

# Final Paper Analyses

*Marilyn Piccirillo*

*12/12/2017*

```
## Loading tidyverse: ggplot2
## Loading tidyverse: tibble
## Loading tidyverse: tidyr
## Loading tidyverse: readr
## Loading tidyverse: purrr
## Loading tidyverse: dplyr

## Warning: package 'tidyr' was built under R version 3.4.2
## Warning: package 'purrr' was built under R version 3.4.2
## Warning: package 'dplyr' was built under R version 3.4.2

## Conflicts with tidy packages -----
## %>%(): ggplot2, psych
## alpha(): ggplot2, psych
## filter(): dplyr, stats
## lag(): dplyr, stats

##
## Attaching package: 'lubridate'

## The following object is masked from 'package:base':
##
##   date

## Loading required package: Matrix

##
## Attaching package: 'Matrix'

## The following object is masked from 'package:tidyr':
##
##   expand

## This is lavaan 0.5-23.1097
## lavaan is BETA software! Please report any bugs.

##
## Attaching package: 'lavaan'

## The following object is masked from 'package:psych':
##
##   cor2cov

## Warning: package 'sjstats' was built under R version 3.4.2

## Warning in checkMatrixPackageVersion(): Package version inconsistency detected.
## TMB was built with Matrix version 1.2.10
## Current Matrix version is 1.2.11
## Please re-install 'TMB' from source or restore original 'Matrix' package

##
## Attaching package: 'sjstats'
```

```
## The following object is masked from 'package:broom':
##
##   bootstrap
## The following object is masked from 'package:psych':
##
##   phi
## Warning: package 'sjPlot' was built under R version 3.4.2
## Warning in read.spss("StressStudy_T1short_rename.sav", use.value.labels
## = FALSE, : StressStudy_T1short_rename.sav: Unrecognized record type 7,
## subtype 10 encountered in system file
## Warning in read.spss("StressStudy_T1short_rename.sav", use.value.labels =
## FALSE, : StressStudy_T1short_rename.sav: Long string missing values record
## found (record type 7, subtype 22), but ignored
## Warning in read.spss("StressStudy_T2short_rename.sav", use.value.labels
## = FALSE, : StressStudy_T2short_rename.sav: Unrecognized record type 7,
## subtype 10 encountered in system file
## Warning in read.spss("StressStudy_T2short_rename.sav", use.value.labels =
## FALSE, : StressStudy_T2short_rename.sav: Long string missing values record
## found (record type 7, subtype 22), but ignored
## Warning in read.spss("StressStudy_T3short_rename.sav", use.value.labels
## = FALSE, : StressStudy_T3short_rename.sav: Unrecognized record type 7,
## subtype 10 encountered in system file
## Warning in read.spss("StressStudy_T3short_rename.sav", use.value.labels =
## FALSE, : StressStudy_T3short_rename.sav: Long string missing values record
## found (record type 7, subtype 22), but ignored
```

Testing a longitudinal path model to replicate the model proposed by Davidson et al. (2011). They examined how interpersonal needs (Percieved burdensomness and thwarted belongingness) mediated the relationship of depression and social anxiety predicting suicidality. The issue with their model is that they used cross-sectional data. I'd like to replicate their model using longitudindial data. There weren't enough people who indicated symptoms of suicidality, so I will use the Acquired Capability for Suicide Scale, which assesses constructs such as fearlessness towards death. ACSS scores are highly correlated with suicidality.

I will test the following mediation relationships:

Depression → Percieved Burdensomness, Thwarted Belongingness → Acquired Capability for Suicide  
 Social Anxiety → Percieved Burdensomness, Thwarted Belongingness → Acquired Capability for Suicide

I expect that the mediation effect will be stronger for Depression → Percieved burdensomness → ACSS and that the mediation effect will be stronger for Social Anxiety → Thwarted Belongingness → ACSS

```
medDep <- '
#Correlate residuals
T1_BDI ~~ T2_BDI
T2_BDI ~~ T3_BDI
T1_Belong ~~ T2_Belong
T2_Belong ~~ T3_Belong
T1_Burden ~~ T2_Burden
```

```

T2_Burden ~~ T3_Burden

T1_ACSS ~~ T2_ACSS
T2_ACSS ~~ T3_ACSS

#Regressions to calculate indirect effects

T1_Burden ~ a1*T1_BDI + c1*T1_ACSS
T1_Belong ~ b1*T1_BDI + d1*T1_ACSS

T2_Belong ~ a1*T1_BDI + c2*T2_ACSS + g1*T1_Belong
T2_Burden ~ b1*T1_BDI + d2*T2_ACSS + h1*T2_Burden

T3_Belong ~ a2*T2_BDI + c3*T3_ACSS + g2*T2_Belong
T3_Burden ~ b2*T2_BDI + d3*T3_ACSS + h2*T2_Burden

T2_ACSS ~ i1*T1_ACSS + e1*T1_BDI + c1*T1_Belong + d1*T1_Burden
T3_ACSS ~ i2*T2_ACSS + e1*T1_BDI + c2*T2_Belong + d2*T2_Burden

#BDI --> Belong (a)
#BDI --> Burden (b)
#Belong --> ACSS (c)
#Burden --> ACSS (d)
#BDI --> ACSS (e)
#AR: BDI--> BDI (f)
#AR: Belong --> Belong (g)
#AR: Burden --> Burden (h)
#AR: ACSS --> ACSS (i)

ind:= a1*i2*c2*g1*g2 + b1*i2*d2*h1*h2
total:= ind + e1 + i2 '

fit.medDep <- sem(medDep, data = Depressionshort, missing = "ml")

## Warning in lavaan::lavaan(model = medDep, data = Depressionshort, missing =
## "ml", : lavaan WARNING: model has NOT converged!

summary(fit.medDep, standardized = TRUE, fit.measures = TRUE)

## ** WARNING ** lavaan (0.5-23.1097) did NOT converge after 114 iterations
## ** WARNING ** Estimates below are most likely unreliable
##
##      Number of observations                553
##
##      Number of missing patterns              8
##
##      Estimator                            ML
##      Minimum Function Test Statistic        NA
##      Degrees of freedom                     NA
##      P-value                               NA

## Warning in .local(object, ...): lavaan WARNING: fit measures not available if model did not converge
##
## Parameter Estimates:

```

```

##
## Information
## Standard Errors
##
## Regressions:
## Estimate Std.Err z-value P(>|z|) Std.lv Std.all
## T1_Burden ~
## T1_BDI (a1) 0.317 NA 0.317 0.464
## T1_ACSS (c1) -0.001 NA -0.001 -0.000
## T1_Belong ~
## T1_BDI (b1) 0.337 NA 0.337 0.395
## T1_ACSS (d1) -0.000 NA -0.000 -0.000
## T2_Belong ~
## T1_BDI (a1) 0.317 NA 0.317 0.360
## T2_ACSS (c2) -0.033 NA -0.033 -0.014
## T1_Belong (g1) 0.441 NA 0.441 0.427
## T2_Burden ~
## T1_BDI (b1) 0.337 NA 0.337 0.576
## T2_ACSS (d2) 0.195 NA 0.195 0.120
## T2_Burden (h1) -0.549 NA -0.549 -0.549
## T3_Belong ~
## T2_BDI (a2) 0.099 NA 0.099 0.096
## T3_ACSS (c3) -0.098 NA -0.098 -0.036
## T2_Belong (g2) 0.693 NA 0.693 0.666
## T3_Burden ~
## T2_BDI (b2) 0.049 NA 0.049 0.102
## T3_ACSS (d3) 0.205 NA 0.205 0.163
## T2_Burden (h2) 0.462 NA 0.462 0.636
## T2_ACSS ~
## T1_ACSS (i1) 0.554 NA 0.554 0.566
## T1_BDI (e1) 0.024 NA 0.024 0.066
## T1_Belong (c1) -0.001 NA -0.001 -0.001
## T1_Burden (d1) -0.000 NA -0.000 -0.000
## T3_ACSS ~
## T2_ACSS (i2) 0.574 NA 0.574 0.609
## T1_BDI (e1) 0.024 NA 0.024 0.070
## T2_Belong (c2) -0.033 NA -0.033 -0.087
##
## Covariances:
## Estimate Std.Err z-value P(>|z|) Std.lv Std.all
## T1_BDI ~~
## T2_BDI 0.107 NA 0.107 0.001
## T2_BDI ~~
## T3_BDI 0.072 NA 0.072 0.001
## .T1_Belong ~~
## .T2_Belong 0.029 NA 0.029 0.000
## .T2_Belong ~~
## .T3_Belong 0.007 NA 0.007 0.000
## .T1_Burden ~~
## .T2_Burden 0.244 NA 0.244 0.007
## .T2_Burden ~~
## .T3_Burden 0.069 NA 0.069 0.004
## .T2_ACSS ~~
## T1_ACSS 0.121 NA 0.121 0.010

```

```
##      .T3_ACSS          0.031      NA          0.031      0.003
##      .T3_Belong ~~
##      .T3_Burden        0.037      NA          0.037      0.002
##
## Intercepts:
##      Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##      .T1_Burden      7.028      NA          7.028      0.949
##      .T1_Belong     16.998      NA         16.998      1.838
##      .T2_Belong      7.579      NA          7.579      0.794
##      .T2_Burden      3.434      NA          3.434      0.541
##      .T3_Belong      5.569      NA          5.569      0.560
##      .T3_Burden      1.567      NA          1.567      0.340
##      .T2_ACSS        2.893      NA          2.893      0.742
##      .T3_ACSS        3.387      NA          3.387      0.921
##      T1_BDI         10.490      NA         10.490      0.966
##      T1_ACSS         8.244      NA          8.244      2.071
##      T2_BDI         8.008      NA          8.008      0.831
##      T3_BDI         5.689      NA          5.689      0.804
##
## Variances:
##      Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##      .T1_Burden     43.042      NA         43.042      0.784
##      .T1_Belong     72.143      NA         72.143      0.844
##      .T2_Belong     51.659      NA         51.659      0.566
##      .T2_Burden     26.000      NA         26.000      0.645
##      .T3_Belong     53.711      NA         53.711      0.544
##      .T3_Burden     11.359      NA         11.359      0.533
##      .T2_ACSS       10.142      NA         10.142      0.667
##      .T3_ACSS        8.353      NA          8.353      0.617
##      T1_BDI        117.914      NA        117.914      1.000
##      T1_ACSS       15.852      NA         15.852      1.000
##      T2_BDI       92.793      NA         92.793      1.000
##      T3_BDI       50.114      NA         50.114      1.000
##
## Defined Parameters:
##      Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##      ind       -0.011      NA        -0.011     -0.011
##      total        0.587      NA         0.587      0.664
```

```
medSIAS <- '
```

```
#Correlate residuals
```

```
T1_SIAS ~~ T2_SIAS
```

```
T2_SIAS ~~ T3_SIAS
```

```
T1_Belong ~~ T2_Belong
```

```
T2_Belong ~~ T3_Belong
```

```
T1_Burden ~~ T2_Burden
```

```
T2_Burden ~~ T3_Burden
```

```
T1_ACSS ~~ T2_ACSS
```

```

T2_ACSS ~~ T3_ACSS

#Regressions to calculate indirect effects

T1_Burden ~ a1*T1_SIAS + c1*T1_ACSS
T1_Belong ~ b1*T1_SIAS + d1*T1_ACSS

T2_Belong ~ a1*T1_SIAS + c2*T2_ACSS + g1*T1_Belong
T2_Burden ~ b1*T1_SIAS + d2*T2_ACSS + h1*T2_Burden

T3_Belong ~ a2*T2_SIAS + c3*T3_ACSS + g2*T2_Belong
T3_Burden ~ b2*T2_SIAS + d3*T3_ACSS + h2*T2_Burden

T2_ACSS ~ i1*T1_ACSS + e1*T1_SIAS + c1*T1_Belong + d1*T1_Burden
T3_ACSS ~ i2*T2_ACSS + e1*T1_SIAS + c2*T2_Belong + d2*T2_Belong

#SIAS --> Belong (a)
#SIAS --> Burden (b)
#Belong --> ACSS (c)
#Burden --> ACSS (d)
#SIAS --> ACSS (e)
#AR: SIAS--> SIAS (f)
#AR: Belong --> Belong (g)
#AR: Burden --> Burden (h)
#AR: ACSS --> ACSS (i)

ind:= a1*i2*c2*g1*g2 + b1*i2*d2*h1*h2
total:= ind + e1 + i2 '

fit.medSIAS <- sem(medSIAS, data = Depressionshort, missing = "ml")

## Warning in lavaan::lavaan(model = medSIAS, data = Depressionshort, missing
## = "ml", : lavaan WARNING: model has NOT converged!

summary(fit.medSIAS, standardized = TRUE, fit.measures = TRUE)

## ** WARNING ** lavaan (0.5-23.1097) did NOT converge after 10000 iterations
## ** WARNING ** Estimates below are most likely unreliable
##
##      Number of observations                553
##
##      Number of missing patterns              6
##
##      Estimator                             ML
##      Minimum Function Test Statistic        NA
##      Degrees of freedom                     NA
##      P-value                                NA

## Warning in .local(object, ...): lavaan WARNING: fit measures not available if model did not converge
##
## Parameter Estimates:
##
##      Information                Observed
##      Standard Errors            Standard

```

```

##
## Regressions:
##      Estimate  Std.Err  z-value  P(>|z|)  Std.lv  Std.all
##  T1_Burden ~
##    T1_SIAS  (a1)    0.161      NA          0.161    0.336
##    T1_ACSS  (c1)    0.013      NA          0.013    0.007
##  T1_Belong ~
##    T1_SIAS  (b1)    0.187      NA          0.187    0.299
##    T1_ACSS  (d1)    0.024      NA          0.024    0.010
##  T2_Belong ~
##    T1_SIAS  (a1)    0.161      NA          0.161    0.258
##    T2_ACSS  (c2)   -0.017      NA         -0.017   -0.007
##    T1_Belong (g1)    0.523      NA          0.523    0.524
##  T2_Burden ~
##    T1_SIAS  (b1)    0.187      NA          0.187    0.500
##    T2_ACSS  (d2)    0.290      NA          0.290    0.201
##    T2_Burden (h1)  -4.825      NA         -4.825   -4.825
##  T3_Belong ~
##    T2_SIAS  (a2)    0.193      NA          0.193    0.301
##    T3_ACSS  (c3)    0.049      NA          0.049    0.019
##    T2_Belong (g2)    0.617      NA          0.617    0.624
##  T3_Burden ~
##    T2_SIAS  (b2)    0.017      NA          0.017    0.055
##    T3_ACSS  (d3)    0.206      NA          0.206    0.169
##    T2_Burden (h2)    0.472      NA          0.472    0.603
##  T2_ACSS ~
##    T1_ACSS  (i1)    0.566      NA          0.566    0.579
##    T1_SIAS  (e1)   -0.017      NA         -0.017   -0.067
##    T1_Belong (c1)    0.013      NA          0.013    0.031
##    T1_Burden (d1)    0.024      NA          0.024    0.044
##  T3_ACSS ~
##    T2_ACSS  (i2)    0.769      NA          0.769    0.826
##    T1_SIAS  (e1)   -0.017      NA         -0.017   -0.072
##    T2_Belong (c2)   -0.017      NA         -0.017   -0.045
##
## Covariances:
##      Estimate  Std.Err  z-value  P(>|z|)  Std.lv  Std.all
##  T1_SIAS ~~
##    T2_SIAS      1.444      NA          1.444    0.007
##  T2_SIAS ~~
##    T3_SIAS      0.805      NA          0.805    0.004
##  .T1_Belong ~~
##    .T2_Belong    0.137      NA          0.137    0.002
##  .T2_Belong ~~
##    .T3_Belong   -0.395      NA         -0.395   -0.008
##  .T1_Burden ~~
##    .T2_Burden    7.078      NA          7.078    0.222
##  .T2_Burden ~~
##    .T3_Burden    0.434      NA          0.434    0.028
##  .T2_ACSS ~~
##    T1_ACSS      0.118      NA          0.118    0.010
##    .T3_ACSS     -2.813      NA         -2.813   -0.314
##  .T3_Belong ~~
##    .T3_Burden    0.545      NA          0.545    0.025

```

```
##
## Intercepts:
##           Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##   .T1_Burden      6.914    NA         6.914    0.966
##   .T1_Belong     16.463    NA        16.463    1.762
##   .T2_Belong      5.695    NA         5.695    0.611
##   .T2_Burden      2.367    NA         2.367    0.424
##   .T3_Belong      2.803    NA         2.803    0.304
##   .T3_Burden      1.503    NA         1.503    0.343
##   .T2_ACSS        2.931    NA         2.931    0.758
##   .T3_ACSS        2.277    NA         2.277    0.633
##   T1_SIAS        20.737    NA        20.737    1.387
##   T1_ACSS         8.244    NA         8.244    2.085
##   T2_SIAS        19.107    NA        19.107    1.331
##   T3_SIAS        17.460    NA        17.460    1.320
##
## Variances:
##           Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##   .T1_Burden     45.404    NA        45.404    0.887
##   .T1_Belong     79.534    NA        79.534    0.911
##   .T2_Belong     50.092    NA        50.092    0.576
##   .T2_Burden     22.352    NA        22.352    0.716
##   .T3_Belong     44.668    NA        44.668    0.526
##   .T3_Burden     10.889    NA        10.889    0.569
##   .T2_ACSS        9.727    NA         9.727    0.651
##   .T3_ACSS        8.234    NA         8.234    0.636
##   T1_SIAS       223.682    NA       223.682    1.000
##   T1_ACSS       15.638    NA        15.638    1.000
##   T2_SIAS       206.074    NA       206.074    1.000
##   T3_SIAS       174.850    NA       174.850    1.000
##
## Defined Parameters:
##           Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##   ind         -0.096         -0.096    -0.145
##   total         0.656         0.656     0.614

#Descriptive stats
table(DepressionMerge$Gender)

##
##   1  2  3  4
## 122 425  1  1

table(DepressionMerge$Ethnicity)

##
##   0  1  2  3  4  5
## 319 109 35 25 37 27

table(DepressionMerge$Hispanic)

##
##   1  2
## 42 511

describe(DepressionMerge$Age)
```



```
##      vars    n mean    sd median trimmed  mad min max range skew kurtosis   se
## X1      1 552 21.1 4.13      20   20.42 1.48  10  61    51 4.62    30.87 0.18
```

```
#Subset data to calculate alphas
```

```
T1BDI <- dplyr::select(DepressionMerge, T1_BDI1:T1_BDI21)
T2BDI <- dplyr::select(DepressionMerge, T2_BDI1:T2_BDI21)
T3BDI <- dplyr::select(DepressionMerge, T3_BDI1:T3_BDI21)
T1SIAS <- dplyr::select(DepressionMerge, T1_SIAS1:T1_SIAS4, T1_SIAS6:T1_SIAS8, T1_SIAS10, T1_SIAS12:T1_SIAS14, T1_SIAS16:T1_SIAS18, T1_SIAS20:T1_SIAS21)
T2SIAS <- dplyr::select(DepressionMerge, T2_SIAS1:T2_SIAS4, T2_SIAS6:T2_SIAS8, T2_SIAS10, T2_SIAS12:T2_SIAS14, T2_SIAS16:T2_SIAS18, T2_SIAS20:T2_SIAS21)
T3SIAS <- dplyr::select(DepressionMerge, T3_SIAS1:T3_SIAS4, T3_SIAS6:T3_SIAS8, T3_SIAS10, T3_SIAS12:T3_SIAS14, T3_SIAS16:T3_SIAS18, T3_SIAS20:T3_SIAS21)
T1ACSS <- dplyr::select(DepressionMerge, T1_ACSS1:T1_ACSS5)
T2ACSS <- dplyr::select(DepressionMerge, T2_ACSS1:T2_ACSS5)
T3ACSS <- dplyr::select(DepressionMerge, T3_ACSS1:T3_ACSS5)
T1Belong <- dplyr::select(DepressionMerge, T1_INQ7:T1_INQ15)
T2Belong <- dplyr::select(DepressionMerge, T2_INQ7:T2_INQ16)
T3Belong <- dplyr::select(DepressionMerge, T3_INQ7:T3_INQ16)
T1Burden <- dplyr::select(DepressionMerge, T1_INQ1:T1_INQ6)
T2Burden <- dplyr::select(DepressionMerge, T2_INQ1:T2_INQ6)
T3Burden <- dplyr::select(DepressionMerge, T3_INQ1:T3_INQ6)
```

```
alpha(T1BDI)
alpha(T2BDI)
alpha(T3BDI)
alpha(T1SIAS)
alpha(T2SIAS)
alpha(T3SIAS)
alpha(T1ACSS, check.keys = TRUE)
alpha(T2ACSS, check.keys = TRUE)
alpha(T3ACSS, check.keys = TRUE)
alpha(T1Burden)
alpha(T2Burden)
alpha(T3Burden)
alpha(T1Belong, check.keys = TRUE)
alpha(T2Belong, check.keys = TRUE)
alpha(T3Belong, check.keys = TRUE)
```