P[A(0) = a|V]}{P[A(0) = a|C(0), V]} \prod_{m=0}^t \frac{P[S(m+1) = 0|\bar{S}(m) = 0, A, V]}{P[S(m+1) = 0|C(m), \bar{S}(m) = 0, A, V]}$$

where t represents a specific follow-up visit, m represents a visit equal to or less than t , $S(m+1)$ represents dropping out of the study immediately after visit m ($0 =$ remains at visit $m+1$, $1 =$ drops out by visit $m+1$), $\bar{S}(m) = 0$ represents no censoring up through month m , $C(m)$ represents covariates measured at visit m , while A and V respectively represent assigned treatment and covariates at baseline.

Example SAS code

Note: The following SAS macros are available at the lead author's faculty page through a link to a [GitHub repository](#). A software manual detailing the functions is provided, along with a toy dataset.

```
%lengthen(input=YodaWide,
output=YodaLong,
save=MyOtherDatasets,
diagnostic=3,
censoring=no,
id=yodaaid,
exposure=studydisc,
times_exposure=0 4 8 15 22 29,
times_covariate=0 4 8 15 22 29,
weight_exposure=cumw,
temporal_covariate = Chgpanssttotal hamdttotal yrmsttotal cgisev pctchgweight,
history=treatgrp);
```

```
%balance(input=YodaLong,
save=YodaWide MyOtherDatasets,
output=YodaTable,
diagnostic=3,
approach=weight,
censoring=no,
scope=recent,
recency=0,
exposure=studydisc,
history=treatgrp,
weight_exposure=cumw,
times_exposure = 0 4 8 15 22 29,
times_covariate = 0 4 8 15 22 29,
sort_order=alphabetical);
```

```
%makeplot(input=YodaTable,
save=YodaLong YodaWide MyOtherDatasets,
Output=YodaPlot,
approach=weight,
metric=SMD,
scope=recent)
```