

REIMS sample analysis

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Set up

Set up MPlusAutomation

```
install.packages("devtools")
library(devtools)

install_github("michaelhallquist/MplusAutomation")
```

Load packages

```
## Version: 1.2
## We work hard to write this free software. Please help us get credit by citing:
##
## Hallquist, M. N. & Wiley, J. F. (2018). MplusAutomation: An R Package for Facilitating Large-Scale L
##
## -- see citation("MplusAutomation").

## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.4      v readr      2.1.5
## v forcats    1.0.0      v stringr   1.5.1
## v ggplot2    3.5.0      v tibble    3.2.1
## v lubridate  1.9.3      v tidyr     1.3.1
## v purrr      1.0.2
## -- Conflicts ----- tidyverse_conflicts() --
## x tidyr::extract() masks MplusAutomation::extract()
## x dplyr::filter()  masks stats::filter()
## x dplyr::lag()     masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
## here() starts at /Users/nathanalexander/Dropbox/Projects/immerse
##
##
## Attaching package: 'data.table'
##
##
## The following objects are masked from 'package:lubridate':
##
##      hour, isoweek, mday, minute, month, quarter, second, wday, week,
##      yday, year
```

```
##
##
## The following objects are masked from 'package:dplyr':
##
##   between, first, last
##
##
## The following object is masked from 'package:purrr':
##
##   transpose
##
##
## here() starts at /Users/nathanalexander/Dropbox/Projects/immerse/reims
```

Data

```
# set data
reims <- read_csv(here("data", "reims_clean.csv"))
```

```
## New names:
## Rows: 103 Columns: 57
## -- Column specification
## ----- Delimiter: "," dbl
## (57): groupflag, age, sex, race, mathperson1, mathperson2, mathperson3, ...
## i Use 'spec()' to retrieve the full column specification for this data. i
## Specify the column types or set 'show_col_types = FALSE' to quiet this message.
## * 'belongrace...19' -> 'belongrace'
```

```
# inspect data
reims
```

```
## # A tibble: 103 x 57
##   groupflag age sex race mathperson1 mathperson2 mathperson3 mathperson4
##   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1      1    18  2    6      0      0      0      0
## 2      1    22  2    1      0      0      0      0
## 3      1    20  1    7      1      1      1      1
## 4      1    20  1    3      1      1      1      1
## 5      1    18  1    4      0      0      0      1
## 6      1    18  2    6      0      0      0      0
## 7      1    19  1    6      0      0      0      0
## 8      1    23  2    4      0      0      0      0
## 9      1    37  2    3      0      0      0      0
## 10     1    27  2    6      0      0      0      0
## # i 93 more rows
## # i 49 more variables: dislikemathclass <dbl>, pursuestem <dbl>,
## #   boysbetter <dbl>, learnrace <dbl>, racegroups <dbl>, knowrace <dbl>,
## #   connectrace1 <dbl>, affectrace <dbl>, proudrace <dbl>, mixrace <dbl>,
## #   unclarrace <dbl>, connectrace2 <dbl>, dontknowrace <dbl>,
## #   belongrace <dbl>, understandrace <dbl>, talkrace <dbl>, priderace <dbl>,
## #   avoidrace <dbl>, practicerace <dbl>, playotherrace <dbl>, ...
```

```
summary(reims)
```

```
##      groupflag      age      sex      race
## Min.   :0.0000  Min.   :14.00  Min.    :1.00  Min.    :1.00
## 1st Qu.:1.0000  1st Qu.:18.00  1st Qu.:1.00  1st Qu.:3.00
## Median :1.0000  Median :20.00  Median :2.00  Median :4.00
## Mean   :0.8835  Mean   :20.82  Mean    :1.65  Mean    :3.99
## 3rd Qu.:1.0000  3rd Qu.:21.00  3rd Qu.:2.00  3rd Qu.:6.00
## Max.    :1.0000  Max.    :47.00  Max.    :3.00  Max.    :7.00
## mathperson1  mathperson2  mathperson3  mathperson4
## Min.   :0.0000  Min.   :0.0000  Min.   :0.0000  Min.   :0.0000
## 1st Qu.:0.0000  1st Qu.:0.0000  1st Qu.:0.0000  1st Qu.:0.0000
## Median :1.0000  Median :1.0000  Median :1.0000  Median :1.0000
## Mean   :0.5922  Mean   :0.5631  Mean   :0.5146  Mean   :0.5146
## 3rd Qu.:1.0000  3rd Qu.:1.0000  3rd Qu.:1.0000  3rd Qu.:1.0000
## Max.    :1.0000  Max.    :1.0000  Max.    :1.0000  Max.    :1.0000
## dislikemathclass  pursuestem  boysbetter  learnrace
## Min.   :0.0000  Min.   :0.0000  Min.   :0.00000  Min.   :0.0000
## 1st Qu.:0.0000  1st Qu.:0.0000  1st Qu.:0.00000  1st Qu.:0.0000
## Median :0.0000  Median :1.0000  Median :0.00000  Median :0.0000
## Mean   :0.2524  Mean   :0.5534  Mean   :0.07767  Mean   :0.4078
## 3rd Qu.:0.5000  3rd Qu.:1.0000  3rd Qu.:0.00000  3rd Qu.:1.0000
## Max.    :1.0000  Max.    :1.0000  Max.    :1.00000  Max.    :1.0000
## racegroups      knowrace      connectrace1  affectrace
## Min.   :0.0000  Min.   :0.0000  Min.   :0.0000  Min.   :0.0000
## 1st Qu.:0.0000  1st Qu.:0.0000  1st Qu.:1.0000  1st Qu.:0.0000
## Median :0.0000  Median :1.0000  Median :1.0000  Median :0.0000
## Mean   :0.3301  Mean   :0.6311  Mean   :0.8641  Mean   :0.4563
## 3rd Qu.:1.0000  3rd Qu.:1.0000  3rd Qu.:1.0000  3rd Qu.:1.0000
## Max.    :1.0000  Max.    :1.0000  Max.    :1.0000  Max.    :1.0000
## proudrace      mixrace      unclarrace      connectrace2
## Min.   :0.0000  Min.   :0.00000  Min.   :0.0000  Min.   :0.0000
## 1st Qu.:1.0000  1st Qu.:0.00000  1st Qu.:0.0000  1st Qu.:0.0000
## Median :1.0000  Median :0.00000  Median :0.0000  Median :1.0000
## Mean   :0.8252  Mean   :0.08738  Mean   :0.2427  Mean   :0.6602
## 3rd Qu.:1.0000  3rd Qu.:0.00000  3rd Qu.:0.0000  3rd Qu.:1.0000
## Max.    :1.0000  Max.    :1.00000  Max.    :1.0000  Max.    :1.0000
## dontknowrace  belongrace  understandrace  talkrace
## Min.   :0.0000  Min.   :0.0000  Min.   :0.0000  Min.   :0.0000
## 1st Qu.:0.0000  1st Qu.:0.0000  1st Qu.:0.0000  1st Qu.:0.0000
## Median :0.0000  Median :1.0000  Median :1.0000  Median :0.0000
## Mean   :0.3107  Mean   :0.5146  Mean   :0.5437  Mean   :0.4563
## 3rd Qu.:1.0000  3rd Qu.:1.0000  3rd Qu.:1.0000  3rd Qu.:1.0000
## Max.    :1.0000  Max.    :1.0000  Max.    :1.0000  Max.    :1.0000
## priderace      avoidrace      practicerace  playotherrace
## Min.   :0.0000  Min.   :0.00000  Min.   :0.0000  Min.   :0.000
## 1st Qu.:0.0000  1st Qu.:0.00000  1st Qu.:0.0000  1st Qu.:1.000
## Median :1.0000  Median :0.00000  Median :1.0000  Median :1.000
## Mean   :0.5631  Mean   :0.05825  Mean   :0.5049  Mean   :0.767
## 3rd Qu.:1.0000  3rd Qu.:0.00000  3rd Qu.:1.0000  3rd Qu.:1.000
## Max.    :1.0000  Max.    :1.00000  Max.    :1.0000  Max.    :1.000
## strongrace    enjoyotherrace  feelrace  racediscrimination
## Min.   :0.0000  Min.   :0.0000  Min.   :0.0000  Min.   :0.0000
```

## 1st Qu.:	0.0000	1st Qu.:	1.0000	1st Qu.:	0.0000	1st Qu.:	0.0000
## Median :	1.0000	Median :	1.0000	Median :	1.0000	Median :	0.0000
## Mean :	0.5049	Mean :	0.8932	Mean :	0.7184	Mean :	0.4369
## 3rd Qu.:	1.0000	3rd Qu.:	1.0000	3rd Qu.:	1.0000	3rd Qu.:	1.0000
## Max. :	1.0000	Max. :	1.0000	Max. :	1.0000	Max. :	1.0000
## academicability_peers		mathability_peers		activities		deciderules	
## Min. :	0.0000	Min. :	0.0000	Min. :	0.000	Min. :	0.0000
## 1st Qu.:	0.0000	1st Qu.:	0.0000	1st Qu.:	0.000	1st Qu.:	0.0000
## Median :	0.0000	Median :	1.0000	Median :	1.000	Median :	0.0000
## Mean :	0.4854	Mean :	0.5146	Mean :	0.534	Mean :	0.2039
## 3rd Qu.:	1.0000	3rd Qu.:	1.0000	3rd Qu.:	1.000	3rd Qu.:	0.0000
## Max. :	1.0000	Max. :	1.0000	Max. :	1.000	Max. :	1.0000
## makeadiff		adultcares		adultnotices		adultlistens	
## Min. :	0.0000	Min. :	0.0000	Min. :	0.0000	Min. :	0.000
## 1st Qu.:	0.0000	1st Qu.:	0.0000	1st Qu.:	0.0000	1st Qu.:	0.000
## Median :	0.0000	Median :	1.0000	Median :	0.0000	Median :	1.000
## Mean :	0.4175	Mean :	0.5437	Mean :	0.4951	Mean :	0.699
## 3rd Qu.:	1.0000	3rd Qu.:	1.0000	3rd Qu.:	1.0000	3rd Qu.:	1.000
## Max. :	1.0000	Max. :	1.0000	Max. :	1.0000	Max. :	1.000
## adultpraise		adultmybest		adultmysuccess		mtchmematter	
## Min. :	0.0000	Min. :	0.0000	Min. :	0.0000	Min. :	0.0000
## 1st Qu.:	0.0000	1st Qu.:	0.0000	1st Qu.:	0.0000	1st Qu.:	0.0000
## Median :	1.0000	Median :	1.0000	Median :	1.0000	Median :	0.0000
## Mean :	0.6699	Mean :	0.7087	Mean :	0.7087	Mean :	0.1456
## 3rd Qu.:	1.0000	3rd Qu.:	1.0000	3rd Qu.:	1.0000	3rd Qu.:	0.0000
## Max. :	1.0000	Max. :	1.0000	Max. :	1.0000	Max. :	1.0000
## mtchlowerstandards		mtchraceexpect		mtchtalkrace		mtchignoreraace	
## Min. :	0.00000	Min. :	0.00000	Min. :	0.0000	Min. :	0.0000
## 1st Qu.:	0.00000	1st Qu.:	0.00000	1st Qu.:	0.0000	1st Qu.:	0.0000
## Median :	0.00000	Median :	0.00000	Median :	0.0000	Median :	0.0000
## Mean :	0.05825	Mean :	0.05825	Mean :	0.2524	Mean :	0.1748
## 3rd Qu.:	0.00000	3rd Qu.:	0.00000	3rd Qu.:	0.5000	3rd Qu.:	0.0000
## Max. :	1.00000	Max. :	1.00000	Max. :	1.0000	Max. :	1.0000
## mtchvalues		mtchrespect		mtchfair		mtchsuccess	
## Min. :	0.000	Min. :	0.0000	Min. :	0.000	Min. :	0.0000
## 1st Qu.:	1.000	1st Qu.:	1.0000	1st Qu.:	1.000	1st Qu.:	1.0000
## Median :	1.000	Median :	1.0000	Median :	1.000	Median :	1.0000
## Mean :	0.835	Mean :	0.8641	Mean :	0.835	Mean :	0.8155
## 3rd Qu.:	1.000	3rd Qu.:	1.0000	3rd Qu.:	1.000	3rd Qu.:	1.0000
## Max. :	1.000	Max. :	1.0000	Max. :	1.000	Max. :	1.0000
## mtchmistakesok		mtchbiased		mtchmadeinteresting		mtchgendebias	
## Min. :	0.0000	Min. :	0.0000	Min. :	0.0000	Min. :	0.00000
## 1st Qu.:	1.0000	1st Qu.:	0.0000	1st Qu.:	0.0000	1st Qu.:	0.00000
## Median :	1.0000	Median :	0.0000	Median :	1.0000	Median :	0.00000
## Mean :	0.8252	Mean :	0.1845	Mean :	0.6602	Mean :	0.06796
## 3rd Qu.:	1.0000	3rd Qu.:	0.0000	3rd Qu.:	1.0000	3rd Qu.:	0.00000
## Max. :	1.0000	Max. :	1.0000	Max. :	1.0000	Max. :	1.00000
## mtchmadeeasy							
## Min. :	0.0000						
## 1st Qu.:	0.0000						
## Median :	1.0000						
## Mean :	0.5922						
## 3rd Qu.:	1.0000						
## Max. :	1.0000						

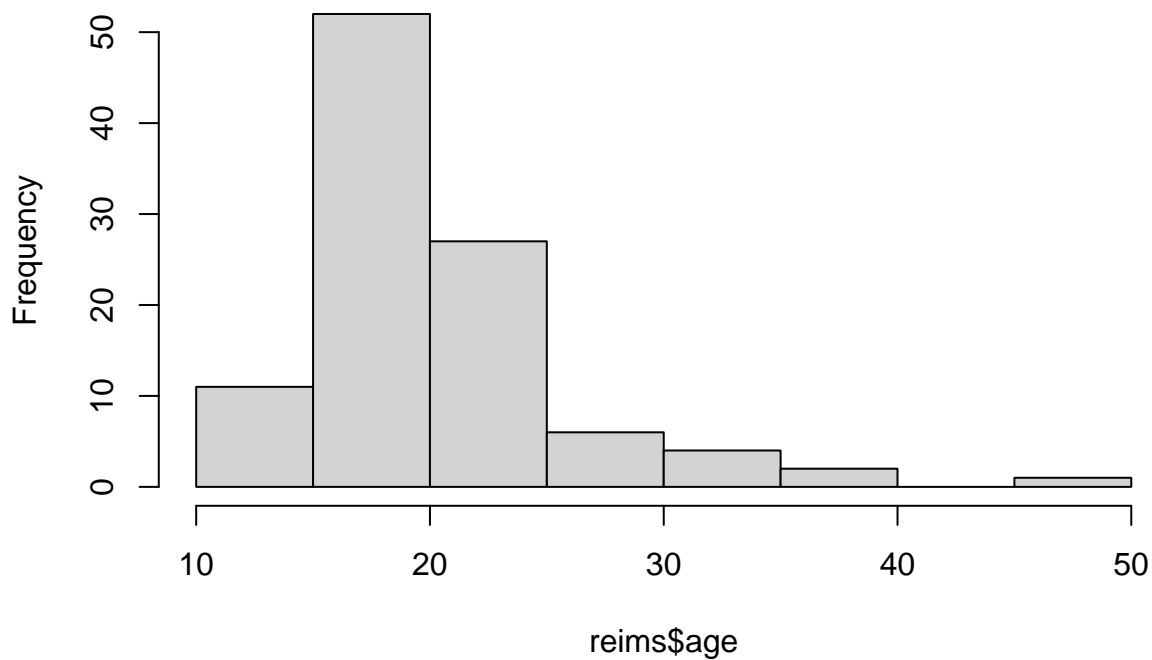
Make updates/edits to data We observe some of the other tags on our data. We also look at the distribution of some values. Take note that we changed the variable names (thanks Dina!) to less than eight characters so that our model could run.

```
# view distribution of indicator variables  
table(reims$groupflag)
```

```
##  
## 0 1  
## 12 91
```

```
hist(reims$age)
```

Histogram of reims\$age



```
table(reims$sex)
```

```
##  
## 1 2 3  
## 37 65 1
```

```
table(reims$race)
```

```
##  
## 1 2 3 4 5 6 7  
## 23 1 12 29 1 30 7
```

```

# subset data
reims1 <- reims %>%
  select(mathperson1, mathperson2, mathperson3, mathperson4, dislikemathclass, pursuestem, boysbetter) %>%
  rename(m1 = mathperson1,
         m2 = mathperson2,
         m3 = mathperson3,
         m4 = mathperson4,
         dislike = dislikemathclass,
         pursue = pursuestem,
         boys = boysbetter)

head(reims1, n=10)

```

```

## # A tibble: 10 x 7
##       m1      m2      m3      m4 dislike pursue  boys
##   <dbl> <dbl> <dbl> <dbl>   <dbl>   <dbl> <dbl>
## 1     0     0     0     0       0     0     0
## 2     0     0     0     0       1     0     0
## 3     1     1     1     1       0     0     0
## 4     1     1     1     1       1     1     1
## 5     0     0     0     1       0     0     0
## 6     0     0     0     0       1     0     0
## 7     0     0     0     0       0     1     0
## 8     0     0     0     0       0     0     0
## 9     0     0     0     0       1     0     0
## 10    0     0     0     0       1     1     0

```

```

## add ids and covariates; tell mplus that what we are

```

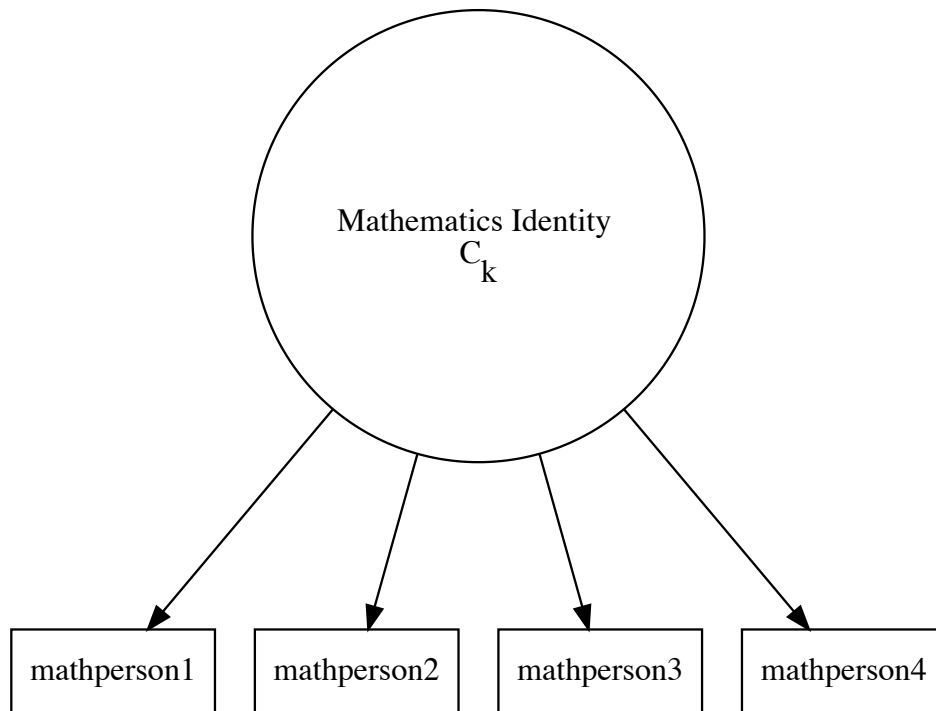
```

# use the reorder function to get the variables that you want to model in the output.

```

Models

Model 1



Model 1 MplusAutomation code

```
input <- mplusObject(  
  TITLE = "REIMS Mathematics Identity Model 1",  
  VARIABLE = "categorical = m1 m2 m3 m4;  
  usevar = m1-m4;  
  classes = c(3);",  
  
  ANALYSIS =  
    "estimator = mlr;  
    type = mixture;",  
  
  OUTPUT = "tech11 tech14;",  
  
  PLOT = "type = plot3;  
    series = m1-m4(*);",  
  
  usevariables = colnames(reims1),  
  rdata = reims1)  
  
output <- mplusModeler(input,  
  dataout = here("mplus", "reims1.dat"),  
  modelout = here("mplus", "reims1.inp"),  
  check = T, run = T, hashfilename = F)
```

```
## When hashfilename = FALSE, writeData cannot be 'ifmissing', setting to 'always'

## The following lines are not empty and do not end in a : or ;.
## 2: REIMS Mathematics Identity Model 1
## 4: FILE = "/Users/nathanalexander/Dropbox/Projects/immerse/reims/mplus/

## Rerun with parseMplus(add = TRUE) to add semicolons to all lines

## The file(s)
## 'reims1.dat'
## currently exist(s) and will be overwritten

## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.
```

Take a look at the item probability plot:

```
source(here("plot_lca.txt")) # custom function created by Dina to plot our lsa output
```

```
##
## Attaching package: 'reshape2'

## The following objects are masked from 'package:data.table':
##
## dcast, melt

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

##
## Attaching package: 'cowplot'

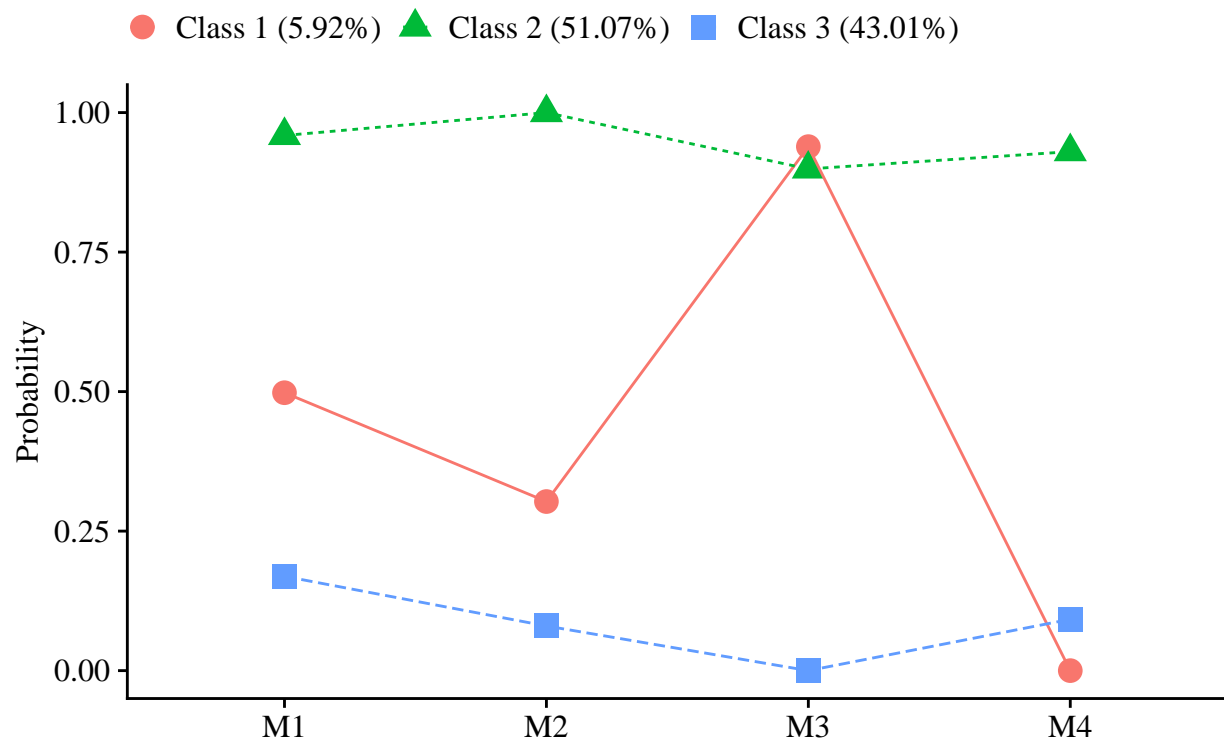
## The following object is masked from 'package:lubridate':
##
## stamp
```

```
model11 <- readModels(here("mplus", "reims1.out")) # read in output
```

```
## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.
```

```
plot_lca(model_name = model11) # there is an error with the non atomic measure columns
```

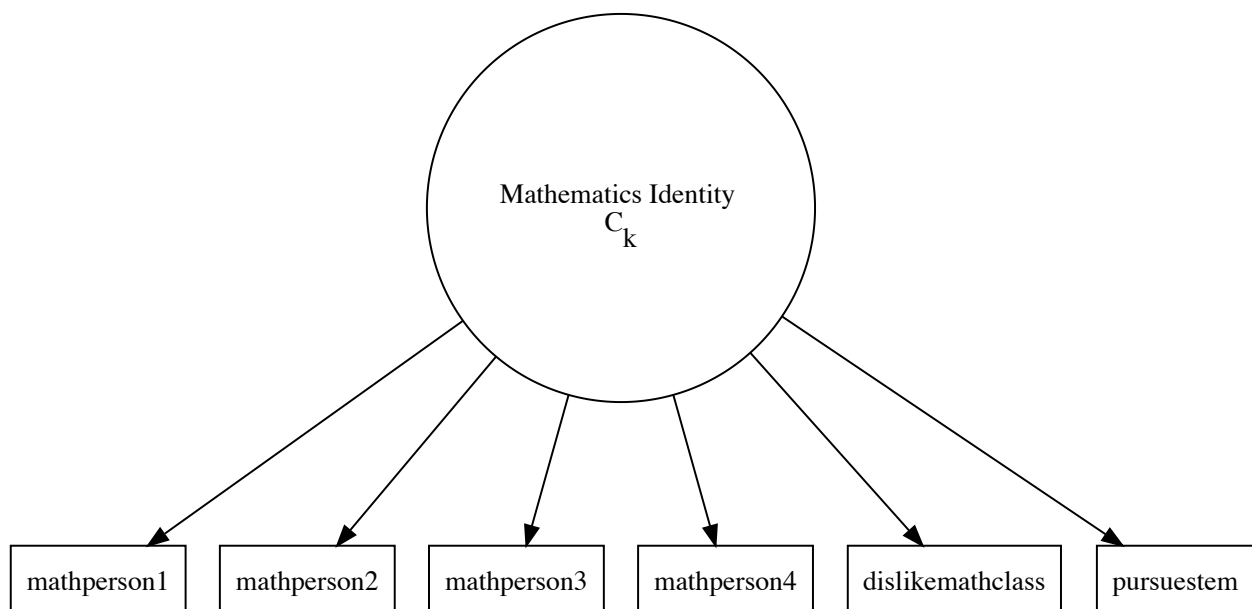

REIMS Mathematics Identity Model 1 Probability Plot



Show probability plot of data and observe the different classes.

Model 2

Let's run a four class model and add three variables.



Model 2 MplusAutomation code

```
input <- mplusObject(
  TITLE = "REIMS Mathematics Identity Model 2",
  VARIABLE = "categorical = m1 m2 m3 m4 dislike pursue;
  usevar = m1-pursue;
  classes = c(4);",

  ANALYSIS =
    "estimator = mlr;
    type = mixture;",

  OUTPUT = "tech11 tech14;",

  PLOT = "type = plot3;
    series = m1-pursue(*);",

  usevariables = colnames(reims1),
  rdata = reims1)

output <- mplusModeler(input,
  dataout = here("mplus", "reims1.dat"),
  modelout = here("mplus", "reims1.inp"),
  check = T, run = T, hashfilename = F)
```

When hashfilename = FALSE, writeData cannot be 'ifmissing', setting to 'always'

The following lines are not empty and do not end in a : or ;.

2: REIMS Mathematics Identity Model 2

4: FILE = "/Users/nathanalexander/Dropbox/Projects/immerse/reims/mplus/

Rerun with parseMplus(add = TRUE) to add semicolons to all lines

The file(s)

'reims1.dat'

currently exist(s) and will be overwritten

No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.

Take a look at the item probability plot:

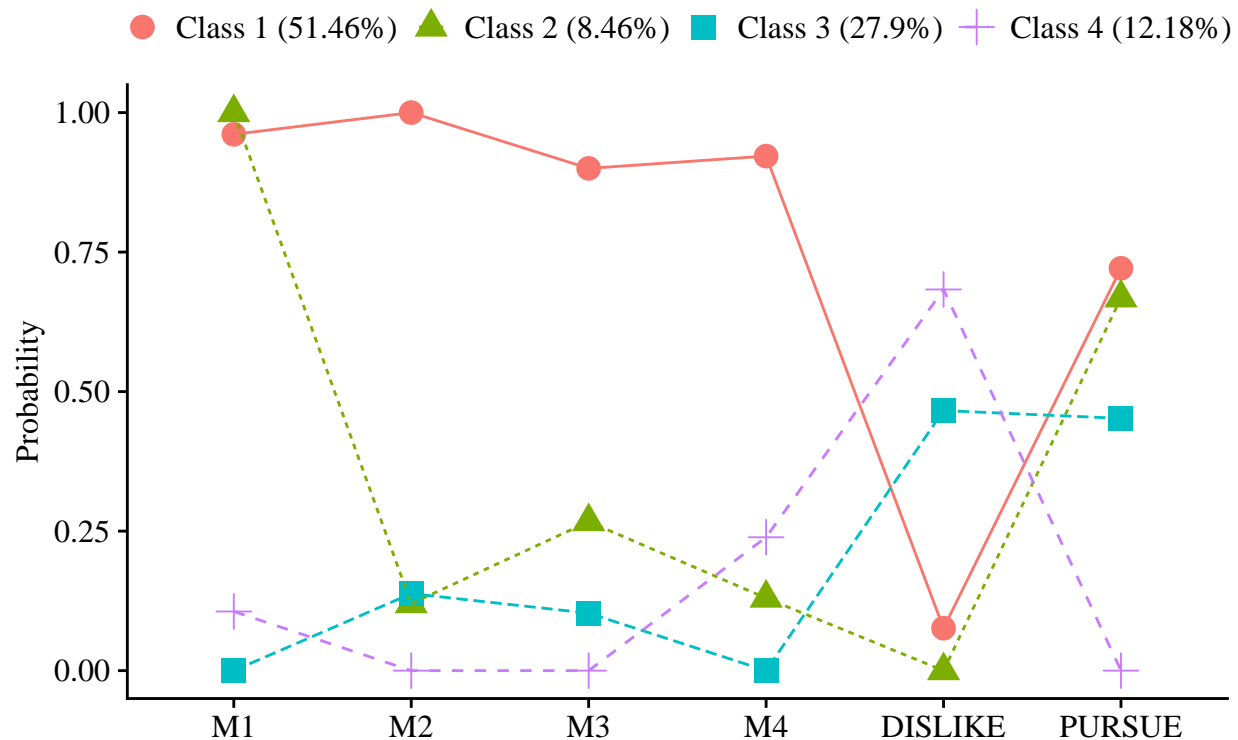
```
source(here("plot_lca.txt")) # custom function created by Dina to plot our lsa output
```

```
model2 <- readModels(here("mplus", "reims1.out")) # read in output
```

No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.

```
plot_lca(model_name = model2) # there is an error with the non atomic measure columns
```

REIMS Mathematics Identity Model 2 Probability Plot



Show probability plot of data and observe the different classes.

Enumeration

We use the `mplusObject` function in the `MPlusAutomation` package and saves all models run.

```
# add new libraries
library(cowplot)
library(glue)
```

Proportion of indicators using R:

```
# set up data to find proportions of binary indicators
df <- reims1 %>%
  pivot_longer(c(m1, m2, m3, m4, dislike, pursue),
    names_to = "Variable")

# create table of variables and counts
t1 <- table(df$Variable, df$value)

# find proportions and round to 3 decimal places
prop <- prop.table(t1, margin = 1) %>%
  round(3)
```

```

# combine everything to one table
dframe <- data.frame(Variables=rownames(t1), Proportion=prop[,2], Count=t1[,2])

# remove row names
row.names(dframe) <- NULL

# Make it a gt() table
prop_table <- dframe %>%
  gt()
prop_table

```

Variables	Proportion	Count
dislike	0.252	26
m1	0.592	61
m2	0.563	58
m3	0.515	53
m4	0.515	53
pursue	0.553	57

```

# save as a word doc
gtsave(prop_table, here("figures", "prop_table.docx"))

```

Use an enumeration function

```

lca_4 <- lapply(1:4, function(k) {
  lca_enum <- mplusObject(

    TITLE = glue("{k}-Class"),

    VARIABLE = glue(
      "categorical m1-pursue;
      usevar = m1-pursue;
      classes = c({k});"),

    ANALYSIS =
      "estimator = mlr;
      type = mixture;
      starts = 500 100;",

    OUTPUT = "tech11 tech14 svalues;",

    usevariables = colnames(reims1),
    rdata = reims1)

  lca_enum_fit <- mplusModeler(lca_enum,
    dataout = glue(here("enum", "reims1.dat")),
    modelout = glue(here("enum", "c{k}_reims1.inp")),
    check = T, run = T, hashfilename = F)})

```

table of fit

We want to begin by extracting the data:

```
output_reims1 <- readModels(here("enum"))
```

```
## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.
## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.
## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.
## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.
```

```
enum_extract <- LatexSummaryTable(
  output_reims1,
  keepCols = c(
    "Title",
    "Parameters",
    "LL",
    "BIC",
    "aBIC",
    "BLRT_PValue",
    "T11_VLMR_PValue",
    "Observations"
  ),
  sortBy = "Title"
) # select first set of models (Class 1 through 4)

allFit <- enum_extract %>%
  mutate(CAIC = -2 * LL + Parameters * (log(Observations) + 1)) %>%
  mutate(AWE = -2 * LL + 2 * Parameters * (log(Observations) + 1.5)) %>%
  mutate(SIC = -.5 * BIC) %>%
  mutate(expSIC = exp(SIC - max(SIC))) %>%
  mutate(BF = exp(SIC - lead(SIC))) %>%
  mutate(cmPk = expSIC / sum(expSIC)) %>%
  dplyr::select(1:5, 9:10, 6:7, 13, 14) %>%
  arrange(Parameters)
```

Then we create a table:

```
fit_table <- allFit %>%
  gt() %>%
  tab_header(title = md("***Model Fit Summary Table**")) %>%
  cols_label(
    Title = "Classes",
    Parameters = md("Par"),
    LL = md("*LL*"),
    T11_VLMR_PValue = "VLMR",
    BLRT_PValue = "BLRT",
    BF = md("BF"),
    cmPk = md("*cmPk*")
  ) %>%
  tab_footnote(
    footnote = md(
      "*Note.* Par = Parameters; *LL* = model log likelihood;
      BIC = Bayesian information criterion;
      aBIC = sample size adjusted BIC; CAIC = consistent Akaike information criterion;
      AWE = approximate weight of evidence criterion;
```

```

BLRT = bootstrapped likelihood ratio test p-value;
VLMR = Vuong-Lo-Mendell-Rubin adjusted likelihood ratio test p-value;
*cmPk* = approximate correct model probability."
),
locations = cells_title()
) %>%
tab_options(column_labels.font.weight = "bold") %>%
fmt_number(c(3:7),
            decimals = 2) %>%
sub_missing(1:11,
            missing_text = "--") %>%
fmt(
  c(8:9, 11),
  fns = function(x)
    ifelse(x < 0.001, "<.001",
           scales::number(x, accuracy = .01))
) %>%
fmt(
  10,
  fns = function (x)
    ifelse(x > 100, ">100",
           scales::number(x, accuracy = .01))
) %>%
tab_style(
  style = list(
    cell_text(weight = "bold")
  ),
  locations = list(cells_body(
    columns = BIC,
    row = BIC == min(BIC[1:6]) # Change this to the number of classes you estimated
  ),
  cells_body(
    columns = aBIC,
    row = aBIC == min(aBIC[1:6])
  ),
  cells_body(
    columns = CAIC,
    row = CAIC == min(CAIC[1:6])
  ),
  cells_body(
    columns = AWE,
    row = AWE == min(AWE[1:6])
  ),
  cells_body(
    columns = cmPk,
    row = cmPk == max(cmPk[1:6])
  ),
  cells_body(
    columns = BF,
    row = BF > 10),
  cells_body(
    columns = T11_VLMR_PValue,

```

```

    row = ifelse(T11_VLMR_PValue < .001 & lead(T11_VLMR_PValue) > .05, T11_VLMR_PValue < .001, NA)),
  cells_body(
    columns = BLRT_PValue,
    row = ifelse(BLRT_PValue < .001 & lead(BLRT_PValue) > .05, BLRT_PValue < .001, NA))
  )
)

fit_table

```

Model Fit Summary Table¹

Classes	Par	<i>LL</i>	BIC	aBIC	CAIC	AWE	BLRT	VLMR	BF	<i>cmPk</i>
1-Class	6	-411.90	851.62	832.66	857.61	897.42	–	–	0.00	<.001
2-Class	13	-294.07	648.40	607.34	661.40	747.65	<.001	<.001	>100	1.00
3-Class	20	-288.38	669.46	606.28	689.46	822.15	0.18	0.09	>100	<.001
4-Class	27	-286.09	697.31	612.03	724.31	903.45	1.00	0.08	–	<.001

¹ Note. Par = Parameters; *LL* = model log likelihood; BIC = Bayesian information criterion; aBIC = sample size adjusted BIC; CAIC = consistent Akaike information criterion; AWE = approximate weight of evidence criterion; BLRT = bootstrapped likelihood ratio test p-value; VLMR = Vuong-Lo-Mendell-Rubin adjusted likelihood ratio test p-value; *cmPk* = approximate correct model probability.

save the table:

```
gtsave(fit_table, here("figures", "fit_table.png"))
```

Information Criterion Plot

```

allFit %>%
  dplyr::select(2:7) %>%
  rowid_to_column() %>%
  pivot_longer(`BIC`:`AWE`,
               names_to = "Index",
               values_to = "ic_value") %>%
  mutate(Index = factor(Index,
                        levels = c ("AWE", "CAIC", "BIC", "aBIC"))) %>%
  ggplot(aes(
    x = rowid,
    y = ic_value,
    color = Index,
    shape = Index,
    group = Index,
    lty = Index
  )) +
  geom_point(size = 2.0) + geom_line(size = .8) +
  scale_x_continuous(breaks = 1:nrow(allFit)) +
  scale_colour_grey(end = .5) +
  theme_cowplot() +
  labs(x = "Number of Classes", y = "Information Criteria Value", title = "Information Criteria") +
  theme(

```

```

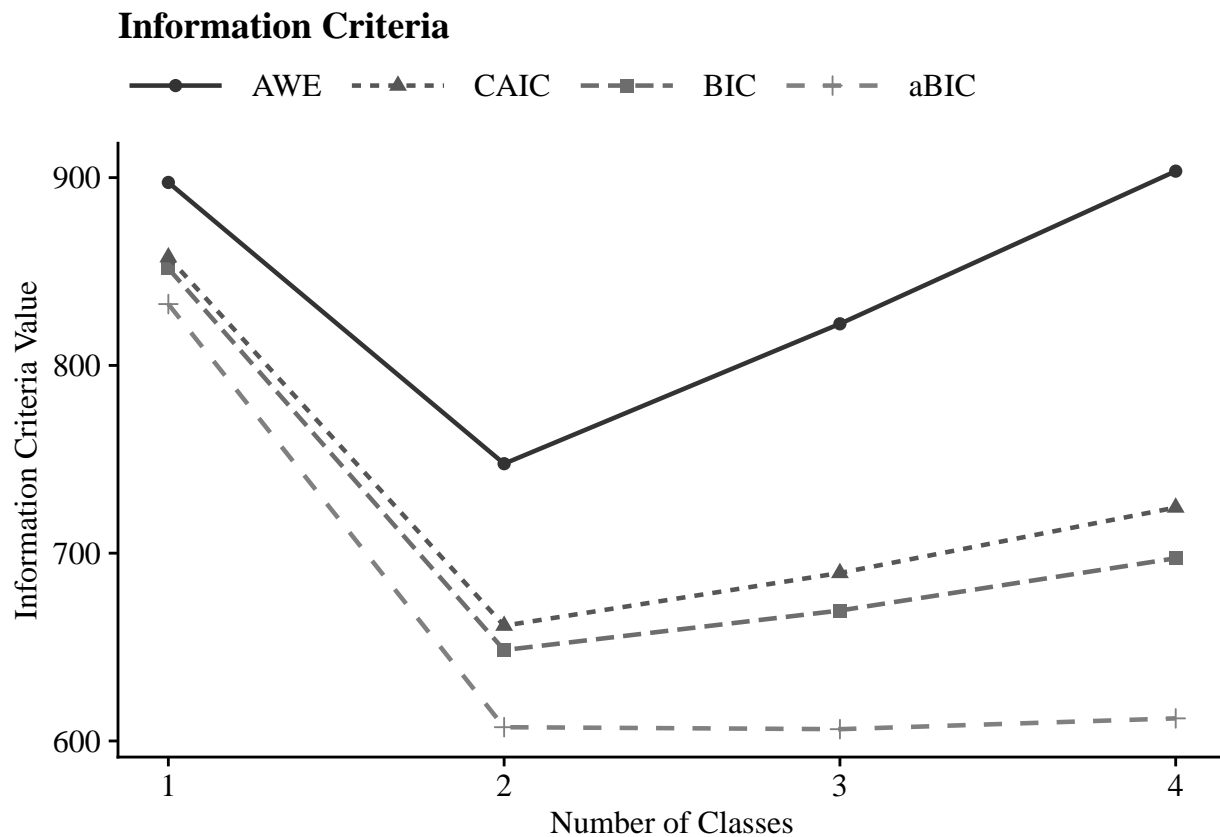
text = element_text(family = "serif", size = 12),
legend.text = element_text(family="serif", size=12),
legend.key.width = unit(3, "line"),
legend.title = element_blank(),
legend.position = "top"
)

```

```

## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use 'linewidth' instead.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.

```



save the figure:

```

ggsave(here("figures", "info_criteria.png"), dpi=300, height=5, width=7, units="in")

```

Compare class solutions

Compare probability plots for $K = 1 : 4$ class solutions

```

model_results <- data.frame()

for (i in 1:length(output_reims1)) {
  temp <- output_reims1[[i]]$parameters$probability.scale %>%

```



```

    mutate(model = paste0(i, "-Class Model"))

  model_results <- rbind(model_results, temp)
}

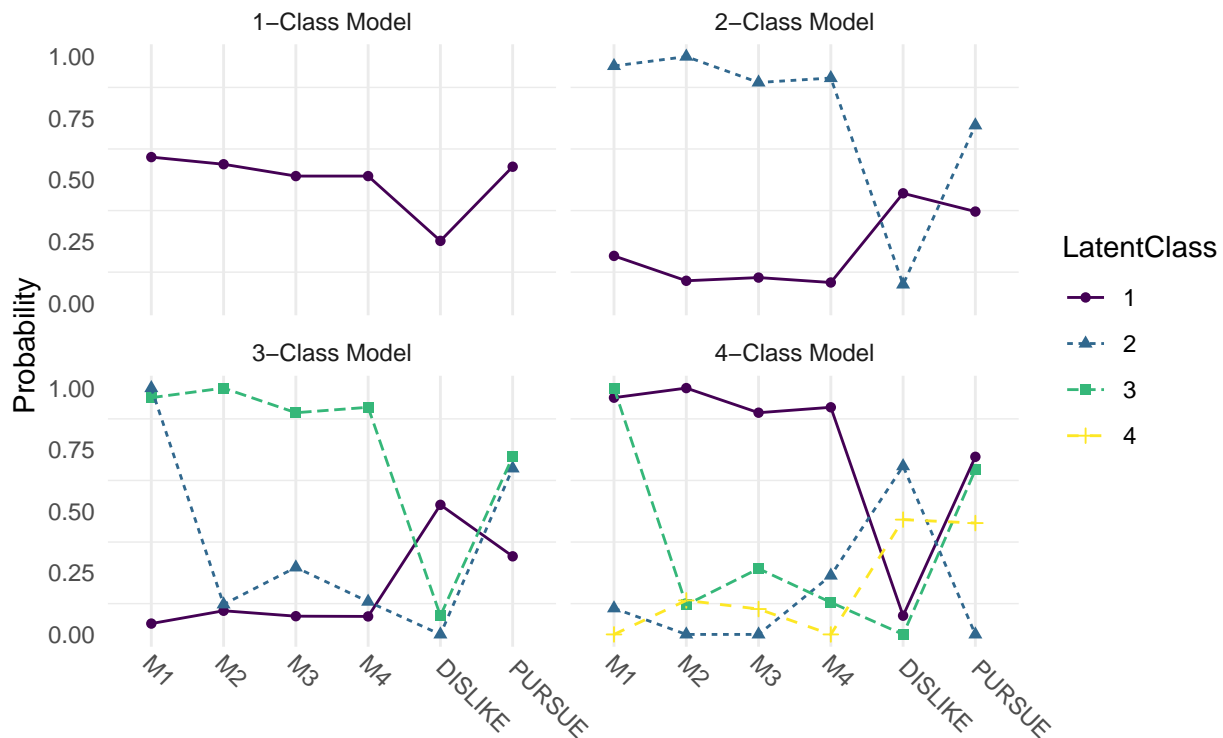
compare_plot <-
  model_results %>%
  filter(category ==2) %>%
  dplyr::select(est, model, LatentClass, param)

compare_plot$param <- fct_inorder(compare_plot$param)

ggplot(
  compare_plot,
  aes(
    x=param,
    y=est,
    color = LatentClass,
    shape = LatentClass,
    group = LatentClass,
    lty = LatentClass
  )
) +
  geom_point() +
  geom_line() +
  scale_color_viridis_d() +
  facet_wrap(~ model, ncol = 2) +
  labs(title = "Mathematics Identity Items", x = " ", y = "Probability") +
  theme_minimal() +
  theme(panel.grid.major.y = element_blank(),
        axis.text.x = element_text(angle = -45, hjust = -.1))

```

Mathematics Identity Items



save the figure:

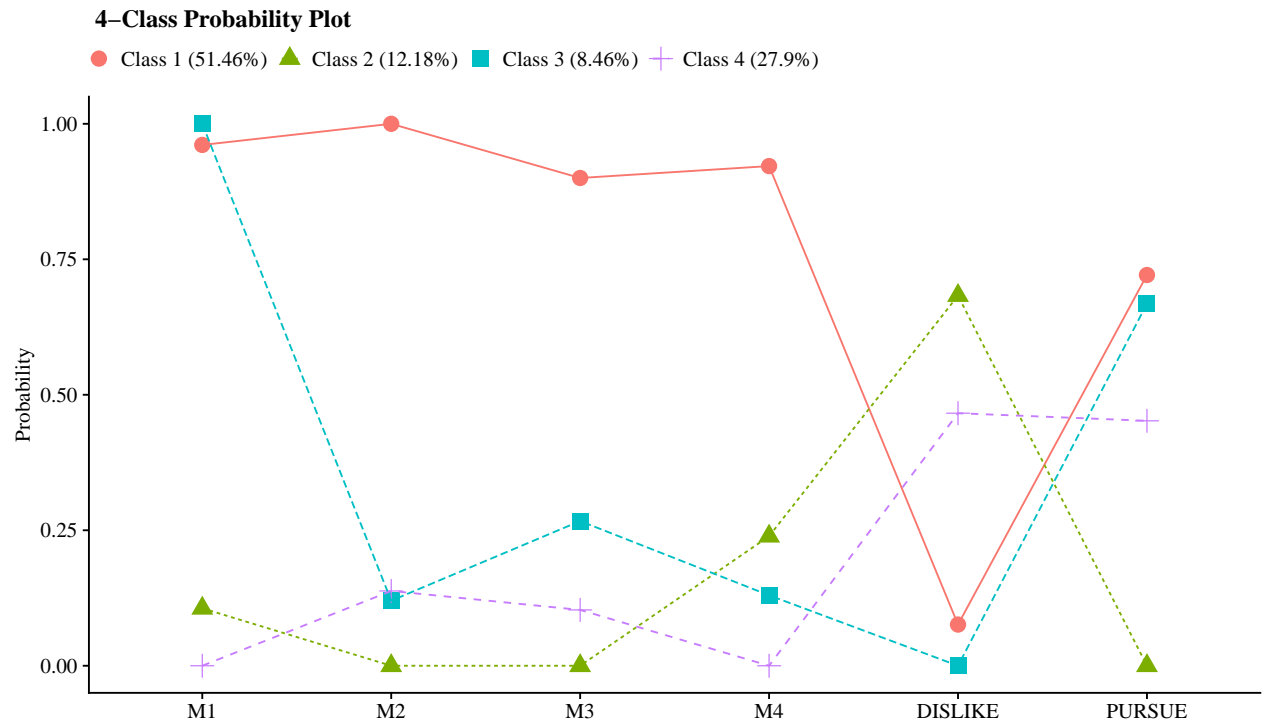
```
ggsave(here("figures", "compare_kclass_plot.png"), dpi=300, height=5, width=7, units="in")
```

4-Class Probability Plot

Use the `plot_lca` function provided in the folder to plot the item probability plot. This function requires one argument: - `model_name`: The name of the Mplus `readModels` object (e.g., `output_1sal$c4_1sal.out`) - this was updated for reims.

```
source("plot_lca.txt")

plot_lca(model_name = output_reims1$c4_reims1.out)
```



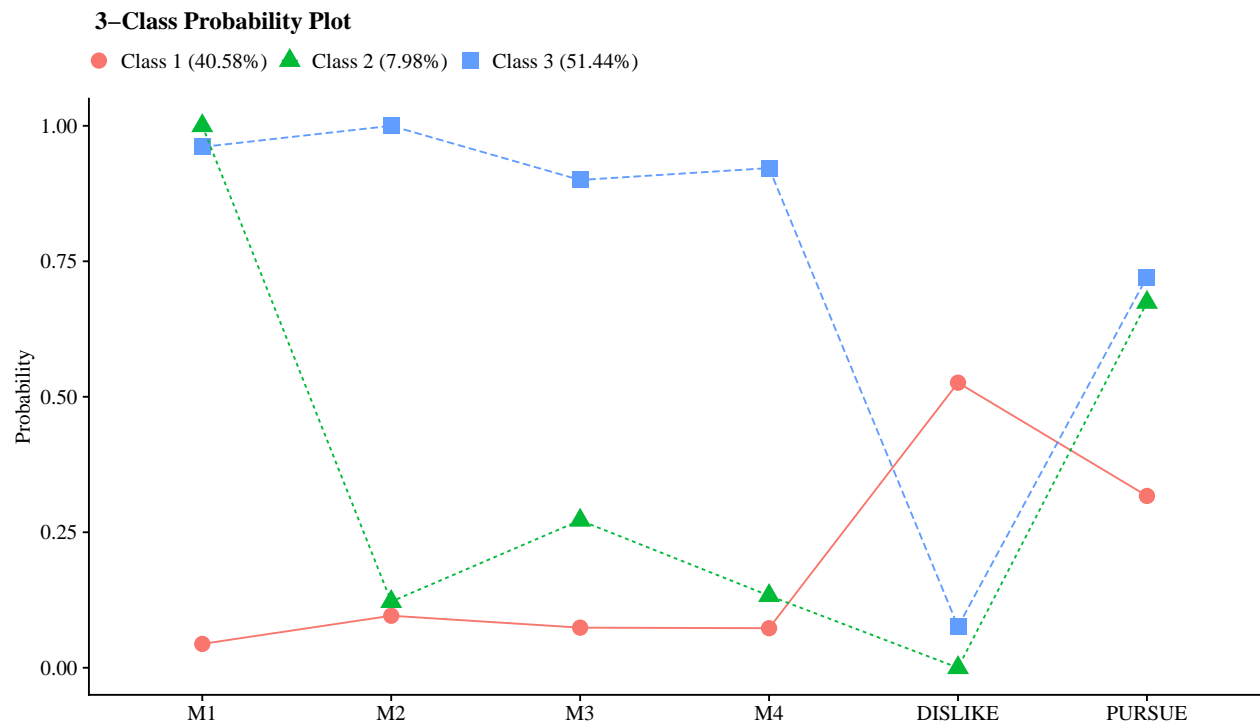
save the figure:

```
ggsave(here("figures", "probability_plot_4class.png"), dpi="retina", height=5, width=7, units="in")
```

3-Class Probability Plot

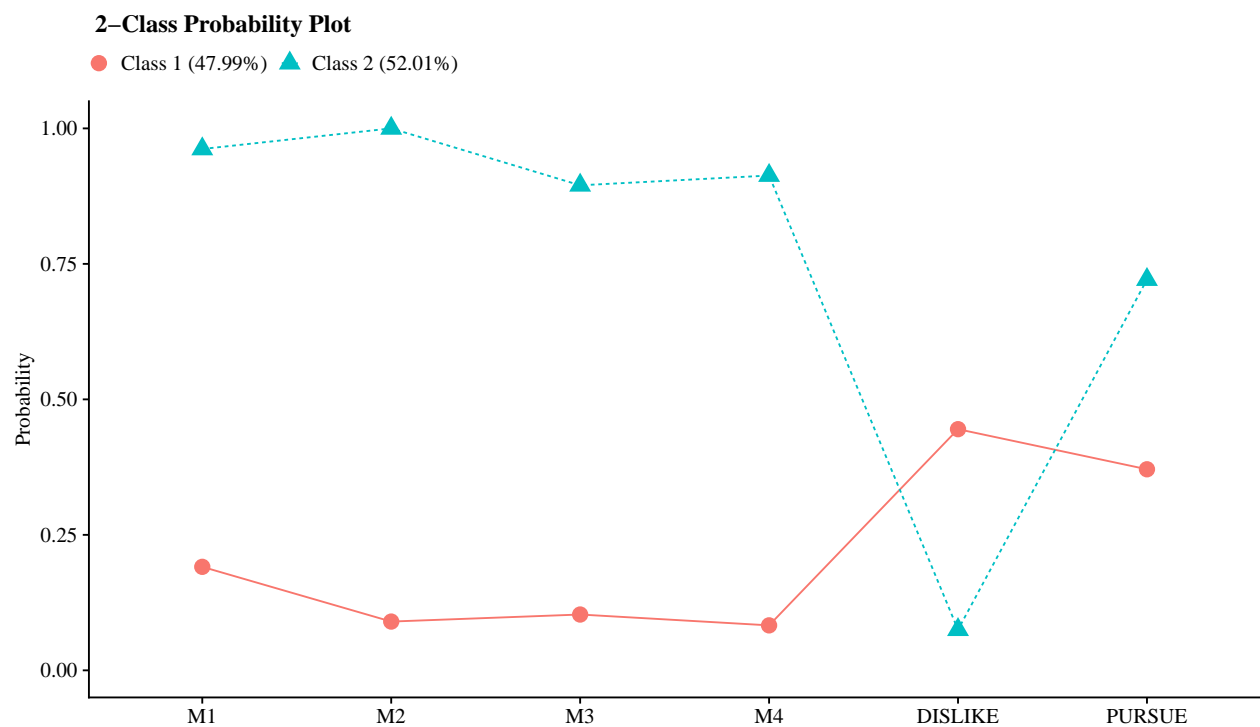
```
source("plot_lca.txt")

plot_lca(model_name = output_reims1$c3_reims1.out)
```



2-Class Probability Plot

```
source("plot_lca.txt")
plot_lca(model_name = output_reims1$c2_reims1.out)
```



save the figure:

```
ggsave(here("figures", "probability_plot_2class.png"), dpi="retina", height=5, width=7, units="in")
```

Observed response patterns

Save response frequencies for the 2-class model with `response` is `_____`.dat under `SAVEDATA`.

```
patterns <- mplusObject(

  TITLE = "LCA - Save response patterns",

  VARIABLE =
  "categorical = m1-pursue;
  usevar = m1-pursue;
  classes = c(4);",

  ANALYSIS =
  "estimator = mlr;
  type = mixture;
  starts = 0;
  processors = 10;
  optseed = 830529;",

  SAVEDATA =
  "File=savedata.dat;
  Save=cprob;

  ! Code to save response frequency data

  response is resp_patterns.dat;",

  OUTPUT = "patterns tech10 tech11 tech14",

  usevariables = colnames(reims1),
  rdata = reims1)

patterns_fit <- mplusModeler(patterns,
  dataout=here("mplus", "patterns.dat"),
  modelout=here("mplus", "patterns.inp") ,
  check=TRUE, run = TRUE, hashfilename = FALSE)
```

read in observed response pattern data and relabel the columns:

```
# read in response frequency data that we just created:
patterns <- read_table(here("mplus", "resp_patterns.dat"),
  col_names=FALSE, na = "*")
```

```
##
## -- Column specification -----
## cols(
##   X1 = col_double(),
```

```
## X2 = col_double(),
## X3 = col_double(),
## X4 = col_double(),
## X5 = col_double(),
## X6 = col_double(),
## X7 = col_double(),
## X8 = col_double(),
## X9 = col_double(),
## X10 = col_double(),
## X11 = col_double(),
## X12 = col_double()
## )
```

```
# extract the column names
names <- names(readModels(here("mplus", "patterns.out"))[['savedata']])
```

```
## Warning in (function (..., deparse.level = 1) : number of columns of result is
## not a multiple of vector length (arg 1)
```

```
## Warning in extractSampstat(outfiletext, curfile): NAs introduced by coercion
```

```
## Warning in extractSampstat(outfiletext, curfile): NAs introduced by coercion
```

```
## Warning in extractSampstat(outfiletext, curfile): NAs introduced by coercion
```

```
## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.
```

```
# add the names back to the dataset
colnames(patterns) <- c("Frequency", names)
```

we then create a table with the response patterns, then top of conditional response pattern for each modal class assignment

```
# Order responses by highest frequency
order_highest <- patterns %>%
  arrange(desc(Frequency))

# Loop `patterns` data to list top 5 conditional response patterns for each class
loop_cond <- lapply(1:max(patterns$C), function(k) {
  order_cond <- patterns %>%
    filter(C == k) %>%
    arrange(desc(Frequency)) %>%
    head(5)
})

# Convert loop into data frame
table_data <- as.data.frame(bind_rows(loop_cond))

# Combine unconditional and conditional responses patterns
response_patterns <- rbind(order_highest[1:5,], table_data)
```

we then use {gt} to make a nicely formatted table.

```

resp_table <- response_patterns %>%
  gt() %>%
    tab_header(
      title = "Observed Response Patterns",
      subtitle = html("Response patterns, estimated frequencies, estimated posterior class probabilities"),
      tab_source_note(
        source_note = md("Data Source: **Racial Ethnic Identity in Mathematics Survey (REIMS)**") %>%
        cols_label(
          Frequency = html("<i>f</i><sub>r</sub>"),
          M1 = "Math Person 1",
          M2 = "Math Person 2",
          M3 = "Math Person 3",
          M4 = "Math Person 4",
          DISLIKE = "Dislike Math",
          PURSUE = "Pursue Math",
          CPROB1 = html("P<sub><i>k</i></sub>=1"),
          CPROB2 = html("P<sub><i>k</i></sub>=2"),
          CPROB3 = html("P<sub><i>k</i></sub>=3"),
          CPROB4 = html("P<sub><i>k</i></sub>=4"), # Change based on number of classes
          C = md("**k**") %>%
        tab_row_group(
          label = "Unconditional response patterns",
          rows = 1:5) %>%
        tab_row_group(
          label = md("**k* = 1 Conditional response patterns"),
          rows = 6) %>% #EDIT THESE VALUES BASED ON THE LAST COLUMN
        tab_row_group(
          label = md("**k* = 2 Conditional response patterns"),
          rows = 7:11) %>% #EDIT THESE VALUES BASED ON THE LAST COLUMN
        tab_row_group(
          label = md("**k* = 3 Conditional response patterns"),
          rows = 12:16) %>% #EDIT THESE VALUES BASED ON THE LAST COLUMN
        tab_row_group(
          label = md("**k* = 4 Conditional response patterns"),
          rows = 17:21) %>% #EDIT THESE VALUES BASED ON THE LAST COLUMN
        row_group_order(
          groups = c("Unconditional response patterns",
                    md("**k* = 1 Conditional response patterns"),
                    md("**k* = 2 Conditional response patterns"),
                    md("**k* = 3 Conditional response patterns"),
                    md("**k* = 4 Conditional response patterns"))) %>%
        tab_footnote(
          footnote = html(
            "<i>Note.</i> <i>f</i><sub>r</sub> = response pattern frequency; P<sub><i>k</i></sub> = posterior
          )
        ) %>%
    cols_align(align = "center") %>%
    opt_align_table_header(align = "left") %>%
    gt::tab_options(table.font.names = "Times New Roman")

resp_table

```

Observed Response Patterns

Response patterns, estimated frequencies, estimated

f_r	Math Person 1	Math Person 2	Math Person 3	Math Person 4	Dislike Math	Pursue Math
Unconditional response patterns						
30	1	1	1	1	0	1
11	0	0	0	0	1	0
9	1	1	1	1	0	0
9	0	0	0	0	0	0
6	0	0	0	0	1	1
$k = 1$ Conditional response patterns						
1	0	1	1	0	1	0
$k = 2$ Conditional response patterns						
3	1	0	0	0	0	1
2	1	1	0	1	0	1
2	1	1	0	1	0	0
2	1	1	1	0	0	0
1	1	1	0	1	1	1
$k = 3$ Conditional response patterns						
30	1	1	1	1	0	1
9	1	1	1	1	0	0
2	1	1	1	1	1	1
2	1	1	1	0	0	1
1	0	1	1	1	0	0
$k = 4$ Conditional response patterns						
11	0	0	0	0	1	0
9	0	0	0	0	0	0
6	0	0	0	0	1	1
5	0	0	0	0	0	1
2	0	0	0	1	1	0

Note: f_r = response pattern frequency; P_k = posterior class probabilities

Data Source: **Racial Ethnic Identity in Mathematics Survey (REIMS)**

save the table

```
gtsave(resp_table, here("figures", "resp_table.png"))
```

Classification Diagnostics

We will use Mplus to calculate k-class confidence intervals, using the 4-class model.

```
classification <- mplusObject(
  TITLE = "LCA - Calculate k-Class 95% CI",
  VARIABLE =
    "categorical = m1-pursue;
```



```

usevar = m1-pursue;
classes = c(4);",

ANALYSIS =
  "estimator = ml;
  type = mixture;
  starts = 0;
  processors = 10;
  optseed = 945065;
  bootstrap = 100;",

MODEL =
  "
  !CHANGE THIS SECTION TO YOUR CHOSEN k-CLASS MODEL

%OVERALL%
[C#1](c1);
[C#2](c2);
[C#3](c3);

Model Constraint:
New(p1 p2 p3 p4);

p1 = exp(c1)/(1+exp(c1)+exp(c2)+exp(c3));
p2 = exp(c2)/(1+exp(c1)+exp(c2)+exp(c3));
p3 = exp(c3)/(1+exp(c1)+exp(c2)+exp(c3));
p4 = 1/(1+exp(c1)+exp(c2)+exp(c3));",

OUTPUT = "cinterval(bcbootstrap)",

usevariables = colnames(reims1),
rdata = reims1)

classification_fit <- mplusModeler(classification,
  dataout=here("mplus", "reims-1.dat"),
  modelout=here("mplus", "class.inp") ,
  check=TRUE, run = TRUE, hashfilename = FALSE)

```

Note from IMMERSE team: Ensure that the classes did not shift during this step (i.g., Class 1 in the enumeration run is now Class 4). Evaluate output and compare the class counts and proportions for the latent classes. Using the OPTSEED function ensures replication of the best loglikelihood value run.

Read in the 4-class model:

```

# Read in the 4-class model and extract information needed
class_output <- readModels(here("mplus", "class.out"))

```

```

## Warning in (function (... , deparse.level = 1) : number of columns of result is
## not a multiple of vector length (arg 1)

```

```

## Warning in extractSampstat(outfiletext, curfile): NAs introduced by coercion
## Warning in extractSampstat(outfiletext, curfile): NAs introduced by coercion
## Warning in extractSampstat(outfiletext, curfile): NAs introduced by coercion

## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.

# Entropy
entropy <- c(class_output$summaries$Entropy, rep(NA, class_output$summaries$NLatentClasses-1))

# 95% k-Class and k-class 95% Confidence Intervals
k_ci <- class_output$parameters$ci.unstandardized %>%
  filter(paramHeader == "New.Additional.Parameters") %>%
  unite(CI, c(low2.5, up2.5), sep=" ", remove = TRUE) %>%
  mutate(CI = paste0("[", CI, "]")) %>%
  rename(kclass=est) %>%
  dplyr::select(kclass, CI)

# AvePPk = Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent C
avePPk <- tibble(avePPk = diag(class_output$class_counts$avgProbs.mostLikely))

# mcaPk = modal class assignment proportion
mcaPk <- round(class_output$class_counts$mostLikely, 3) %>%
  mutate(model = paste0("Class ", class)) %>%
  add_column(avePPk, k_ci) %>%
  rename(mcaPk = proportion) %>%
  dplyr::select(model, kclass, CI, mcaPk, avePPk)

# OCCk = odds of correct classification
OCCk <- mcaPk %>%
  mutate(OCCk = round((avePPk/(1-avePPk))/(kclass/(1-kclass)), 3))

# Put everything together
class_df <- data.frame(OCCk, entropy)

```

now we use {gt} to make a nicely formatted table

```

class_table <- class_df %>%
  gt() %>%
  tab_header(
    title = "Model Classification Diagnostics for the 4-Class Solution" %>%
  cols_label(
    model = md("*k*-Class"),
    kclass = md("*k*-Class Proportions"),
    CI = "95% CI",
    mcaPk = md("*mcaPk*"),
    avePPk = md("*AvePPk*"),
    OCCk = md("*OCCk*"),
    entropy = "Entropy" %>%
  sub_missing(7,
    missing_text = "") %>%
  cols_align(aligned = "center") %>%

```

```

opt_align_table_header(align = "left") %>%
gt::tab_options(table.font.names = "Times New Roman")

class_table

```

Model Classification Diagnostics for the 4-Class Solution

<i>k</i> -Class	<i>k</i> -Class Proportions	95% CI	<i>mcaPk</i>	<i>AvePPk</i>	<i>OCCk</i>	Entropy
Class 1	0.51	[0.395, 0.591]	0.515	0.986	67.667	0.894
Class 2	0.30	[0.175, 0.399]	0.301	0.897	20.320	
Class 3	0.01	[0, 0.029]	0.010	0.997	32901.000	
Class 4	0.18	[0.08, 0.299]	0.175	0.814	19.937	

save the table:

```

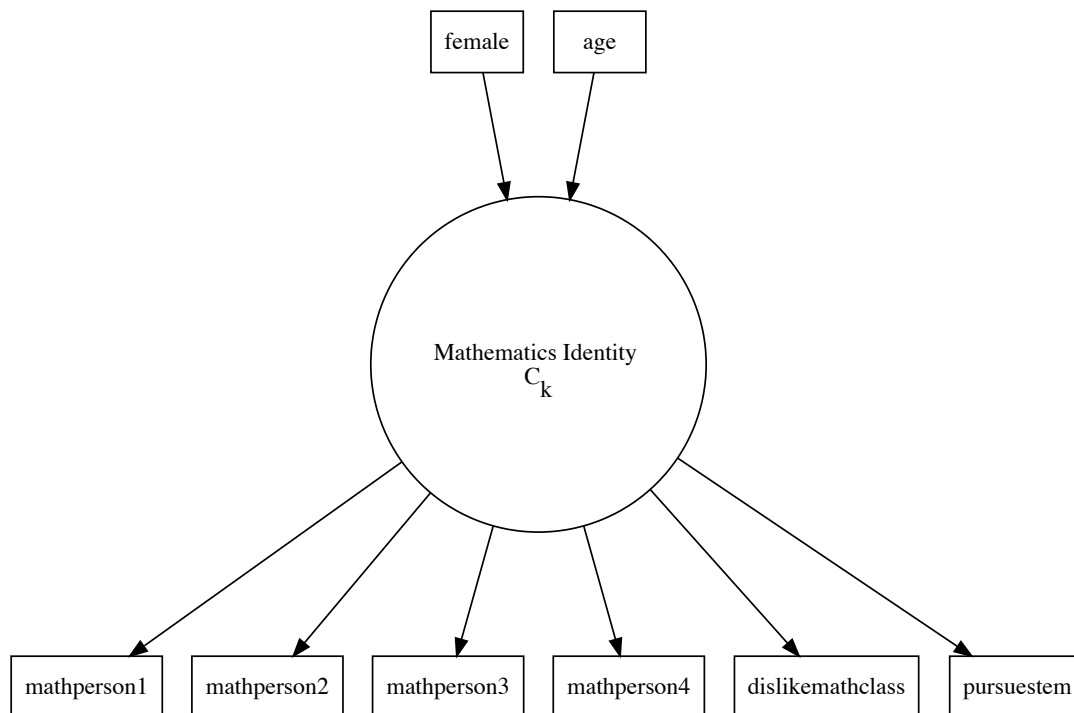
gtsave(class_table, here("figures", "class_table.png"))

```

Auxillary variables

Model 3

We will now add auxiliary variables to our model and name it model 3.



Manual 3-step latent class regression with 3 covariates

Integrate covariates and with the mixture model

```
# indicators and variables for full model build

tribble(
  ~"Name", ~"Variable Description",
  #-----/-----/,
  "mathperson1", "I am a math person.",
  "mathperson2", "My peers think I am a math person.",
  "mathperson3", "My teachers think I am a math person.",
  "mathperson4", "There are a lot of people who think I am a math person.",
  "dislikemathclass", "In general, I dislike math classes.",
  "pursuestem", "If given the chance, I will pursue a STEM career.",
  "female", "Self-reported sex (0=male, 1=female)",
  "age", "Self-reporeted age") %>%
gt() %>%
tab_header(title = md("**LCA Indicators & Auxiliary Variables: Mathematics Identity Example**"), subti
tab_row_group(group = "", rows = 1:6) %>%
tab_row_group(group = "Auxiliary Variables", rows = 7:8) %>%
row_group_order(groups = c("", "Auxiliary Variables")) %>%
tab_options(column_labels.font.weight = "bold",
             row_group.font.weight = "bold")
```

```
## Warning: Since gt v0.3.0 the 'group' argument has been deprecated.
## * Use the 'label' argument to specify the group label.
## This warning is displayed once every 8 hours.
```

LCA Indicators & Auxiliary Variables: Mathematics Identity Example

Name	Variable Description
mathperson1	I am a math person.
mathperson2	My peers think I am a math person.
mathperson3	My teachers think I am a math person.
mathperson4	There are a lot of people who think I am a math person.
dislikemathclass	In general, I dislike math classes.
pursuestem	If given the chance, I will pursue a STEM career.
Auxiliary Variables	
female	Self-reported sex (0=male, 1=female)
age	Self-reporeted age

```
# add female variable from original data set
# the variables were not included in the subset data
reims$sex
```

```
##      [1] 2 2 1 1 1 2 1 2 2 2 2 1 1 2 1 1 2 2 1 2 2 1 2 1 1 2 2 2 2 2 2 2 2 2 3 2
##     [38] 2 1 2 2 1 1 2 2 1 2 2 2 2 2 2 1 1 1 2 2 2 2 2 1 1 2 1 1 2 1 2 2 2 2 1 2
##     [75] 2 2 1 2 1 1 2 1 2 2 2 1 2 2 2 2 2 1 2 1 2 1 1 2 2 1 2 1 1
```

```
reims1$female <- reims$sex
reims1$age <- reims$age
```

Step 1: Class enumeration with auxiliary specification

```
step1 <- mplusObject(
  TITLE = "STEP 1 - THREE-STEP USING LSAL",
  VARIABLE =
    "categorical = m1-pursue;
    usevar = m1-pursue;
    classes = c(3);

    auxiliary = ! list all potential covariates and distals here

    female ! covariate",

  ANALYSIS =
    "estimator = mlr;
    type = mixture;
    starts = 500 100;",

  SAVEDATA =
    "File=3step_savedata.dat;
    Save=cprob;",

  OUTPUT = "residual tech11 tech14",

  PLOT =
    "type = plot3;
    series = m1-pursue(*);",

  usevariables = colnames(reims1),
  rdata = reims1)

step1_fit <- mplusModeler(step1,
  dataout=here("manual", "Step1.dat"),
  modelout=here("manual", "one.inp"),
  check=T, run=T, hashfilename = F)
```

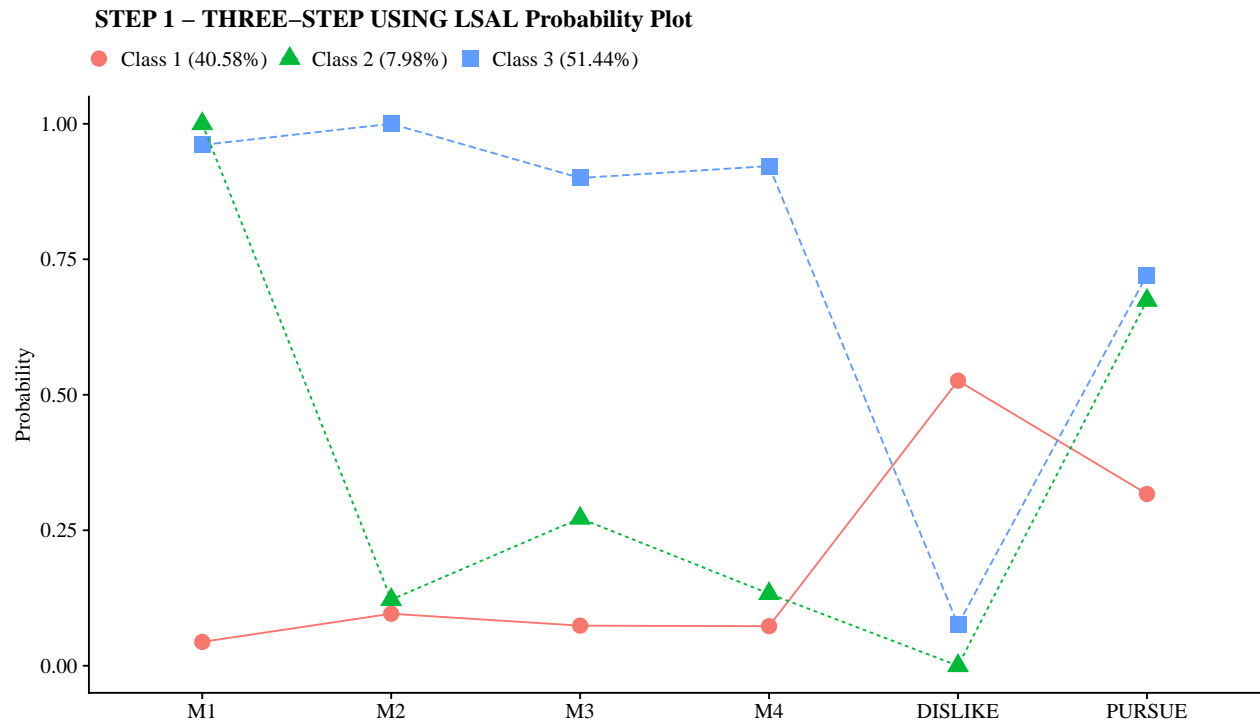
save the plot

```
source("plot_lca.txt")

output_reims1 <- readModels(here("manual", "one.out"))
```

```
## No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.
```

```
plot_lca(model_name = output_reims1)
```



Step 2: Determine measurement error

we want to extract the logits for the classification probabilities for the most likely latent class.

```
logic_cprobs <- as.data.frame(output_reims1[["class_counts"]]  
                             [["logitProbs.mostLikely"]])
```

we now extract the saved data set.

```
savedata <- as.data.frame(output_reims1[["savedata"]])
```

we then rename the column in savedata named "c" and change to "N"

```
colnames(savedata)[colnames(savedata)=="C"] <- "N"
```

Step 3: Add auxiliary variables to analysis

Model 1 with female as covariate (to see if there are differences by sex). There are no distal outcomes included in this sample model.

I will model the auxiliary variable with 3 classes first.

```
step3 <- mplusObject(  
  TITLE = "STEP 3 - 3STEP LSAY",  
  VARIABLE =
```

```

"nominal=N;
usevar = n;
classes = c(3);

usevar = female;",

ANALYSIS =
"estimator = mlr;
type = mixture;
starts = 0;",

MODEL =
  glue(
    " %OVERALL%

    C on female (f1-f2); ! covariate as predictor of C

    %C#1%
    [n#1@{logit_cprobs[1,1]}};
    [n#2@{logit_cprobs[1,2]}};

    %C#2%
    [n#1@{logit_cprobs[2,1]}};
    [n#2@{logit_cprobs[2,2]}};

    %C#3%
    [n#1@{logit_cprobs[3,1]}};
    [n#2@{logit_cprobs[3,2]}}];",

MODELTEST = "
m1=m2;
m2=m3;
!f1=f2;
!f2=f3;
",

MODELCONSTRAINT =
  "New (diff12 diff13 diff23);

  diff12 = m1-m2; ! test pairwise distal mean differences
  diff13 = m1-m3;
  diff23 = m2-m3;",

usevariables = colnames(savedata),
rdata = savedata)

step3_fit <- mplusModeler(step3,
  dataout=here("manual", "Step3.dat"),
  modelout=here("manual", "three.inp"),
  check=TRUE, run = TRUE, hashfilename = FALSE)

```

This file is based on resources provided by the Institute of Mixture Modeling for Equity-Oriented Researchers, Scholars, and Educators (2024). IMMERSE In-Person Training Workshop (IES No. 305B220021). Institute of Education Sciences. <https://immerse-ucsb.github.io/pre-training>