

COS-D419 Factor Analysis and Structural Equation Models 2023, Assignment 4

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1 SEM & teacher burnout

1.1 Exercise 4.1

Draw the graph of the initial, full structural equation model. Make sure that you have included all the specified paths.

Estimate the initial model using the robust MLM estimator (*robust variant of the ML estimator, to be precise!*) and present a brief summary of the model fit.

2 Preparation

2.1 Read in the data set

Start by downloading the data file from Moodle to Project folder.

```
library(tidyverse)
library(readr)
library(here)

#This week's file name
latest.name <- "ALLSEC.CSV"

#read in the data
mbi <- read_csv(file.path(here(),
                           'data',
                           latest.name))
```

2.2 Write functions

To control length of reports, codes already shown in the previous homework were not showing in the current report. Yet they are available in .rmd report.

2.2.1 to check unique values

2.2.2 to generate CFA results with improved readability

2.2.3 to generate functions for improving aesthetics of correlation matrix

2.2.4 to generate a function for histogram overlapping with density plot

2.2.5 to generate a function for violin overlapping with box plot

2.2.6 To generate a function describing continuous data set

2.2.7 Write a function to print a table with concerned parameters

```
#write a function for minus calculation
minus <- function(x,y) {x - y}

#write a function to print a table with concerned parameters
concern.table <- function(sem, nofpath, model){
  options(scipen = 999)
  #This is for structural path residual variance
  ##regression path estimates
  sem.parameter <- parameterEstimates(sem, standardized=TRUE) |> # obtain estimates
  filter(op == "~") |> #select "is measured by" rows
  mutate(Parameter = paste0(rhs, "→", lhs)) |>
  select(Parameter,
         'B'=est, #estimates
```

```

    'Beta'=std.all, #estimates standardized
    SE=se, #standard error
    Z=z, #z statistics
    'p-value'=pvalue #p value
  )

##round the p-value column
sem.parameter$`p-value` <- sem.parameter$`p-value` |>
  round(3)

##add a conditional logic to the p-value column that >0.05 cell shows in red
sem.parameter$`p-value` <- cell_spec(sem.parameter$`p-value`,
  color = ifelse(
    sem.parameter$`p-value` > 0.05,
    "red",
    "black")
  )

#This is for the residual variance of dependence variable
##obtain estimates
variance <- parameterEstimates(sem, standardized=TRUE) |>
  filter(op == "~~") #select "is correlated with" rows
##subset needed rows (variance row)
variance <- variance |>
  filter(rhs %in% c("F8SELF", "F9ELC", "F10EE", "F11DP", "F12PA"))

#variance[minus(sum(32,nofpath), 5-1):sum(32,nofpath),] #32 is the n of indicators;
#12 is the number of factors;
#5 is the new of row I plan to show

##select&rename columns
sem.tab.variance <- variance |> select(
  Parameter=rhs, #right hand side column
  'B'=est, #estimates
  'Beta'=std.all, #standardized estimates
  SE=se, #standard error
  Z=z, #z statistics
  'p-value'=pvalue #p value
)

##remove the row names
rownames(sem.tab.variance) <- NULL
##round the p-value column
sem.tab.variance$`B` <- sem.tab.variance$`B` |>
  round(3)
##add a conditional logic to the p-value column that >0.05 cell shows in red
sem.tab.variance$`B` <- cell_spec(sem.tab.variance$`B`,
  color = ifelse(
    sem.tab.variance$`B` < 0,
    "red",
    "black")
  )

#bind the two table

```

```

concern.table <- rbind(sem.parameter, sem.tab.variance)
concern.table[1:nofpath, 2] <-
  as.character(round(as.numeric(concern.table[1:nofpath, 2]),3))
concern.table[sum(nofpath, 1):sum(nofpath, 5),6] <-
  as.character(round(
    as.numeric(
      concern.table[sum(nofpath,1):sum(nofpath, 5),6]
    ),
    3)
  )

#further aesthetics
concern.table |>
  select("Parameter*" = Parameter,
    'B|' = B, #estimates
    'Beta|' = Beta, #estimates standardized
    SE, #standard error
    Z, #z statistics
    'p-value') |> #p value
kable(digits = 3, #rounded to 3
  #format="latex", #Latex markdown
  booktabs=TRUE, #Latex booktabs
  linesep = "",
  align = "lrrrrr",
  caption=
    paste(
      "Residual variance of structural regression path and select factors for",
      model),
    escape = F) |> #caption
kable_styling(latex_options = "striped") |> #gray every other row
pack_rows("Strutural regression path",
  1,nofpath) |>
pack_rows("Selected factors",
  sum(nofpath,1), sum(nofpath,5)) |>
footnote(general = "Values highlighted in red should be taken note of",
  symbol = c("> indicates regression path",
    "Crude estimates",
    "Standardized estimates"))
}

```

2.2.8 To generate a function for calculating chi square difference was defined.

```

chi.diff <- function(sem1, sem2){ #2 augments input: the 1st is the nested model
  #the 2nd is comparison model (less restricted)
  measure0 <- fitMeasures(sem1, #extract robust fit indices for the 1st model
    c("df.scaled",
      "chisq.scaling.factor",
      "chisq.scaled")) |>
  as.vector() #turn to vector to facilitate calculation
  measure1 <- fitMeasures(sem2, #extract robust fit indices for the 2nd model
    c("df.scaled",

```

```

                                "chisq.scaling.factor",
                                "chisq.scaled")) |>
  as.vector()
cd <- # this is the formula from p 14, week 4 slide
  (measure0[1]*measure0[2]-measure1[1]*measure1[2])/(measure0[1]-measure1[1])
TRd <- # this is the formula from p 15, week 4 slide
  (measure0[3]*measure0[2]-measure1[3]*measure1[2])/cd
TRd <- TRd |> round(3) # round the value to 3 decimal places
}

```

2.2.9 To generate a function for plotting full sem diagram

```

#plot the diagram
full.sem.diagram <- function(model, nofpath, fig.num, quotedmodel){
  semPaths(semPlotModel(model),
    style = "lisrel",
    rotation = 2,
    sizeLat = 6,
    sizeLat2 = 5,
    sizeMan = 5,
    sizeMan2 = 2,
    residScale = 4,
    shapeMan = "rectangle",
    edge.color = c(rep("black",34),
                    rep("blue", minus(nofpath,1)),
                    "orange", # this is the new free parameter
                    rep("gray",32),
                    rep("red",5)),

    residuals = T,
    layout = m,
    nCharNodes=0,
    optimizeLatRes = T,
    exoVar = F)
  #add title
  title(main = list(paste("Figure",
                          fig.num,
                          "Modified model (",
                          quotedmodel,
                          ") of teacher burnout"),
                  cex = 1.5, font =1),
    outer = F, line = -1)
  #add notes
  title(sub =
    "Notes: Orange arrow is the regression path set free to estimate in the current model",
    line = 0, adj = 0.7)
}

```

2.2.10 To generate model comparison tables.

```
###function start here

model.comparison.table <- function(sem, mycaption, newpathstext){
  sem.compare <- matrix(NA, ncol =8, nrow = length(sem))

  for (i in 1:length(sem)){

    sem.compare[i,] <- fitMeasures(eval(parse(text=sem[i])),#obtain specified measured.
                                   c("chisq.scaled",
                                       "df.scaled",
                                       "pvalue.scaled",
                                       "cfi.scaled",
                                       "tli.scaled",
                                       "rmsea.scaled",
                                       "srmr_bentler",
                                       "chisq.scaling.factor")) |>
    round(3)
  }

  colnames(sem.compare) <- c("chisq.scaled",
                              "df.scaled",
                              "pvalue.scaled",
                              "cfi.scaled",
                              "tli.scaled",
                              "rmsea.scaled",
                              "srmr_bentler",
                              "chisq.scaling.factor")

  sem.compare <- sem.compare |> data.frame()

  #add column "chi square" and place values into it
  sem.compare$"ΔChi-square" <- rep(NA, length(sem))
  for (j in 2:length(sem)){ #calculate chisq difference
    sem.compare$"ΔChi-square"[j] <- chi.diff(eval(parse(text=sem[j-1])),
                                              eval(parse(text=sem[j]))
                                              )
    )
  }

  #turn named vector to data frame
  sem.compare<- sem.compare |>
  select("Chi-square" = chisq.scaled, "DF"=df.scaled,
         "p value"=pvalue.scaled, "ΔChi-square", "CFI"= cfi.scaled,
         "TLI"= tli.scaled, "RMSEA"= rmsea.scaled,
         "SRMR"=srmr_bentler, "CSF"=chisq.scaling.factor)

  #There is no chisq-diff value for the first model, so place a "--" in the cell
  sem.compare[1,4] <- "--"
```

```

#add row names
symbols <- c("1", "2*", "3†", "4‡", "5§", "6¶", "7**", "8‡‡")
for (x in 1:length(sem)){
  rownames(sem.compare)[x] <- paste0("Model", symbols[x])
}

#table aesthetics fine-tune and print table

sem.compare |>
  kable(booktab =T,
        #format = "markdown",
        caption = mycaption,
        align = "r",
        linesep = "") |>
  kable_styling() |>
  footnote(symbol = newpathstext)
}

```

2.3 Inspect the data

2.3.1 Data structure

Have a quick overview of the data structure

```

library(knitr)
library(broom)
dim(mbi);mbi %>% apply(2, function(x)class(x));

```

```
## [1] 1430 32
```

```

##  ROLEA1  ROLEA2  ROLEC1  ROLEC2  WORK1  WORK2  CCLIM1  CCLIM2
## "numeric" "numeric" "numeric" "numeric" "numeric" "numeric" "numeric" "numeric"
##  CCLIM3  CCLIM4  DEC1  DEC2  SSUP1  SSUP2  PSUP1  PSUP2
## "numeric" "numeric" "numeric" "numeric" "numeric" "numeric" "numeric" "numeric"
##  SELF1  SELF2  SELF3  ELC1  ELC2  ELC3  ELC4  ELC5
## "numeric" "numeric" "numeric" "numeric" "numeric" "numeric" "numeric" "numeric"
##  EE1  EE2  EE3  DP1  DP2  PA1  PA2  PA3
## "numeric" "numeric" "numeric" "numeric" "numeric" "numeric" "numeric" "numeric"

```

The data set contains 22 numeric variables of 372 obs. Their values appear to follow a consistent pattern covering the integer from 1 to 7, except for Items 4, 7, 17 and 21, which did not include a value of 1.

2.3.2 Descriptive statistics of measured variables

```

library(finalfit);library(kableExtra);

descriptive(mbi) |>
  pack_rows(index =

```

```

c("Factor 1*: Role Ambiguity \n(high score means negative)" = 2,
  "Factor 2*: Role conflict \n(high score means negative)" = 2,
  "Factor 3*: Work overload \n(high score means negative)" = 2,
  "Factor 4‡: classroom climate" = 4,
  "Factor 5*: Decision-making" = 2,
  "Factor 6*: Superior support" = 2,
  "Factor 7*: Peer support" = 2,
  "Factor 8‡: Self-esteem" = 3,
  "Factor 9‡: External locus of control" = 5,
  "Factor 10‡: Emotional Exhaustion \n(high score means negative)" = 3,
  "Factor 11‡: Depersonalization \n(high score means negative)" = 2,
  "Factor 12‡: Personal Accomplishment" = 3)) |>
footnote(general =
  "Indicators variables were formulated through item parcels.",
symbol = c("These indicators are parcels from Teacher Stress Scale instrument",
  "These indicators are parcels from BMI instrument",
  "These parcels consist of items from single unidimensional scales")
)

```

Table 1: Descriptive statistics for measurements

	n	n of NA	Central tendency		Dispersion tendency			
			Mean	Median	SD	Min	Max	Q1~Q3
Factor 1*: Role Ambiguity (high score means negative)								
ROLEA1	1430	0	2.4	2.3	0.9	1.0	6.0	1.7 ~ 3.0
ROLEA2	1430	0	2.1	2.0	1.0	1.0	6.0	1.5 ~ 2.5
Factor 2*: Role conflict (high score means negative)								
ROLEC1	1430	0	3.0	3.0	1.1	1.0	6.0	2.3 ~ 3.7
ROLEC2	1430	0	3.0	3.0	1.2	1.0	6.0	2.0 ~ 4.0
Factor 3*: Work overload (high score means negative)								
WORK1	1430	0	3.2	3.3	1.2	1.0	6.0	2.3 ~ 4.0
WORK2	1430	0	2.2	2.0	1.1	1.0	6.0	1.5 ~ 3.0
Factor 4‡: classroom climate								
CCLIM1	1430	0	3.0	3.0	0.5	1.0	4.0	2.7 ~ 3.3
CCLIM2	1430	0	2.7	2.7	0.6	1.0	4.0	2.3 ~ 3.0
CCLIM3	1430	0	2.9	3.0	0.5	1.0	4.0	2.7 ~ 3.3
CCLIM4	1430	0	3.1	3.0	0.7	1.0	4.0	2.5 ~ 3.5
Factor 5*: Decision-making								
DEC1	1430	0	4.0	4.0	1.0	1.0	6.0	3.3 ~ 4.7
DEC2	1430	0	4.2	4.5	1.3	1.0	6.0	3.5 ~ 5.5
Factor 6*: Superior support								
SSUP1	1430	0	4.3	4.3	1.2	1.0	6.0	3.7 ~ 5.3
SSUP2	1430	0	4.4	4.5	1.3	1.0	6.0	3.5 ~ 5.5
Factor 7*: Peer support								
PSUP1	1430	0	4.6	4.7	1.0	1.0	6.0	4.0 ~ 5.3
PSUP2	1430	0	4.6	4.5	0.9	1.0	6.0	4.0 ~ 5.0
Factor 8‡: Self-esteem								

Table 1: Descriptive statistics for measurements (*continued*)

	n	n of NA	Central tendency		Dispersion tendency			
			Mean	Median	SD	Min	Max	Q1~Q3
SELF1	1430	0	3.6	3.7	0.4	1.0	4.0	3.3 ~ 4.0
SELF2	1430	0	3.6	3.8	0.5	1.0	4.0	3.4 ~ 4.0
SELF3	1430	0	3.5	3.7	0.5	1.0	4.0	3.3 ~ 4.0
Factor 9‡: External locus of control								
ELC1	1430	0	2.9	3.0	0.6	1.0	4.8	2.6 ~ 3.4
ELC2	1430	0	3.0	3.0	0.6	1.0	5.0	2.5 ~ 3.5
ELC3	1430	0	2.8	2.8	0.5	1.0	4.8	2.4 ~ 3.2
ELC4	1430	0	2.2	2.2	0.6	1.0	4.5	1.8 ~ 2.5
ELC5	1430	0	2.5	2.4	0.6	1.0	4.8	2.0 ~ 3.0
Factor 10†: Emotional Exhaustion (high score means negative)								
EE1	1430	0	3.9	4.0	1.3	1.0	7.0	3.0 ~ 4.7
EE2	1430	0	3.5	3.3	1.3	1.0	7.0	2.7 ~ 4.3
EE3	1430	0	3.2	3.0	1.3	1.0	7.0	2.0 ~ 4.0
Factor 11†: Depersonalization (high score means negative)								
DP1	1430	0	2.3	2.0	1.1	1.0	6.7	1.3 ~ 3.0
DP2	1430	0	2.1	1.5	1.2	1.0	7.0	1.0 ~ 2.5
Factor 12‡: Personal Accomplishment								
PA1	1430	0	5.7	6.0	0.9	2.0	7.0	5.3 ~ 6.3
PA2	1430	0	5.8	6.0	1.0	2.0	7.0	5.5 ~ 6.5
PA3	1430	0	5.8	6.0	1.0	2.0	7.0	5.3 ~ 6.7

Note:

Indicators variables were formulated through item parcels.

* These indicators are parcels from Teacher Stress Scale instrument

† These indicators are parcels from BMI instrument

‡ These parcels consist of items from single unidimensional scales

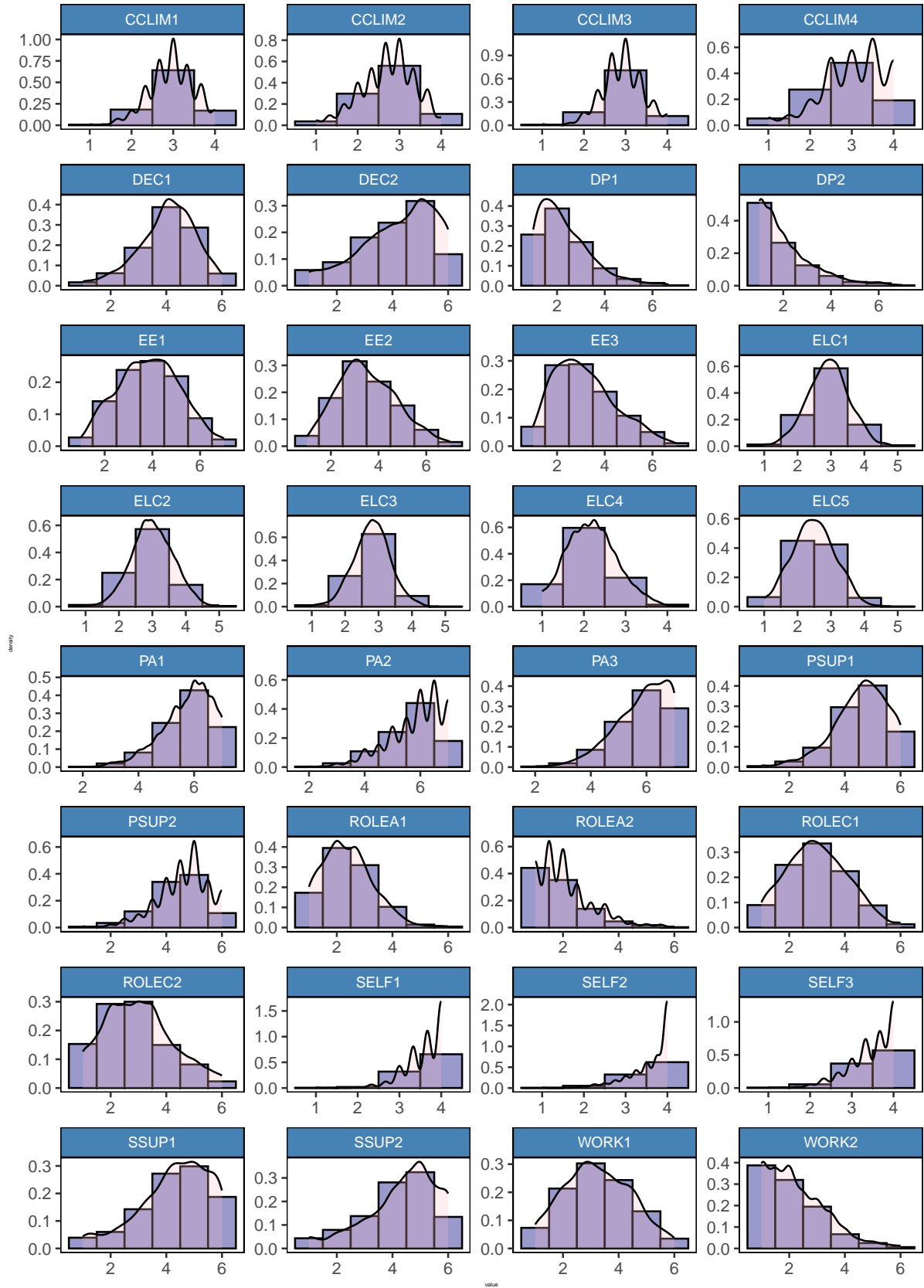
2.3.3 Visualization

(1) Histogram

Distribution of the data was examined via Histogram

```
corr.density(mbi, fig.num = 1)
```

Figure 1 Distribution of selected items



Ridge-line plots were generated for TSS and MBI indicators, respectively. By partially overlaying, it is a demonstration viable for comparing multiple distributions.

```
library(ggribes)
library(viridis)
library(hrbrthemes)
library(patchwork)
a <- mbi |>
  select(
    starts_with("EE")|starts_with("DP")|starts_with("PA")
  ) |>
  pivot_longer(everything(), names_to = "variable", values_to = "value") |>
  ggplot(aes(x = value, y = variable, fill = ..x..)) +
  geom_density_ridges_gradient(scale = 3, rel_min_height = 0.01) +
  scale_fill_viridis(name = "parcelled score", option = "C") +
  labs(title =
    'Fig2 (a). Distribution of indicator scores from BMI instrument') +
  labs(x = "Indicator scores", y = "Indicators") +
  theme(
    legend.position="none",
    panel.spacing = unit(0.1, "lines"),
    plot.title = element_text(size = 12),
    panel.grid.major = element_blank(),
    panel.background = element_rect(color = "black",
                                     fill = "white")
  )

b <- mbi |>
  select(
    starts_with("ROL")|starts_with("WOR")|starts_with("DEC")|contains("SUP")
  ) |>
  pivot_longer(everything(), names_to = "variable", values_to = "value") |>
  ggplot(aes(x = value, y = variable, fill = ..x..)) +
  geom_density_ridges_gradient(scale = 3, rel_min_height = 0.01) +
  scale_fill_viridis(name = "parcelled score", option = "C") +
  labs(title =
    'Fig2 (b). Distribution of indicator scores from TSS instrument') +
  labs(x = "Indicator scores", y = "Indicators")+
  theme(
    legend.position="none",
    panel.spacing = unit(0.1, "lines"),
    plot.title = element_text(size = 12),
    panel.grid.major = element_blank(),
    panel.background = element_rect(color = "black",
                                     fill = "white")
  )

a/b
```

Fig2 (a). Distribution of indicator scores from BMI instrument

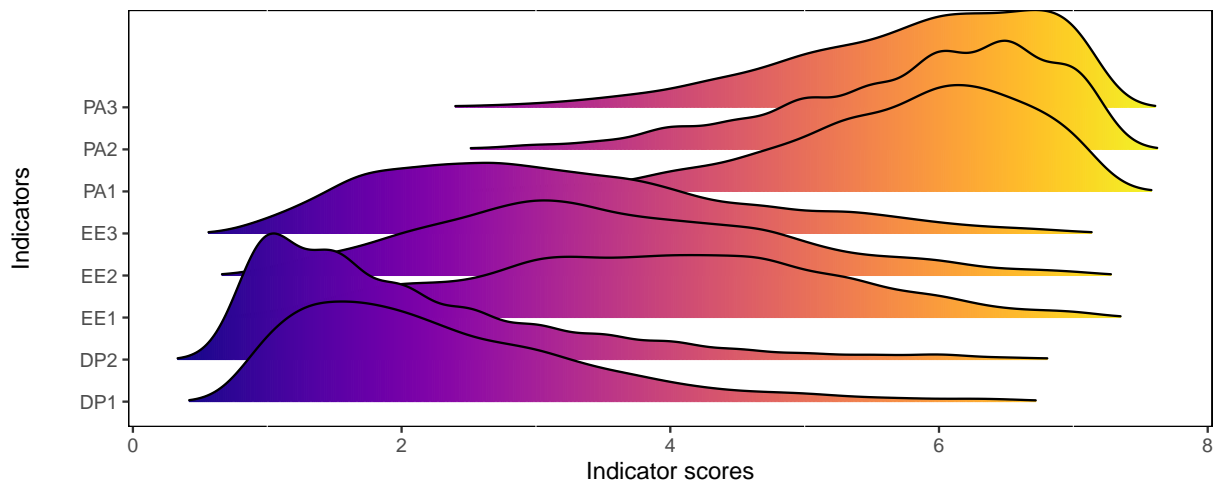
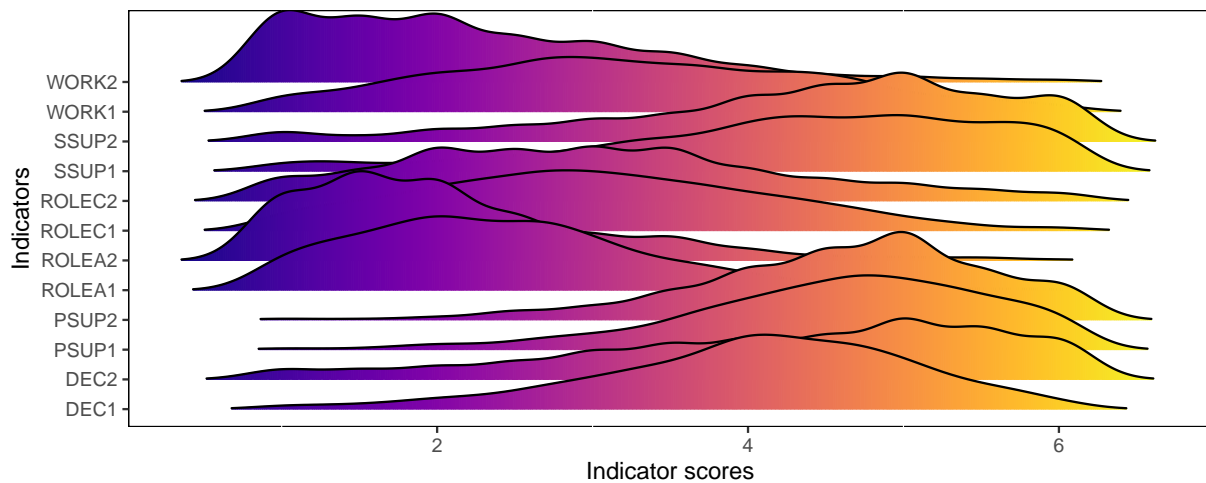


Fig2 (b). Distribution of indicator scores from TSS instrument



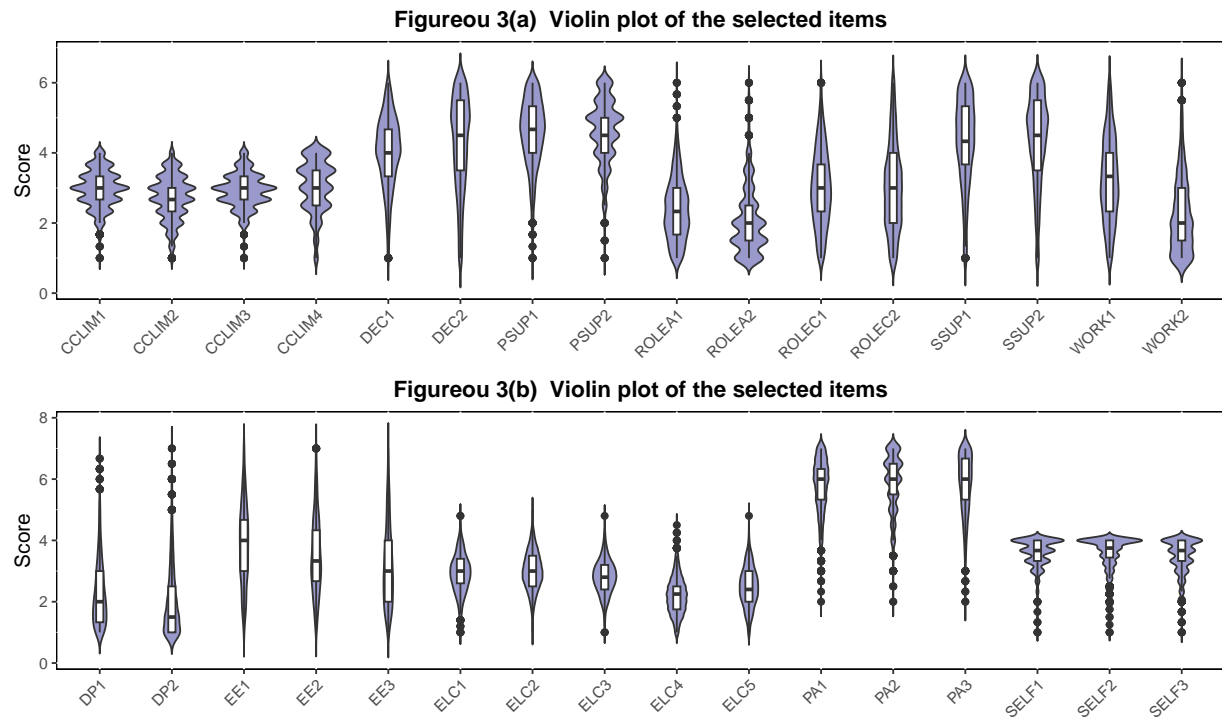
Clearly, within each instrument, indicators for same factor tend to show similar distribution features.

(2) Violin plot

Violin plot also provides information on distribution, plus ideas on out-liers.

```
a <- mbi |>
  select(1:16) |>
  violin.box(fig.num = "3(a)")

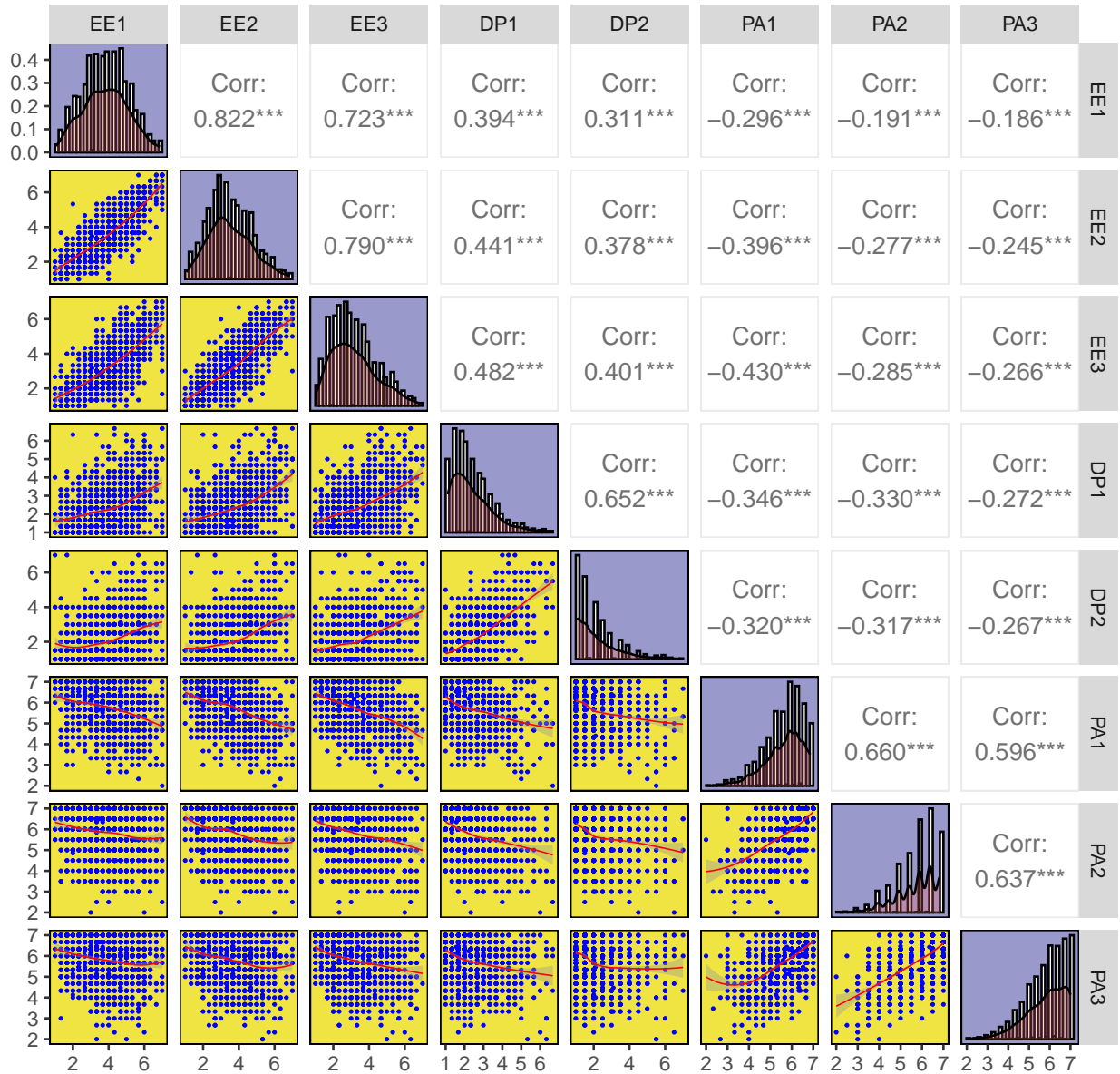
b <- mbi |>
  select(17:32) |>
  violin.box(fig.num = "3(b)")
library(patchwork)
a/b
```



(3) Correlation among items

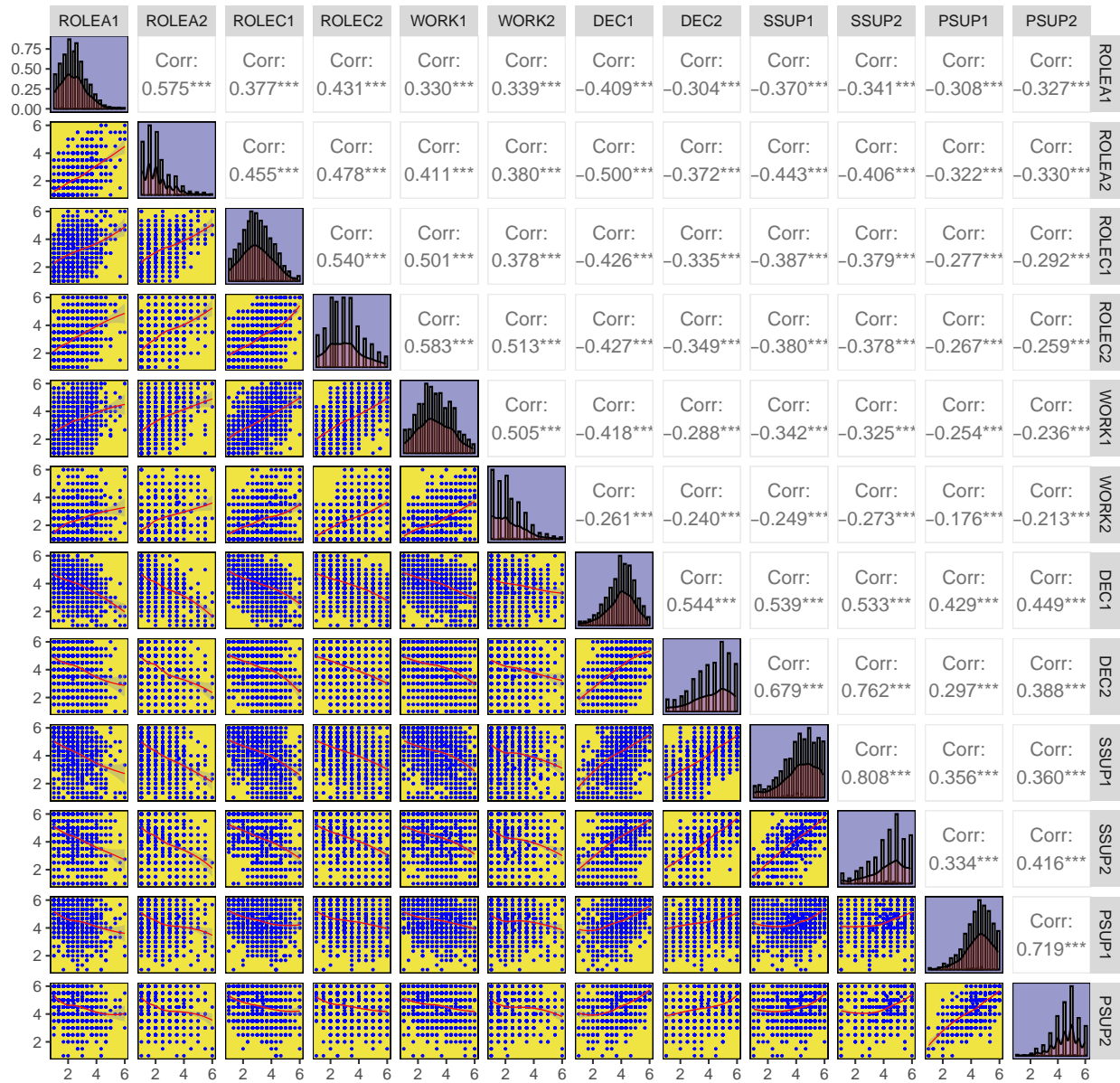
```
#draw it
mbi |> select(starts_with("EE")|starts_with("DP")|starts_with("PA")) |>
  ggpairs(lower =
    list(continuous = my.fun.smooth), #lower half show points with fitted line
    diag =
    list(continuous = my.fun.density), #diagonal grids show density plots
    title = "Fig4(a). Relationships among parcels of MBI instrument") + #title
  theme (plot.title = element_text(size = 12, #adjust title visuals
    face = "bold"))
```

Fig4(a). Relationships among parcels of MBI instrument



```
mbi |> select(starts_with("ROL")|starts_with("WOR")|starts_with("DEC")|contains("SUP")) |>
ggpairs(lower =
  list(continuous = my.fun.smooth), #lower half show points with fitted line
  diag =
    list(continuous = my.fun.density), #diagonal grids show density plots
  title = "Fig4(b). Relationships among parcels of TSS instrument") + #title
theme (plot.title = element_text(size = 14, #adjust title visuals
  face = "bold"))
```

Fig4(b). Relationships among parcels of TSS instrument



3 Testing the for the validity of causal structure of burnout

3.1 Initial full structural equation model (hypothesized model modified according to CFA)

This full structural equation model was a hypothesized model. I have established causal structure linking several stressor variables considered to contribute to the presence of burnout (Fig. 5). These postulated causal relations linking variables were supported in theory and/or empirical research. I wanted to test the hypothesis that the causal pattern was true. The findings would contribute to the understanding of key determinants of teacher burnout. Since the hypothesis was proven not true, I performed post-hoc analysis to improve the causal structure step by step, until best fitting, albeit most parsimonious, model of any set of tested models had been achieved.

Note that “an important preliminary step in the analysis of full latent variable models is to test first for the validity of the measurement model before making any attempt to evaluate the structural model. Accordingly, CFA procedures are used in testing the validity of the indicator variables. Once it is known that the measurement model is operating adequately, one can then have more confidence in findings related to assessment of the hypothesized structural model.” The current analysis started at the point where CFA had already been done. As described by Byrne, the analysis produced fit indices showing exceptionally good fit to the data; nonetheless, CFA model for the TSS was re-specified to include two additional parameters, both about allowing for cross-loading terms (DEC2 cross-loads onto Factor 1; DEC2 cross-loads onto Factor 5). These parameters set free were incorporated into the initial hypothesized model (Fig. 5).

3.1.1 Define the initial model

```
library(semPlot)#install.packages("semPlot")

initial_model <- '
# Factors:
F1ROLA =~ ROLEA1 + ROLEA2 + DEC2
F2ROLC =~ ROLEC1 + ROLEC2
F3WORK =~ WORK1 + WORK2
F4CLIM =~ CCLIM1 + CCLIM2 + CCLIM3 + CCLIM4
F5DEC =~ DEC1 + DEC2
F6SSUP =~ SSUP1 + SSUP2 + DEC2
F7PSUP =~ PSUP1 + PSUP2
F8SELF =~ SELF1 + SELF2 + SELF3
F9ELC =~ ELC1 + ELC2 + ELC3 + ELC4 + ELC5
F10EE =~ EE1 + EE2 + EE3
F11DP =~ DP1 + DP2
F12PA =~ PA1 + PA2 + PA3
# Regressions:
F8SELF ~ F5DEC + F6SSUP + F7PSUP
F9ELC ~ F5DEC
F10EE ~ F2ROLC + F3WORK + F4CLIM
F11DP ~ F2ROLC + F10EE
F12PA ~ F1ROLA + F8SELF + F9ELC + F10EE + F11DP
'
```

3.1.2 Visualize the initial model

To approximate the visual effect on slides, the coordinates for each nodes were defined on a 60 by 72 matrix.

```
#generate a matrix
m <- matrix(NA, 60, 72)

#define positions of the factors
m[12, 68] <- "F1ROLA"
m[12, 40] <- "F2ROLC"
m[12, 28] <- "F3WORK"
m[12, 12] <- "F4CLIM"
m[21, 12] <- "F5DEC"
m[40, 12] <- "F6SSUP"
m[53, 9] <- "F7PSUP"
```



```

m[44,24] <- "F8SELF"
m[52,40] <- "F9ELC"
m[37,48] <- "F10EE"
m[26,60] <- "F11DP"
m[48,64] <- "F12PA"

#define the positions of the indicators (parcelled items)
m[4, 72] <- "ROLEA1"
m[4, 64] <- "ROLEA2"
m[4, 48] <- "ROLEC1"
m[4, 40] <- "ROLEC2"
m[4, 32] <- "WORK1"
m[4, 24] <- "WORK2"
m[4, 16] <- "CCLIM1"
m[5, 10] <- "CCLIM2"
m[10, 4] <- "CCLIM3"
m[15, 4] <- "CCLIM4"
m[20, 4] <- "DEC1"
m[27, 6] <- "DEC2"
m[36, 4] <- "SSUP1"
m[40, 4] <- "SSUP2"
m[59, 6] <- "PSUP1"
m[59, 13] <- "PSUP2"
m[48, 32] <- "SELF1"
m[52, 28] <- "SELF2"
m[51, 21] <- "SELF3"
m[56, 50] <- "ELC1"
m[60, 48] <- "ELC2"
m[60, 42] <- "ELC3"
m[60, 35] <- "ELC4"
m[56, 31] <- "ELC5"
m[43, 45] <- "EE1"
m[39, 40] <- "EE2"
m[35, 38] <- "EE3"
m[20, 64] <- "DP1"
m[20, 58] <- "DP2"
m[52, 71] <- "PA1"
m[56, 64] <- "PA2"
m[53, 57] <- "PA3"

```

The diagram of the initial model was generated.

```

semPaths(semPlotModel(initial_model),
         style = "lisrel",
         rotation = 2,
         sizeLat = 6,
         sizeLat2 = 5,
         sizeMan = 5,
         sizeMan2 = 2,
         residScale = 4,
         shapeMan = "rectangle",
         edge.color = c(rep("black",34),
                       rep("blue",14)),

```

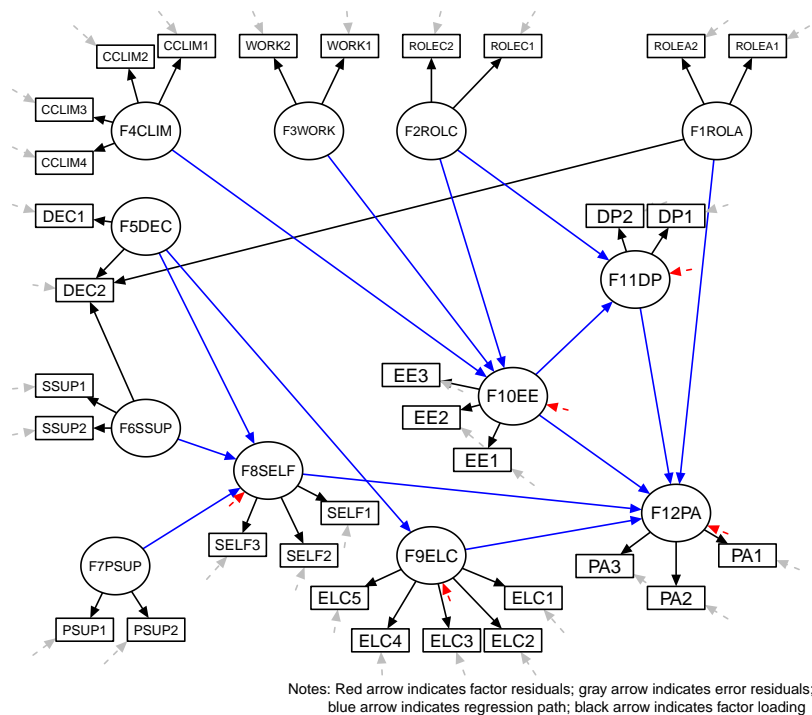
```

                                rep("gray",32),
                                rep("red",5)),

    residuals = T,
    layout = m,
    nCharNodes=0,
    optimizeLatRes = T,
    exoVar = F)
title(main = list("Figure 5. Hypothesized model of teacher burnout",
                  cex = 1.5, font =1),
      outer = F, line = -1)
title(sub = "Notes: Red arrow indicates factor residuals; gray arrow indicates error residuals;
            blue arrow indicates regression path; black arrow indicates factor loading",
      line = 0, adj = 0.7)

```

Figure 5. Hypothesized model of teacher burnout



3.1.3 Estimate the SEM model (initial)

```

library(lavaan)
modell1 <- initial_model # defined above

# Estimate the model with the robust (MLM) estimator:
sem1 <- sem(modell1, data = mbi, estimator = "MLM", mimic = "Mplus")

```

```
# Numerical summary of the model:
sem.summary.mlm.a(sem1, 12, 32, "mlm", "Model fit indices for initial model")
```

Table 2: Model fit indices for initial model

Measure	Value
chi square	1554.942
df	427.000
p value	0.000
CFI	0.945
TLI	0.936
RMSEA	0.043
RMSEA p value	1.000
SRMR	0.051
CSF	1.117

```
#summary(sem1, fit.measures = TRUE, standardized = TRUE)
```

```
#print concern table for model 1
concern.table(sem1, 14, "model1")
```

3.1.4 Comments on the result (initial model)

Here we see in tabel 2 that the re-scaled chi square value (i.e., the MLM chi square) is 1541.844 with 427 degrees of freedom. The reported chi square scaling factor value for the MLM estimator indicates that if the MLM chi square were multiplied by 1.127, it would approximate the uncorrected ML chi square value (1737.658).

Given the large number of parameters estimated in this model, the reported results are understandably lengthy. Thus, in the interest of space, I report findings pertinent to only the structural parameters, as well as a few residual variances. These results are presented in table 3.

Examination of estimated parameters in the model revealed all to be statistically significant except for those highlighted in red in table 3. These non-significant parameters the following structural regression paths: (a) F10 on F2 (Role Conflict → Emotional Exhaustion), F10 on F3 (Work Overload → Emotional Exhaustion), F10 on F4 (Classroom Climate → Emotional Exhaustion), and (b) F12 on F1 (Role Ambiguity → Personal Accomplishment).

For the negative variance for the residual associated with Factor 10 (highlighted in red in table 3), I left it as it was for the time being. The plan is over the progression of the addition of new regression path, the anomolies would disappear on itself.

3.1.5 Model mis-specification

The sense-making for the choice of parameter to set free was placed in the table 4. The path regression to be set free in next model was highlighted in red, with the logics behind selecting it listing in the last column.

A review of the MIs reveals some evidence of misfit in the model. Because we are interested solely in the causal paths of the model at this point, only MIs related to these parameters are included in table 5.

Table 3: Residual variance of structural regression path and select factors for model1

Parameter*	B [†]	Beta [‡]	SE	Z	p-value
Structural regression path					
F5DEC→F8SELF	0.475	0.990	0.050	9.462	0
F6SSUP→F8SELF	-0.155	-0.490	0.025	-6.194	0
F7PSUP→F8SELF	-0.066	-0.150	0.029	-2.307	0.021
F5DEC→F9ELC	-0.288	-0.476	0.021	-13.784	0
F2ROLC→F10EE	-8.707	-5.765	6.310	-1.380	0.168
F3WORK→F10EE	8.082	6.325	5.321	1.519	0.129
F4CLIM→F10EE	-0.93	-0.272	0.658	-1.414	0.157
F2ROLC→F11DP	0.258	0.203	0.048	5.356	0
F10EE→F11DP	0.373	0.444	0.033	11.377	0
F1ROLA→F12PA	-0.071	-0.062	0.046	-1.548	0.122
F8SELF→F12PA	0.472	0.217	0.085	5.562	0
F9ELC→F12PA	-0.208	-0.121	0.049	-4.264	0
F10EE→F12PA	-0.064	-0.098	0.025	-2.564	0.01
F11DP→F12PA	-0.218	-0.281	0.031	-7.115	0
Selected factors					
F8SELF	0.079	0.682	0.007	10.672	0
F9ELC	0.143	0.774	0.011	13.432	0
F10EE	-0.432	-0.332	0.770	-0.562	0.574
F11DP	0.605	0.658	0.047	12.880	0
F12PA	0.383	0.695	0.023	16.815	0

Note:

Values highlighted in red should be taken note of

* → indicates regression path

[†] Crude estimates

[‡] Standardized estimates

```

#extract needed variables
MI.model1 <- modindices(sem1,
                        standardized = TRUE,
                        sort. = TRUE,
                        maximum.number = 20) |>
filter(op == "~") |>
filter(lhs %in% #When these factors are predicted variables, it is related
      c("F8SELF", # to the topic of this study
        "F9ELC",
        "F10EE",
        "F11DP",
        "F12PA"))

#adapt to publication style
MI.model1 <- MI.model1 |>
mutate(Parameter = paste(rhs, "→", lhs)) |>
select(Parameter, MI = mi, EPC = epc, "std EPC" = sepc.all) |>
filter(MI>10) |> #for saving space, the number of parameters was managed
mutate("Logics" = c("Sensible and meaningful",
                    "illogical (wrong direction of correlation)",
                    "Sensibe but not meaningful for this study"))

#add footnote symbol to parameters
for (i in 1:nrow(MI.model1)){
  symbol <- c("∗", "†", "‡") #symbols for footnotes
  Parameter <- unlist(MI.model1$Parameter)
  MI.model1$Parameter[i] <- paste0(Parameter[i], symbol[i])
}

#Visualize the table
MI.model1 |>
kable(digits = 3,
      booktab = T,
      linesep = "",
      caption = "Selected modification indices for initial model") |>
kable_styling(latex_options = "striped") |>
row_spec(1, color = "red") |> #this is the row with parameter to set free
footnote(general =
  "Parameter highlighted in red is selected for modification",
  symbol = c("Poorer Classroom climate leads to worsening depersonalization",
             "Less workload leads to worsening depersonalization",
             "Higher accomplishment results in increased self-esteem"))

```

3.2 Post hos analysis (Model 2)

3.2.1 Compare SEM model 2 with initial model

I defined model 2 as per the conclusion from last section: set the regression path leading from F4 to F11 (Poorer Classroom climate leads to worsening depersonalization) free to estimate (Fig 6). The model fit indices and its comparison with the preceding model (model1) was tabulated in table 5.

Table 4: Selected modification indices for initial model

Parameter	MI	EPC	std EPC	Logics
F4CLIM → F11DP*	112.597	-0.974	-0.339	Sensible and meaningful
F3WORK → F11DP†	85.318	-4.799	-4.467	illogical (wrong direction of correlation)
F12PA → F8SELF‡	49.986	0.329	0.719	Sensibe but not meaningful for this study

Note:

Parameter highlighted in red is selected for modification

* Poorer Classroom climate leads to worsening depersonalization

† Less workload leads to worsening depersonalization

‡ Higher accomplishment results in increased self-esteem

Table 5: Comparison of new and preceding models

	Chi-square	DF	p value	ΔChi-square	CFI	TLI	RMSEA	SRMR	CSF
Model1	1554.942	427	0	–	0.945	0.936	0.043	0.051	1.117
Model2*	1450.985	426	0	91.67	0.950	0.941	0.041	0.046	1.117

* Model1 + structural path 'F4→F11' set free to estimate

```

model2 <- paste(initial_model, "F11DP ~ F4CLIM")

# Estimate the model with the robust (MLM) estimator:
sem2 <- sem(model2, data = mbi, estimator = "MLM", mimic = "Mplus")

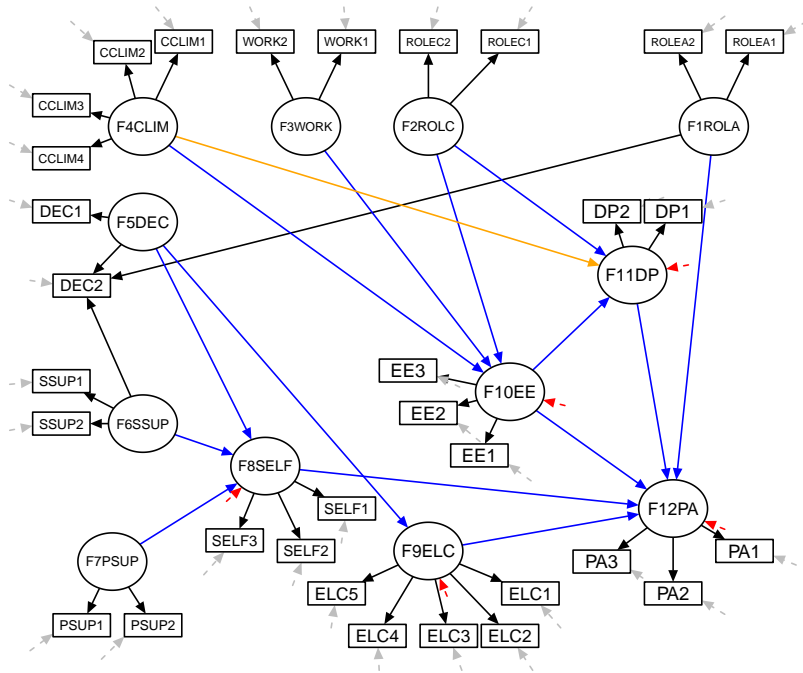
sem <- c("sem1", "sem2")
newpathstext <- " Model1 + structural path 'F4→F11' set free to estimate"

model.comparison.table (sem,
                        mycaption = "Comparison of new and preceding models",
                        newpathstext = newpathstext)

#plot the diagram
full.sem.diagram(model2, 15, "6", "model2")

```

Figure 6 Modified model (model2) of teacher burnout



Notes: Orange arrow is the regression path set free to estimate in the current model

3.2.2 Examine the parameters of concern for model 2

```
#regression path estimates
concern.table(sem2, 15, "model2")
```

3.2.3 Comment on the result (model 2)

The estimation of Model 2 yielded an overall MLM chi square(426) value of 1450.985 (scaling correction factor = 1.117); CFI and TLI values of 0.950 and 0.945, respectively; a RMSEA value of 0.041; and a SRMR value of 0.046 (Table 5). Although improvement in model fit for Model 2, compared with the originally hypothesized model, would appear to be somewhat minimal on the basis of the CFI, TLI, RMSEA, and SRMR values, the corrected chi-square difference test was found to be significant (MLM Δ chi square[1] = 91.67), which finalized the decision (Table 5).

Notes from Byrne's book: "the thrust of these post-hoc analyses is to fine-tune our hypothesized structure such that it includes all viable and statistically significant structural paths, and, at the same time, eliminates all non-significant paths. Consequently, as long as the chi square-difference test is statistically significant, and the newly added parameters are substantively meaningful, I consider the post-hoc analyses to be appropriate."

The anomaly of negative residual variance remained in the output for Model 2 (table 6). The estimated structural regression paths for the three factors hypothesized to influence Factor 10 (Factors 2, 3, and 4) and F1 to influence Factor 12 remained statistically non-significant (table 6).

Besides, The negative residual variance of F10 remained as it was in model 2 (table 6).

Table 6: Residual variance of structural regression path and select factors for model2

Parameter*	B†	Beta‡	SE	Z	p-value
Strutural regression path					
F5DEC→F8SELF	0.474	0.989	0.050	9.453	0
F6SSUP→F8SELF	-0.155	-0.489	0.025	-6.198	0
F7PSUP→F8SELF	-0.066	-0.150	0.029	-2.308	0.021
F5DEC→F9ELC	-0.288	-0.476	0.021	-13.791	0
F2ROLC→F10EE	-8.396	-5.553	5.945	-1.412	0.158
F3WORK→F10EE	7.856	6.151	5.034	1.561	0.119
F4CLIM→F10EE	-0.563	-0.164	0.633	-0.888	0.374
F2ROLC→F11DP	0.173	0.135	0.046	3.726	0
F10EE→F11DP	0.299	0.354	0.031	9.627	0
F1ROLA→F12PA	-0.069	-0.060	0.046	-1.489	0.137
F8SELF→F12PA	0.473	0.217	0.084	5.599	0
F9ELC→F12PA	-0.204	-0.118	0.049	-4.190	0
F10EE→F12PA	-0.058	-0.089	0.024	-2.411	0.016
F11DP→F12PA	-0.228	-0.297	0.030	-7.533	0
F4CLIM→F11DP	-0.969	-0.334	0.100	-9.642	0
Selected factors					
F8SELF	0.079	0.682	0.007	10.678	0
F9ELC	0.143	0.774	0.011	13.421	0
F10EE	-0.377	-0.289	0.720	-0.524	0.6
F11DP	0.535	0.574	0.044	12.120	0
F12PA	0.38	0.689	0.023	16.857	0

Note:

Values highlighted in red should be taken note of

* → indicates regression path

† Crude estimates

‡ Standardized estimates

3.2.4 Model mis-specification (model 2)

```
#extract needed variables
MI.model2 <- modindices(sem2,
  standardized = TRUE,
  sort. = TRUE,
  maximum.number = 25) |>
filter(op == "~") |>
filter(lhs %in% #When these factors are predicted variables, it is related
  c("F8SELF", # to the topic of this study
    "F9ELC",
    "F10EE",
    "F11DP",
    "F12PA"))

#adapt to publication style
MI.model2 <- MI.model2 |>
mutate(Parameter = paste(rhs, "→", lhs)) |>
select(Parameter, MI = mi, EPC = epc, "std EPC" = sepc.all) |>
filter(MI>10) |>
mutate("Logics" = c("Sensible but not meaningful for this study",
  "Sensible and meaningful",
  "Sensibe and meaningful, but MI is lower than F5→F12"))

#add footnote symbol to parameters
for (i in 1:nrow(MI.model2)){
  symbol <- c("∗", "†", "‡")
  Parameter <- unlist(MI.model2$Parameter)
  MI.model2$Parameter[i] <- paste0(Parameter[i], symbol[i])
}

#Visualize the table
MI.model2 |>
kable(digits = 3,
  booktab = T,
  linesep = "",
  caption = "Selected modification indices for initial model") |>
kable_styling(latex_options = "striped") |>
row_spec(2, color = "red") |>
footnote(general =
  "Parameter highlighted in red is selected for modification",
  symbol = c("Higher accomplishment leads to increased self-esteem",
    "Invovlement of more decision making gives sense of accomplishment",
    "People with high esteem will less likely get emotionally exhausted"))
```

3.3 Post hos analysis (Model 3)

3.3.1 Compare SEM model 3 with preceding models

I defined model 3 as per the conclusion from last section: on the basis of model 2, I set the regression path leading from F5 to F12 (Invovlement of more decision making gives sense of accomplishment) free to estimate

Table 7: Selected modification indices for initial model

Parameter	MI	EPC	std EPC	Logics
F12PA → F8SELF*	49.859	0.330	0.720	Sensible but not meaningful for this study
F5DEC → F12PA†	47.088	0.480	0.458	Sensible and meaningful
F8SELF → F10EE‡	42.770	-1.029	-0.306	Sensibe and meaningful, but MI is lower than F5→F12

Note:

Parameter highlighted in red is selected for modification

* Higher accomplishment leads to increased self-esteem

† Invovlement of more decision making gives sense of accomplishment

‡ People with high esteem will less likely get emotionally exhausted

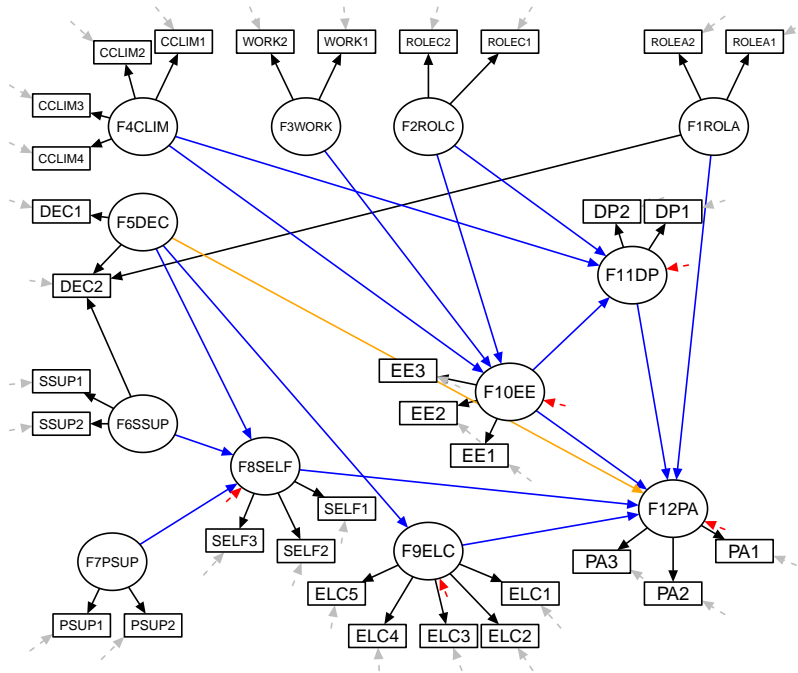
(Fig 7). The model fit indices and their comparison with the preceding models were tabulated in table 8.

```
#add new freely estimated parameter to the preceding model
model3 <- paste(model2, "\nF12PA ~ F5DEC")

# Estimate the model with the robust (MLM) estimator:
sem3 <- sem(model3, data = mbi, estimator = "MLM", mimic = "Mplus")

full.sem.diagram(model3, 16, "7", "model3")
```

Figure 7 Modified model (model3) of teacher burnout



Notes: Orange arrow is the regression path set free to estimate in the current model

Table 8: Comparison of new (model3) and preceding models

	Chi-square	DF	p value	Δ Chi-square	CFI	TLI	RMSEA	SRMR	CSF
Model1	1554.942	427	0	–	0.945	0.936	0.043	0.051	1.117
Model2*	1450.985	426	0	91.67	0.950	0.941	0.041	0.046	1.117
Model3†	1406.517	425	0	46.866	0.952	0.944	0.040	0.044	1.117

* Model1 + structural path 'F4→F11' set free to estimate

† Model2 + structural path 'F5→F12' set free to estimate

```
sem <- c("sem1", "sem2", "sem3")
newpathstext <- c(" Model1 + structural path 'F4→F11' set free to estimate",
                  " Model2 + structural path 'F5→F12' set free to estimate")

model.comparison.table (sem,
                        mycaption = "Comparison of new (model3) and preceding models",
                        newpathstext = newpathstext)
```

3.3.2 Examine the parameters of concern for model 3

```
concern.table(sem3, 16, "model3")
```

3.3.3 Comment on the result (model 3)

Model 3 yielded an overall MLM chi square(425) value of 1406.517 (scaling correction factor = 1.117), with CFI = 0.952, TLI = 0.944, RMSEA = 0.040, and SRMR = 0.044. Again, the MLM chi square difference between Models 2 and 3 is statistically significant (MLM chi square[1] = 46.866), albeit differences in the other fit indices across Models 2 and 3 were once again minimal. See table 8.

As expected, the estimate for the newly incorporated path from Decision Making to Personal Accomplishment (F12 on F5) was found to be statistically significant. However, the three previous non-significant path remained p values > 0.05 (two leading to F12; one leading to F10). Besides, The negative residual variance of F10 remained as it was in model 3 (see tab 9, estimate highlighted in red).

3.3.4 Model mis-specification (model 3)

```
#extract needed variables
MI.model3 <- modindices(sem3,
                        standardized = TRUE,
                        sort. = TRUE,
                        maximum.number = 50) |>
filter(op == "~") |>
filter(lhs %in% #When these factors are predicted variables, it is related
       c("F8SELF", # to the topic of this study
         "F9ELC",
         "F10EE",
         "F11DP",
```

Table 9: Residual variance of structural regression path and select factors for model3

Parameter*	B [†]	Beta [‡]	SE	Z	p-value
Strutural regression path					
F5DEC→F8SELF	0.473	0.989	0.050	9.432	0
F6SSUP→F8SELF	-0.157	-0.496	0.025	-6.363	0
F7PSUP→F8SELF	-0.061	-0.138	0.028	-2.197	0.028
F5DEC→F9ELC	-0.288	-0.477	0.021	-13.722	0
F2ROLC→F10EE	-6.345	-4.198	3.393	-1.870	0.062
F3WORK→F10EE	6.113	4.797	2.870	2.130	0.033
F4CLIM→F10EE	-0.576	-0.168	0.488	-1.181	0.237
F2ROLC→F11DP	0.176	0.137	0.046	3.779	0
F10EE→F11DP	0.3	0.353	0.031	9.611	0
F1ROLA→F12PA	0.333	0.285	0.085	3.917	0
F8SELF→F12PA	0.469	0.213	0.085	5.488	0
F9ELC→F12PA	-0.153	-0.088	0.053	-2.919	0.004
F10EE→F12PA	0.025	0.038	0.028	0.887	0.375
F11DP→F12PA	-0.2	-0.259	0.030	-6.575	0
F4CLIM→F11DP	-0.963	-0.330	0.100	-9.599	0
F5DEC→F12PA	0.504	0.480	0.083	6.054	0
Selected factors					
F8SELF	0.077	0.669	0.007	10.651	0
F9ELC	0.143	0.772	0.011	13.405	0
F10EE	-0.124	-0.095	0.410	-0.303	0.762
F11DP	0.545	0.578	0.045	12.103	0
F12PA	0.36	0.644	0.023	15.840	0

Note:

Values highlighted in red should be taken note of

* → indicates regression path

[†] Crude estimates

[‡] Standardized estimates

Table 10: Selected modification indices for initial model

Parameter	MI	EPC	std EPC	Logics
F8SELF → F10EE*	43.412	-1.021	-0.304	Sensible and meaningful, but not as sensible as F2→F9
F2ROLC → F9ELC†	41.343	0.191	0.336	Highly sensible and meaningful
F8SELF → F9ELC‡	39.841	-0.284	-0.225	Sensible and meaningful, but MI < other 2

Note:

Parameter highlighted in red is selected for modification

* Lower self-esteem will causes higher emotional exhaustion

† Higher role conflict increases external locus of control

‡ Lower self-esteem leads to higher external locus of control

```

    "F12PA"))

#adapt to publication style
MI.model3 <- MI.model3 |>
  mutate(Parameter = paste(rhs, "→", lhs)) |>
  select(Parameter, MI = mi, EPC = epc, "std EPC" = sepc.all) |>
  filter(MI>10) |>
  head(3) |>
  mutate("Logics" = c("Sensible and meaningful, but not as sensible as F2→F9",
                      "Highly sensible and meaningful",
                      "Sensible and meaningful, but MI < other 2"))

#add footnote symbol to parameters
for (i in 1:nrow(MI.model3)){
  symbol <- c("∗", "†", "‡")
  Parameter <- unlist(MI.model3$Parameter)
  MI.model3$Parameter[i] <- paste0(Parameter[i], symbol[i])
}

#Visualize the table
MI.model3 |>
  kable(digits = 3,
        booktab = T,
        linesep = "",
        caption = "Selected modification indices for initial model") |>
  kable_styling(latex_options = "striped") |>
  row_spec(2, color = "red") |>
  footnote(general =
    "Parameter highlighted in red is selected for modification",
    symbol = c("Lower self-esteem will causes higher emotional exhaustion",
              "Higher role conflict increases external locus of control",
              "Lower self-esteem leads to higher external locus of control"))

```

Quoted from Byrne, “Again, I believe it is worthwhile to note why two alternate MI values, close in value to the one chosen here, are considered to be inappropriate. I refer to results related to the structural paths of F10 on F8 (MI = 38.868) and of F9 on F8 (MI = 35.670). In both cases, the flow of causal direction is incorrect.” I don’t agree with her comments on this wrong causal direction. To me, it is fairly logical, though not perfectly straightforward, that Lower self-esteem will cause higher emotional exhaustion (F8→F10, negative

sign of EPC), and that lower self-esteem leads to higher external locus of control (F8→F9, negative sign of EPC). See table 10. However, I agree with Bynre's decision on most appropriate parameter F2→F9 (Higher role conflict increases external locus of control). The causal effect for this parameter is most intuitive –role conflict is a type of external factors that has negative influence. Considering its MI (41.343) is so close to the highest seen in this model (43.412), substantive meaningfulness, in my opinion, should have the final say when statistics disagrees moderately. As such, I will continue with setting F2 → F9 free to estimate in model 4. See next section.

3.4 Post hos analysis (Model 4)

3.4.1 Compare SEM model 4 with preceding models

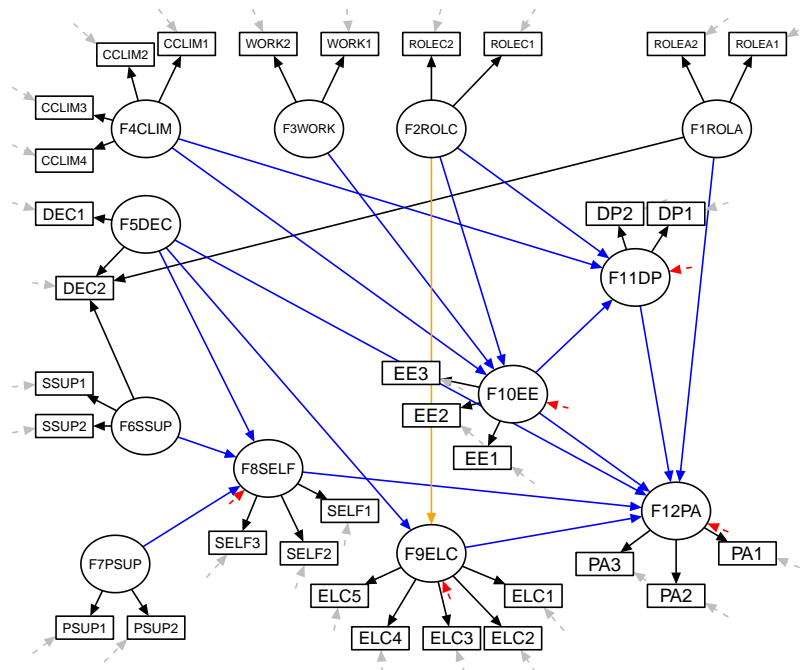
I defined model 4 as per the conclusion from last section: on the basis of model 3, I set the regression path leading from F2 to F9 (Higher role conflict increases external locus of control) free to estimate (Fig 8). The model fit indices and their comparison with the preceding models were tabulated in table 11.

```
#add new freely estimated parameter to the preceding model
model4 <- paste(model3, "\nF9ELC ~ F2ROLC")

# Estimate the model with the robust (MLM) estimator:
sem4 <- sem(model4, data = mbi, estimator = "MLM", mimic = "Mplus")

full.sem.diagram(model4, 17, "8", "model4")
```

Figure 8 Modified model (model4) of teacher burnout



Notes: Orange arrow is the regression path set free to estimate in the current model

Table 11: Comparison of new (model 4) and preceding models

	Chi-square	DF	p value	Δ Chi-square	CFI	TLI	RMSEA	SRMR	CSF
Model1	1554.942	427	0	–	0.945	0.936	0.043	0.051	1.117
Model2*	1450.985	426	0	91.67	0.950	0.941	0.041	0.046	1.117
Model3†	1406.517	425	0	46.866	0.952	0.944	0.040	0.044	1.117
Model4‡	1366.129	424	0	52.949	0.954	0.946	0.039	0.041	1.118

* Model1 + structural path 'F4→F11' set free to estimate

† Model2 + structural path 'F5→F12' set free to estimate

‡ Model3 + structural path 'F2→F9' set free to estimate

```
sem <- c("sem1", "sem2", "sem3", "sem4")
newpathstext <- c(" Model1 + structural path 'F4→F11' set free to estimate",
                  " Model2 + structural path 'F5→F12' set free to estimate",
                  " Model3 + structural path 'F2→F9' set free to estimate")

model.comparison.table (sem,
                        mycaption = "Comparison of new (model 4) and preceding models",
                        newpathstext = newpathstext)
```

3.4.2 Examine the parameters of concern for model 4

```
concern.table(sem4, 17, "model4")
```

3.4.3 Comment on the result (model 4)

The estimation of Model 4 yielded a MLM chi square value of 1366.129 (scaling correction factor = 1.118) with 424 degrees of freedom. Values related to the CFI, TLI, RMSEA, and SRMR were .954, .946, 0.039, and 0.041, respectively. Again, the difference in fit between this model (Model 4) and its predecessor (Model 3) was statistically significant (MLM chi square[1] = 52.949). See table 11.

As expected, the newly specified parameter (F9 on F2) was found to be statistically significant (Estimate = 0.189; Z-statistics = 7.157). However, once again the three paths leading from F2, F3, and F4 to F10, and from F10 to F12 were all found to be non-significant. Finally, once again, the negative residual associated with Factor 10 appeared. See table 12.

3.4.4 Model mis-specification

```
#extract needed variables
MI.model4 <- modindices(sem4,
                        standardized = TRUE,
                        sort. = TRUE,
                        maximum.number = 50) |>
filter(op == "~") |>
filter(lhs %in% #When these factors are predicted variables, it is related
       c("F8SELF", # to the topic of this study)
```

Table 12: Residual variance of structural regression path and select factors for model4

Parameter*	B [†]	Beta [‡]	SE	Z	p-value
Strutural regression path					
F5DEC→F8SELF	0.566	1.177	0.067	8.412	0
F6SSUP→F8SELF	-0.205	-0.648	0.033	-6.147	0
F7PSUP→F8SELF	-0.1	-0.228	0.036	-2.784	0.005
F5DEC→F9ELC	-0.125	-0.208	0.027	-4.638	0
F2ROLC→F10EE	-8.853	-5.875	6.316	-1.402	0.161
F3WORK→F10EE	8.26	6.453	5.359	1.541	0.123
F4CLIM→F10EE	-0.777	-0.226	0.659	-1.179	0.239
F2ROLC→F11DP	0.176	0.138	0.047	3.740	0
F10EE→F11DP	0.3	0.354	0.032	9.537	0
F1ROLA→F12PA	0.271	0.232	0.075	3.626	0
F8SELF→F12PA	0.494	0.225	0.087	5.714	0
F9ELC→F12PA	-0.166	-0.095	0.052	-3.178	0.001
F10EE→F12PA	0.013	0.020	0.027	0.478	0.633
F11DP→F12PA	-0.202	-0.262	0.030	-6.671	0
F4CLIM→F11DP	-0.958	-0.329	0.100	-9.575	0
F5DEC→F12PA	0.432	0.410	0.068	6.387	0
F2ROLC→F9ELC	0.189	0.336	0.026	7.157	0
Selected factors					
F8SELF	0.078	0.672	0.007	10.445	0
F9ELC	0.136	0.744	0.010	12.968	0
F10EE	-0.441	-0.338	0.762	-0.579	0.563
F11DP	0.544	0.578	0.045	12.082	0
F12PA	0.363	0.651	0.023	16.089	0

Note:

Values highlighted in red should be taken note of

* → indicates regression path

[†] Crude estimates

[‡] Standardized estimates

Table 13: Selected modification indices for initial model

Parameter	MI	EPC	std EPC	Logics
F8SELF → F9ELC*	38.685	-0.261	-0.208	Sensible and meaningful
F12PA → F8SELF†	37.407	0.302	0.662	Sensible but not meaningful for the study
F12PA → F9ELC‡	30.392	-0.265	-0.464	Sensible and not meaningful for the study

Note:

Parameter highlighted in red is selected for modification

* Lower self-esteem causes severer external locus of control

† Higher personal accomplishment increases self-esteem

‡ Higher personal accomplishment decreases external locus of control

```

      "F9ELC",
      "F10EE",
      "F11DP",
      "F12PA"))

#adapt to publication style
MI.model4 <- MI.model4 |>
  mutate(Parameter = paste(rhs, "→", lhs)) |>
  select(Parameter, MI = mi, EPC = epc, "std EPC" = sepc.all) |>
  filter(MI>10) |>
  head(3) |>
  mutate("Logics" = c("Sensible and meaningful",
                      "Sensible but not meaningful for the study",
                      "Sensible and not meaningful for the study"))

#add footnote symbol to parameters
for (i in 1:nrow(MI.model4)){
  symbol <- c("∗", "†", "‡")
  Parameter <- unlist(MI.model4$Parameter)
  MI.model4$Parameter[i] <- paste0(Parameter[i], symbol[i])
}

#Visualize the table
MI.model4 |>
  kable(digits = 3,
        booktab = T,
        linesep = "",
        caption = "Selected modification indices for initial model") |>
  kable_styling(latex_options = "striped") |>
  row_spec(1, color = "red") |>
  footnote(general =
    "Parameter highlighted in red is selected for modification",
    symbol = c("Lower self-esteem causes severer external locus of control",
              "Higher personal accomplishment increases self-esteem",
              "Higher personal accomplishment decreases external locus of control"))

```

The regression path F8→F9 shows up again but this time it has the highest MI among its peers. See table 13. In the preceding model (model 3) I did not adopt it only because its relatively lower MI albeit good

sensibility and meaningfulness. Given that other two competitor parameters in the current model were bad in meaningfulness (because they regress risk factors on the outcome of interest, but we want the opposite), we could decide on adopting the path of F8→F9 (Lower self-esteem causes severer external locus of control) without much ado.

3.5 Post hos analysis (Model 5)

3.5.1 Compare SEM model 5 with preceding models

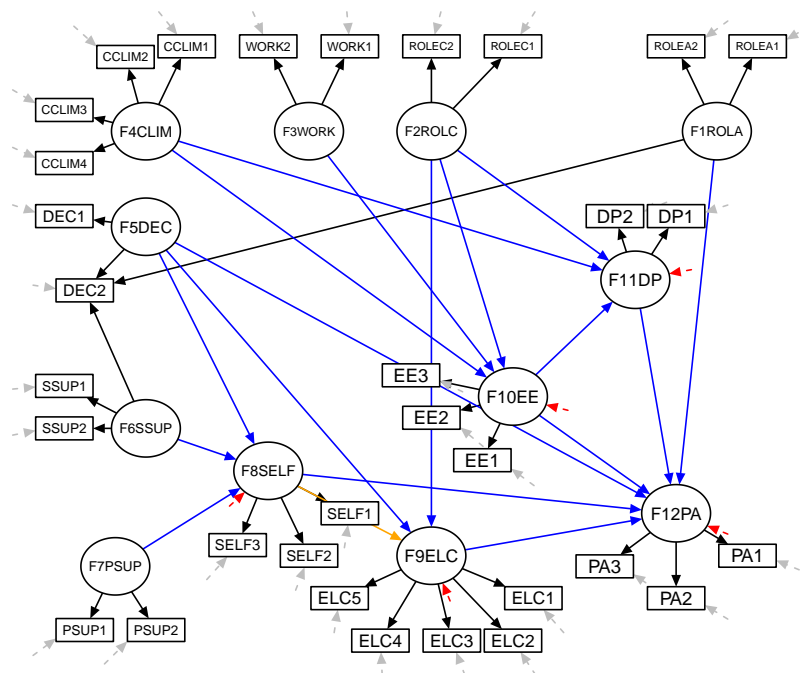
I defined model 5 as per the conclusion from last section: on the basis of model 4, I set the regression path leading from F8 to F9 (Lower self-esteem causes severer external locus of control) free to estimate (Fig 9). The model fit indices and their comparison with the preceding models were tabulated in table 14.

```
#add new freely estimated parameter to the preceding model
model5 <- paste(model4, "\nF9ELC ~ F8SELF")

# Estimate the model with the robust (MLM) estimator:
sem5 <- sem(model5, data = mbi, estimator = "MLM", mimic = "Mplus")

full.sem.diagram(model5, 18, "9", "model5")
```

Figure 9 Modified model (model5) of teacher burnout



Notes: Orange arrow is the regression path set free to estimate in the current model

```
sem <- c("sem1", "sem2", "sem3", "sem4", "sem5")
newpathstext <- c(" Model1 + structural path 'F4→F11' set free to estimate",
```

Table 14: Comparison of new (model 5) and preceding models

	Chi-square	DF	p value	Δ Chi-square	CFI	TLI	RMSEA	SRMR	CSF
Model1	1554.942	427	0	–	0.945	0.936	0.043	0.051	1.117
Model2*	1450.985	426	0	91.67	0.950	0.941	0.041	0.046	1.117
Model3†	1406.517	425	0	46.866	0.952	0.944	0.040	0.044	1.117
Model4‡	1366.129	424	0	52.949	0.954	0.946	0.039	0.041	1.118
Model5§	1331.636	423	0	29.877	0.955	0.948	0.039	0.039	1.117

* Model1 + structural path 'F4→F11' set free to estimate

† Model2 + structural path 'F5→F12' set free to estimate

‡ Model3 + structural path 'F2→F9' set free to estimate

§ Model5 + structural path 'F8→F9' set free to estimate

```

      " Model2 + structural path 'F5→F12' set free to estimate",
      " Model3 + structural path 'F2→F9' set free to estimate",
      " Model5 + structural path 'F8→F9' set free to estimate")

model.comparison.table (sem,
  mycaption =
    "Comparison of new (model 5) and preceding models",
  newpathstext = newpathstext)

```

3.5.2 Examine the parameters of concern for model 5

```
concern.table(sem5, 18, "model5")
```

3.5.3 Comment on the result (model 5)

The estimation of Model 5 yielded a MLM chi square(423) value of 1331.636 (scaling correction factor = 1.117), with other fit indices as follows: CFI = 0.955, TLI = 0.948, RMSEA = 0.039, and SRMR = 0.039. Again, the difference in fit between this model (Model 5) and its predecessor (Model 4) was found to be statistically significant (MLM chi square[1] = 29.877). See table 14.

Once again, the newly specified parameter (F9 on F8) was found to be statistically significant and accompanied by the correct sign (Estimate = -0.257; Z-statistics = -5.337). However, as with Model 4, again two of the three paths leading to F10 (regressions of F2 → F10; F4 → F10), and one leading from F10 to F12 remained statistically non-significant. Besides, the negative residual associated with Factor 10 remained unchanged. See table 15.

3.5.4 Model mis-specification

```

#extract needed variables
MI.model5 <- modindices(sem5,
  standardized = TRUE,
  sort. = TRUE,

```

Table 15: Residual variance of structural regression path and select factors for model5

Parameter*	B [†]	Beta [‡]	SE	Z	p-value
Strutural regression path					
F5DEC→F8SELF	0.591	1.221	0.074	8.038	0
F6SSUP→F8SELF	-0.22	-0.696	0.037	-5.960	0
F7PSUP→F8SELF	-0.109	-0.249	0.039	-2.826	0.005
F5DEC→F9ELC	-0.076	-0.125	0.027	-2.803	0.005
F2ROLC→F10EE	-6.508	-4.323	3.440	-1.892	0.058
F3WORK→F10EE	6.262	4.907	2.914	2.149	0.032
F4CLIM→F10EE	-0.718	-0.209	0.492	-1.460	0.144
F2ROLC→F11DP	0.173	0.135	0.047	3.688	0
F10EE→F11DP	0.302	0.356	0.031	9.640	0
F1ROLA→F12PA	0.259	0.222	0.074	3.512	0
F8SELF→F12PA	0.521	0.237	0.090	5.815	0
F9ELC→F12PA	-0.162	-0.092	0.055	-2.957	0.003
F10EE→F12PA	0.013	0.019	0.027	0.463	0.643
F11DP→F12PA	-0.202	-0.262	0.030	-6.680	0
F4CLIM→F11DP	-0.959	-0.329	0.100	-9.579	0
F5DEC→F12PA	0.417	0.392	0.065	6.386	0
F2ROLC→F9ELC	0.174	0.310	0.026	6.744	0
F8SELF→F9ELC	-0.257	-0.205	0.048	-5.337	0
Selected factors					
F8SELF	0.081	0.705	0.008	10.678	0
F9ELC	0.13	0.718	0.010	12.662	0
F10EE	-0.171	-0.131	0.428	-0.400	0.689
F11DP	0.544	0.578	0.045	12.079	0
F12PA	0.365	0.653	0.023	16.233	0

Note:

Values highlighted in red should be taken note of

* → indicates regression path

[†] Crude estimates

[‡] Standardized estimates

Table 16: Selected modification indices for initial model

Parameter	MI	EPC	std EPC	Logics
F8SELF → F10EE*	30.096	-1.068	-0.318	Sensible and meaningful
F11DP → F8SELF†	25.671	-0.066	-0.188	Sensible but not meaningful for the study

Note:

Parameter highlighted in red is selected for modification

* Lower self-esteem causes severer emotional exhaustion

† Higher depersonalization causes severer role conflict

```

maximum.number = 50) |>
filter(op == "~") |>
filter(lhs %in% #When these factors are predicted variables, it is related
  c("F8SELF", # to the topic of this study
    "F9ELC",
    "F10EE",
    "F11DP",
    "F12PA"))

#adapt to publication style
MI.model5 <- MI.model5 |>
mutate(Parameter = paste(rhs, "→", lhs)) |>
select(Parameter, MI = mi, EPC = epc, "std EPC" = sepc.all) |>
head(2) |>
mutate("Logics" = c("Sensible and meaningful",
  "Sensible but not meaningful for the study"))

#add footnote symbol to parameters
for (i in 1:nrow(MI.model5)){
  symbol <- c("∗", "†", "‡")
  Parameter <- unlist(MI.model5$Parameter)
  MI.model5$Parameter[i] <- paste0(Parameter[i], symbol[i])
}

#Visualize the table
MI.model5 |>
kable(digits = 3,
  booktab = T,
  linesep = "",
  caption = "Selected modification indices for initial model") |>
kable_styling(latex_options = "striped") |>
row_spec(1, color = "red") |>
footnote(general =
  "Parameter highlighted in red is selected for modification",
  symbol = c("Lower self-esteem causes severer emotional exhaustion",
    "Higher depersonalization causes severer role conflict"))

```

This output reveals the structural path leading from Self-Esteem to Emotional Exhaustion (F8 → F10) as having the largest MI value. Given that the fact that it seems reasonable that teachers who exhibit high levels of self-esteem may exhibit low levels of emotional exhaustion, the model was re-estimated once again, with this path freely estimated. See tab .One other interesting reason to do so, cited from Byrne is

that “Because Factor 10 has been problematic regarding the estimation of its residual in yielding a negative variance, it would seem likely that if this parameter were to be included in the model, this undesirable result may finally be resolved.”

3.6 Post hos analysis (Model 6)

3.6.1 Compare SEM model 6 with preceding models

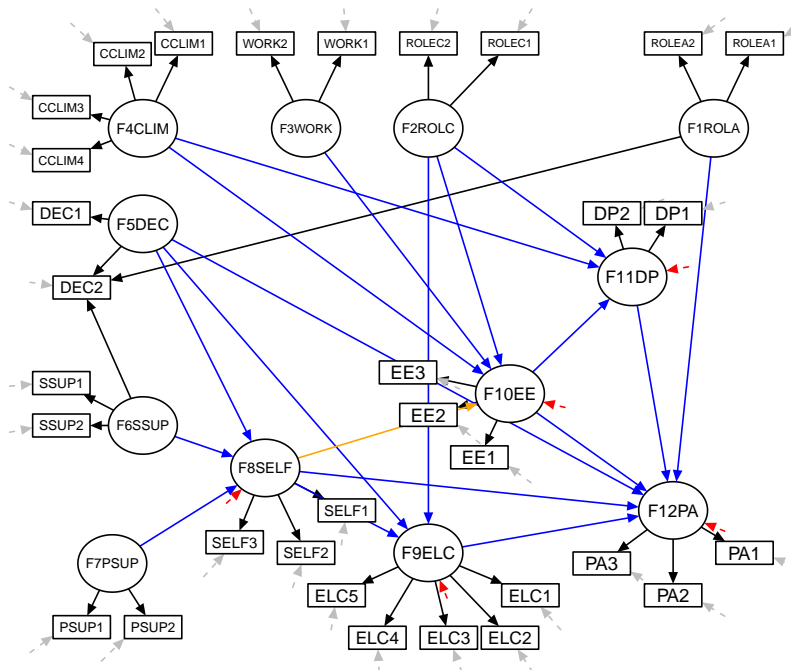
I defined model 6 as per the conclusion from last section: on the basis of model 5, I set the regression path leading from F8 to F9 (Lower self-esteem causes severer emotional exhaustion) free to estimate (Fig 10). The model fit indices and their comparison with the preceding models were tabulated in table 17.

```
#add new freely estimated parameter to the preceding model
model6 <- paste(model5, "\nF10EE ~ F8SELF")

# Estimate the model with the robust (MLM) estimator:
sem6 <- sem(model6, data = mbi, estimator = "MLM", mimic = "Mplus")

full.sem.diagram(model6, 19, "10", "model6")
```

Figure 10 Modified model (model6) of teacher burnout



Notes: Orange arrow is the regression path set free to estimate in the current model

```
sem <- c("sem1", "sem2", "sem3", "sem4", "sem5", "sem6")
newpathstext <- c(" Model1 + structural path 'F4→F11' set free to estimate",
  " Model2 + structural path 'F5→F12' set free to estimate",
```

Table 17: Comparison of new (model 6) and preceding models

	Chi-square	DF	p value	Δ Chi-square	CFI	TLI	RMSEA	SRMR	CSF
Model1	1554.942	427	0	—	0.945	0.936	0.043	0.051	1.117
Model2*	1450.985	426	0	91.67	0.950	0.941	0.041	0.046	1.117
Model3†	1406.517	425	0	46.866	0.952	0.944	0.040	0.044	1.117
Model4‡	1366.129	424	0	52.949	0.954	0.946	0.039	0.041	1.118
Model5§	1331.636	423	0	29.877	0.955	0.948	0.039	0.039	1.117
Model6¶	1297.489	422	0	23.358	0.957	0.949	0.038	0.039	1.116

* Model1 + structural path 'F4→F11' set free to estimate

† Model2 + structural path 'F5→F12' set free to estimate

‡ Model3 + structural path 'F2→F9' set free to estimate

§ Model5 + structural path 'F8→F9' set free to estimate

¶ Model6 + structural path 'F8→F10' set free to estimate

```

      " Model3 + structural path 'F2→F9' set free to estimate",
      " Model5 + structural path 'F8→F9' set free to estimate",
      " Model6 + structural path 'F8→F10' set free to estimate")

model.comparison.table (sem,
  mycaption =
    "Comparison of new (model 6) and preceding models",
  newpathstext = newpathstext)

```

3.6.2 Examine the parameters of concern for model 6

```
concern.table(sem6, 19, "model6")
```

3.6.3 Comment on the result (model 6)

The estimation of Model 6 yielded a MLM chi square(422) value of 1297.489 (scaling correction factor = 1.116), with the remaining fit indices as follows: CFI = 0.957, TLI = 0.949, RMSEA = 0.038, and SRMR = 0.039. As might be expected given the newly specified path involving Factor 10, the difference in fit between this model (Model 6) and its predecessor (Model 5) was found to be statistically significant (MLM chi square[1] = 23.358). See tab 17.

An important finding here is that the alarming negative residual variance of Factor 10 finally disappeared. The estimated value for the newly specified path (F10 on F8) was statistically significant (Estimate = -0.884; Est/SE = -8.116). Furthermore, one of the previously determined non-significant paths loading from F4 onto F10 turned significant. Nonetheless, the regression path leading from F10 to F12, which has been consistently non-significant, remained so. In addition, however, the parameter, F8 on F7 (Peer Support → Self-Esteem), was found to be statistically non-significant. See tab 18.

3.6.4 Model mis-specification

Table 18: Residual variance of structural regression path and select factors for model6

Parameter*	B [†]	Beta [‡]	SE	Z	p-value
Strutural regression path					
F5DEC→F8SELF	0.357	0.770	0.051	6.964	0
F6SSUP→F8SELF	-0.118	-0.371	0.026	-4.479	0
F7PSUP→F8SELF	-0.014	-0.031	0.028	-0.492	0.623
F5DEC→F9ELC	-0.083	-0.143	0.031	-2.665	0.008
F2ROLC→F10EE	-1.168	-0.782	0.394	-2.963	0.003
F3WORK→F10EE	1.554	1.285	0.333	4.674	0
F4CLIM→F10EE	-0.568	-0.167	0.131	-4.351	0
F2ROLC→F11DP	0.167	0.132	0.047	3.545	0
F10EE→F11DP	0.307	0.360	0.032	9.583	0
F1ROLA→F12PA	0.368	0.314	0.087	4.211	0
F8SELF→F12PA	0.481	0.218	0.086	5.612	0
F9ELC→F12PA	-0.151	-0.086	0.055	-2.750	0.006
F10EE→F12PA	-0.014	-0.022	0.026	-0.563	0.573
F11DP→F12PA	-0.202	-0.261	0.031	-6.561	0
F4CLIM→F11DP	-0.945	-0.326	0.099	-9.526	0
F5DEC→F12PA	0.481	0.472	0.074	6.525	0
F2ROLC→F9ELC	0.166	0.297	0.028	5.920	0
F8SELF→F9ELC	-0.253	-0.202	0.048	-5.300	0
F8SELF→F10EE	-0.884	-0.264	0.109	-8.116	0
Selected factors					
F8SELF	0.087	0.751	0.008	11.198	0
F9ELC	0.13	0.719	0.010	12.649	0
F10EE	0.48	0.371	0.064	7.546	0
F11DP	0.545	0.581	0.045	12.053	0
F12PA	0.356	0.633	0.023	15.618	0

Note:

Values highlighted in red should be taken note of

* → indicates regression path

[†] Crude estimates

[‡] Standardized estimates

Table 19: Selected modification indices for initial model

Parameter	MI	EPC	std EPC	Logics
F11DP → F8SELF*	31.478	-0.076	-0.216	Sensible but not meaningfulfor the study
F2ROLC → F12PA†	22.794	0.339	0.345	Not sensible
F8SELF → F11DP‡	22.713	-0.409	-0.144	Sensible and meaningful

Note:

Parameter highlighted in red is selected for modification

* Higher depersonalization causes lower self-esteem

† Higher role conflict improves personal accomplishment

‡ Higher self-esteem causes lower depersonalization

```
#extract needed variables
MI.model6 <- modindices(sem6,
                        standardized = TRUE,
                        sort. = TRUE,
                        maximum.number = 50) |>
  filter(op == "~")

#adapt to publication style
MI.model6 <- MI.model6 |>
  mutate(Parameter = paste(rhs, "→", lhs)) |>
  select(Parameter, MI = mi, EPC = epc, "std EPC" = sepc.all) |>
  mutate("Logics" = c("Sensible but not meaningfulfor the study",
                      "Not sensible",
                      "Sensible and meaningful"))

#add footnote symbol to parameters
for (i in 1:nrow(MI.model6)){
  symbol <- c("","†","‡")
  Parameter <- unlist(MI.model6$Parameter)
  MI.model6$Parameter[i] <- paste0(Parameter[i], symbol[i])
}

#Visualize the table
MI.model6 |>
  kable(digits = 3,
        booktab = T,
        linesep = "",
        caption = "Selected modification indices for initial model") |>
  kable_styling(latex_options = "striped") |>
  row_spec(3, color = "red") |>
  footnote(general =
    "Parameter highlighted in red is selected for modification",
    symbol = c("Higher depersonalization causes lower self-esteem",
              "Higher role conflict improves personal accomplishment",
              "Higher self-esteem causes lower depersonalization"
    ))
```

Interestingly, Byrne's comments on F8→F11 is that "the parameter represents a causal path flowing from

F8 (Self-Esteem) to F11 (Depersonalization), and the fact that the EPC statistic has a negative sign makes interpretation of this path illogical as it conveys the notion that high self-esteem leads to low levels of depersonalization.” However, I was not able to make sense out of it since it seems to me that Higher self-esteem (a positive perception) lowers depersonalization (a negative perception) is a plausible and meaningful relation. People with strong confidence in themselves would less likely to feel being detached and numbness. Furthermore, Byrne believed $F2 \rightarrow F12$ (Higher role conflict improves personal accomplishment) “makes sense substantively”, which I was not able to grasp the idea. Actually, I believed this regression path is the most appropriate candidate for modification in current model, given the one with highest MI ($F11 \rightarrow F8$, Higher depersonalization causes lower self-esteem) was sensible but not digging into the problem under scope of the present study (We want it opposite). As such, at the point I decided to go for a different track than Byrne by modifying $F8 \rightarrow F11$ instead of $F2 \rightarrow F12$. See tab 19.

3.7 Post hos analysis (Model 7)

3.7.1 Compare SEM model 7 with preceding models

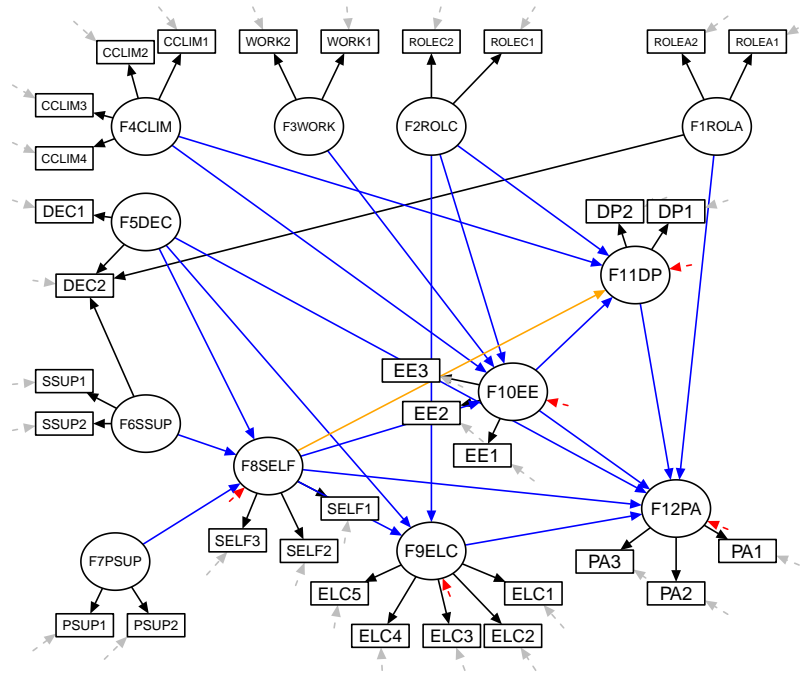
I defined model 7 as per the conclusion from last section: on the basis of model 6, I set the regression path leading from F8 to F11 (Higher role conflict improves personal accomplishment) free to estimate (Fig 11). The model fit indices and their comparison with the preceding models were tabulated in table 20.

```
#add new freely estimated parameter to the preceding model
model7 <- paste(model6, "\nF11DP ~ F8SELF")

# Estimate the model with the robust (MLM) estimator:
sem7 <- sem(model7, data = mbi, estimator = "MLM", mimic = "Mplus")

full.sem.diagram(model7, 20, "11", "model7")
```

Figure 11 Modified model (model7) of teacher burnout



Notes: Orange arrow is the regression path set free to estimate in the current model

```
sem <- c("sem1", "sem2", "sem3", "sem4", "sem5", "sem6", "sem7")
newpathstext <- c("Model1 + structural path 'F4→F11' set free to estimate",
  " Model2 + structural path 'F5→F12' set free to estimate",
  " Model3 + structural path 'F2→F9' set free to estimate",
  " Model5 + structural path 'F8→F9' set free to estimate",
  " Model6 + structural path 'F8→F10' set free to estimate",
  "Model7 + structural path 'F2→F12' set free to estimate")

model.comparison.table (sem,
  mycaption =
    "Comparison of new (model 7) and preceding models",
  newpathstext = newpathstext)
```

3.7.2 Examine the parameters of concern for model 7

```
concern.table(sem7, 20, "model7")
```

3.7.3 Comments on the results(model 7)

The estimation of Model 7 yielded a MLM chi square(421) value of 1267.208 (scaling correction factor = 1.116), with the remaining fit indices as follows: CFI = 0.958, TLI = 0.951, RMSEA = 0.038, and SRMR

Table 20: Comparison of new (model 7) and preceding models

	Chi-square	DF	p value	Δ Chi-square	CFI	TLI	RMSEA	SRMR	CSF
Model1	1554.942	427	0	—	0.945	0.936	0.043	0.051	1.117
Model2*	1450.985	426	0	91.67	0.950	0.941	0.041	0.046	1.117
Model3†	1406.517	425	0	46.866	0.952	0.944	0.040	0.044	1.117
Model4‡	1366.129	424	0	52.949	0.954	0.946	0.039	0.041	1.118
Model5§	1331.636	423	0	29.877	0.955	0.948	0.039	0.039	1.117
Model6¶	1297.489	422	0	23.358	0.957	0.949	0.038	0.039	1.116
Model7**	1276.208	421	0	23.057	0.958	0.951	0.038	0.038	1.116

* Model1 + structural path 'F4→F11' set free to estimate

† Model2 + structural path 'F5→F12' set free to estimate

‡ Model3 + structural path 'F2→F9' set free to estimate

§ Model5 + structural path 'F8→F9' set free to estimate

¶ Model6 + structural path 'F8→F10' set free to estimate

** Model7 + structural path 'F2→F12' set free to estimate

= 0.038. Again, the difference in fit between this model (Model 7) and its predecessor (Model 6) was found to be statistically significant (MLM chi square[1] = 23.057). See tab 20.

Here again, the new parameter added to the model (F11 on F8) was found to be statistically significant (Estimate = -0.42; Z-statistics = -4.688). However, the structural regression paths F8 (Self-Esteem) on F7 (Peer Support) and F12 (Personal Accomplishment) on F10 (Emotional exhaustion) remained non-signification. See tab 21.

At this point, I believe that I have exhausted the search for a better fitting model that best represents the data for secondary teachers. Thus, I start to delete regression path that are originally specified in the hypothesized model but remain statistically non-significant as per the results for Model 7. Accordingly, Model 8 is specified in which the two non-significant parameters noted above (F7 → F8; F10 → F12) are not included. Note that one of the excluded paths is different from Byrne's decision, where she deleted F12 on F1 instead of F12 on F10. See next section.

3.8 Final Model

3.8.1 Estimate the final model (model 8)

#add new freely estimated parameter to the preceding model

```
model8 <- '
  F1ROLA =~ ROLEA1 + ROLEA2 + DEC2
  F2ROLC =~ ROLEC1 + ROLEC2
  F3WORK =~ WORK1 + WORK2
  F4CLIM =~ CCLIM1 + CCLIM2 + CCLIM3 + CCLIM4
  F5DEC =~ DEC1 + DEC2
  F6SSUP =~ SSUP1 + SSUP2 + DEC2
  F7PSUP =~ PSUP1 + PSUP2
  F8SELF =~ SELF1 + SELF2 + SELF3
  F9ELC =~ ELC1 + ELC2 + ELC3 + ELC4 + ELC5
  F10EE =~ EE1 + EE2 + EE3
  F11DP =~ DP1 + DP2
  F12PA =~ PA1 + PA2 + PA3
```

Table 21: Residual variance of structural regression path and select factors for model7

Parameter*	B [†]	Beta [‡]	SE	Z	p-value
Strutural regression path					
F5DEC→F8SELF	0.347	0.751	0.050	6.905	0
F6SSUP→F8SELF	-0.113	-0.357	0.026	-4.373	0
F7PSUP→F8SELF	-0.01	-0.023	0.027	-0.367	0.713
F5DEC→F9ELC	-0.082	-0.143	0.031	-2.656	0.008
F2ROLC→F10EE	-1.185	-0.792	0.399	-2.969	0.003
F3WORK→F10EE	1.567	1.295	0.336	4.662	0
F4CLIM→F10EE	-0.576	-0.169	0.131	-4.386	0
F2ROLC→F11DP	0.139	0.109	0.047	2.927	0.003
F10EE→F11DP	0.256	0.301	0.033	7.803	0
F1ROLA→F12PA	0.37	0.314	0.088	4.208	0
F8SELF→F12PA	0.461	0.208	0.085	5.390	0
F9ELC→F12PA	-0.153	-0.086	0.055	-2.772	0.006
F10EE→F12PA	-0.018	-0.028	0.025	-0.747	0.455
F11DP→F12PA	-0.198	-0.255	0.031	-6.365	0
F4CLIM→F11DP	-0.934	-0.322	0.099	-9.434	0
F5DEC→F12PA	0.484	0.473	0.074	6.527	0
F2ROLC→F9ELC	0.166	0.297	0.028	5.920	0
F8SELF→F9ELC	-0.255	-0.204	0.048	-5.327	0
F8SELF→F10EE	-0.879	-0.262	0.108	-8.104	0
F8SELF→F11DP	-0.42	-0.147	0.089	-4.688	0
Selected factors					
F8SELF	0.087	0.756	0.008	11.233	0
F9ELC	0.13	0.719	0.010	12.665	0
F10EE	0.482	0.371	0.064	7.535	0
F11DP	0.538	0.572	0.045	11.840	0
F12PA	0.356	0.627	0.023	15.685	0

Note:

Values highlighted in red should be taken note of

* → indicates regression path

[†] Crude estimates

[‡] Standardized estimates

```

F8SELF ~ F5DEC + F6SSUP
F9ELC ~ F5DEC
F10EE ~ F2ROLC + F3WORK + F4CLIM
F11DP ~ F2ROLC + F10EE
F12PA ~ F1ROLA + F8SELF + F9ELC + F11DP
F11DP ~ F4CLIM
F12PA ~ F5DEC
F9ELC ~ F2ROLC
F9ELC ~ F8SELF
F10EE ~ F8SELF
F12PA ~ F2ROLC
'

# Estimate the model with the robust (MLM) estimator:
sem8 <- sem(model8, data = mbi, estimator = "MLM", mimic = "Mplus")
#summary(sem8, fit.measures = T, standardized = T)

```

I established the final model (model 8) by deleting two paths ($F7 \rightarrow F8$; $F10 \rightarrow F12$) from model 7. See picture 12

Estimation of this final model (Model 8) yielded an overall MLM chi square(423) value of 1291.217 (scaling correction factor = 1.116). This 423 degrees of freedom seems very suspicious since Byrne's final model also had 423 degrees of freedom. However, my final model deleted one less factor than hers. My intuition is the degree of freedom could be different too. Hence, I tried to manually count of the df of my final model to see if there was anything going wrong.

*My final number of manifest variable was 32. Hence I had $32 \times 32 / 2 = 528$ pieces of information**

I had 95 estimated parameters:

22 factor loadings ; 44 variances (7 factors, 32 error residual, 5 factor residual); 21 factor covariances (factor 1-7) 18 structural regression paths.

So I have $528 - (22 + 44 + 21 + 18) = 423$ degrees of freedom.

So nothing went wrong.

The final model have a CFI value of 0.957, representing a very good fit to the data for secondary teachers. The remaining TLI, RMSEA, and SRMR values further substantiate these results. In contrast to all previous models, it is also found that the corrected chi-square difference between the present model and Model 7 to be statistically significant (MLM chi square[1] = -14.863). See table 22.

It is also interesting to test the corrected chi-square difference between the final model and initial model:

```
print(chi.diff(sem1,sem8))
```

```
## [1] 240.081
```

A chi square value of 240.081 indicated significant difference. The final model fit the data significantly better than the initial model.

```

sem <- c("sem1", "sem2", "sem3", "sem4", "sem5", "sem6", "sem7", "sem8")
newpathstext <- c(" Model1 + structural path 'F4→F11' set free to estimate",
                  " Model2 + structural path 'F5→F12' set free to estimate",
                  " Model3 + structural path 'F2→F9' set free to estimate",
                  " Model5 + structural path 'F8→F9' set free to estimate",

```

Table 22: Comparison of final(model 8) and preceding models

	Chi-square	DF	p value	Δ Chi-square	CFI	TLI	RMSEA	SRMR	CSF
Model1	1554.942	427	0	—	0.945	0.936	0.043	0.051	1.117
Model2*	1450.985	426	0	91.67	0.950	0.941	0.041	0.046	1.117
Model3†	1406.517	425	0	46.866	0.952	0.944	0.040	0.044	1.117
Model4‡	1366.129	424	0	52.949	0.954	0.946	0.039	0.041	1.118
Model5§	1331.636	423	0	29.877	0.955	0.948	0.039	0.039	1.117
Model6¶	1297.489	422	0	23.358	0.957	0.949	0.038	0.039	1.116
Model7**	1276.208	421	0	23.057	0.958	0.951	0.038	0.038	1.116
Model8††	1291.217	423	0	-14.863	0.957	0.950	0.038	0.039	1.116

* Model1 + structural path 'F4→F11' set free to estimate

† Model2 + structural path 'F5→F12' set free to estimate

‡ Model3 + structural path 'F2→F9' set free to estimate

§ Model5 + structural path 'F8→F9' set free to estimate

¶ Model6 + structural path 'F8→F10' set free to estimate

** Model7 + structural path 'F2→F12' set free to estimate

†† Model8 - structural path with F7 → F8 and F10 → F12 set fixed

```

      " Model6 + structural path 'F8→F10' set free to estimate",
      "Model7 + structural path 'F2→F12' set free to estimate",
      "Model8 - structural path with F7 → F8 and F10 → F12 set fixed")

model.comparison.table (sem,
                        mycaption =
                          "Comparison of final(model 8) and preceding models",
                        newpathstext = newpathstext)

```

3.8.2 Visualize the final model

For the sake of plotting, I deleted the F7 with two of its indicators from the model (which I shouldn't, since they still have a position in the measurement model, but I haven't found a better way to hide a node using semplot). However, this is only for demonstrating the structure of final model. It won't cause any misunderstanding, I think.

Figure 12 displays the final model, which highlights in orange all the regression paths added in addition to the initial model.

```

#broom cells
m[53,9] <- NA
m[59,6] <- NA
m[59,13] <- NA

model.plot <- '
  F1ROLA =~ ROLEA1 + ROLEA2 + DEC2
  F2ROLA =~ ROLEC1 + ROLEC2
  F3WORK =~ WORK1 + WORK2
  F4CLIM =~ CCLIM1 + CCLIM2 + CCLIM3 + CCLIM4
  F5DEC =~ DEC1 + DEC2

```

```

F6SSUP =~ SSUP1 + SSUP2 + DEC2
F8SELF =~ SELF1 + SELF2 + SELF3
F9ELC =~ ELC1 + ELC2 + ELC3 + ELC4 + ELC5
F10EE =~ EE1 + EE2 + EE3
F11DP =~ DP1 + DP2
F12PA =~ PA1 + PA2 + PA3
F8SELF ~ F5DEC + F6SSUP
F9ELC ~ F5DEC
F10EE ~ F2ROLC + F3WORK + F4CLIM
F11DP ~ F2ROLC + F10EE
F12PA ~ F1ROLA + F8SELF + F9ELC + F11DP
F11DP ~ F4CLIM
F12PA ~ F5DEC
F9ELC ~ F2ROLC
F9ELC ~ F8SELF
F10EE ~ F8SELF
F12PA ~ F2ROLC

```

Estimate the model with the robust (MLM) estimator:

```

sem.plot <- sem(model.plot, data = mbi, estimator = "MLM", mimic = "Mplus")
#summary(sem8, fit.measures = T, standardized = T)

```

```

semPaths(semPlotModel(model.plot),
  style = "lisrel",
  rotation = 2,
  sizeLat = 6,
  sizeLat2 = 5,
  sizeMan = 5,
  sizeMan2 = 2,
  residScale = 4,
  shapeMan = "rectangle",
  edge.color = c(rep("black",32),
                 rep("blue",14),
                 rep("orange",4),
                 rep("gray",32),
                 rep("red",5)),
  residuals = T,
  layout = m,
  nCharNodes=0,
  optimizeLatRes = T,
  exoVar = F)
title(main = list("Figure 12. Final model of teacher burnout",
  cex = 1.5, font = 1),
  outer = F, line = -1)
title(sub = "Notes: Red arrow indicates factor residuals; gray arrow indicates error residuals;
  black arrow indicates factor loading; blue arrow indicates originally hypothesized
  regression path; orange arrow indicates regression paths added over pos hoc analysis",
  line = 0, adj = 0.7)

```


Figure 12. Final model of teacher burnout

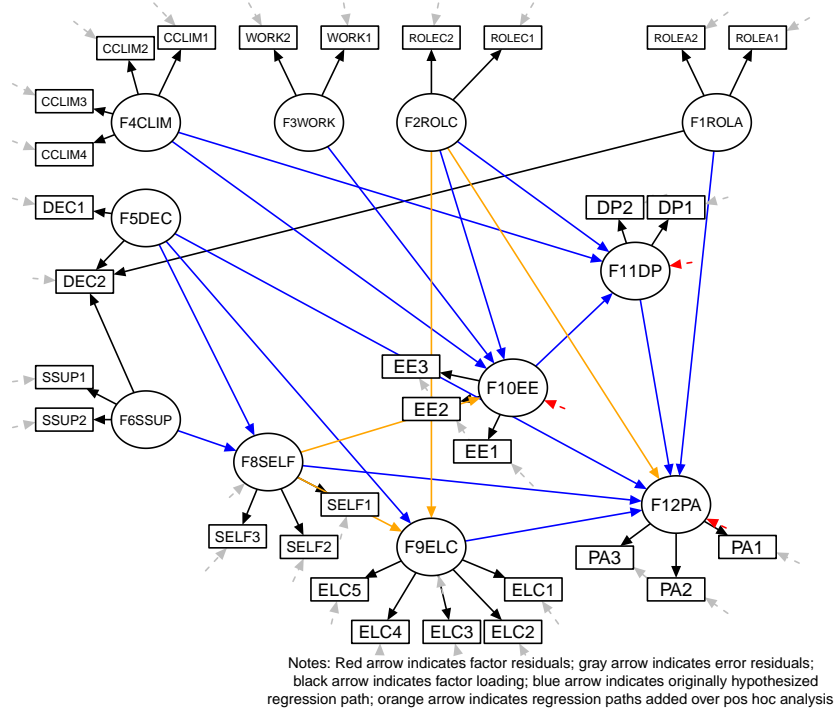


Figure 13 displays the theoretical structure of the final model, showing how the psychological determinants influencing teacher's burnout.

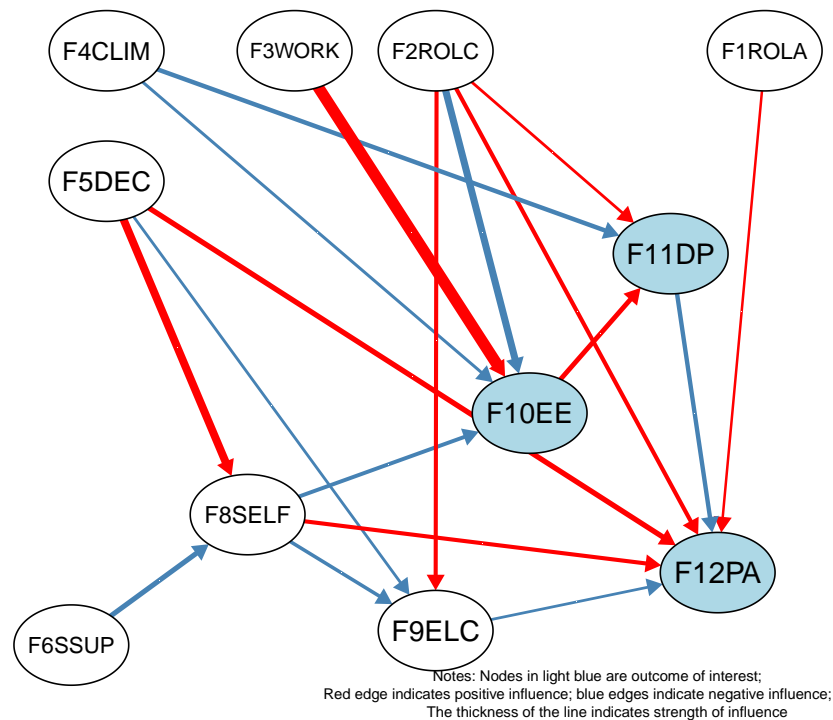
```
grps <- list(c("F10EE", "F11DP", "F12PA"),
             c("F1ROLA", "F2ROLC", "F3WORK", "F4CLIM", "F5DEC", "F6SSUP", "F8SELF", "F9ELC"))
a <- "Figure 13. Summary of final model structural paths representing determinants
of burnout for teachers"
m[40,12] <- NA
m[53,9] <- "F6SSUP"
semPaths(sem8,
  style = "lisrel",
  #"path",
  #"par",
  "std",
  rotation = 2,
  sizeLat = 10,
  sizeLat2 = 7,
  sizeMan = 5,
  sizeMan2 = 2,
  residScale = 4,
  shapeMan = "rectangle",
  #edge.color = c(rep("black",5),
  #               rep("blue",5),
  #               rep("gray",4)),
  residuals = F,
```

```

layout = m,
nCharNodes=0,
optimizeLatRes = T,
exoVar = F,
covAtResiduals = F,
intercepts = F,
exoCov = F,
structural = T,
posCol = c("red","red"),
negCol = c("steelblue","steelblue"),
fade = F,
edge.label.cex = 0.01,
group =grps,
color = c("lightblue", "white")
)
title(main = list(a,
                  cex = 1.5, font =1),
      outer = F)
title(sub = "Notes: Nodes in light blue are outcome of interest;
           Red edge indicates positive influence; blue edges indicate negative influence;
           The thickness of the line indicates strength of influence",
      line = 0, adj = 0.7)

```

Figure 13. Summary of final model structural paths representing determinants of burnout for teachers



4 Summary

In the exercise, I practiced post-hoc analysis of full SEM with teacher burnout data set. Over the process, I established 8 models step by step and compared their model fit indices, regression path estimates, and selected alarming variances;

Over the exercises, I mostly followed the step of the slides and Byrne's book. However, in models 7 and 8, I started to go on a different tract, leading to a different final model.

I learnt something that I would definitely misunderstand or mistake as I know if I hadn't done this exercise. Like, when removing factors for the final model, we should keep the factor loading (measurement model) of the deleted factors in the model;

There are some visual effects I want to realize using Semplot that I finally failed to, such as showing the edge as broken arrows. Need to dig more next week.

Calculating degrees of freedom is very challenging. I understand every bit of it when the process is shown to me. However, when I need to calculate without any external aide, I always forgot this or that.

I found SEM is a beautiful blending of statistics and mindful sense-making. Indeed, they always go together. Yet, not often so well balanced as they are in SEM, for the scope of statistics I am familiar with.