

# Mixture Models with Covariates & Distal Outcomes

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## Preparation

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### Data source:

1. The first example utilizes a dataset on undergraduate *Cheating* available from the `poLCA` package (Dayton, 1998): [See documentation here](#)
  2. The second examples utilizes the public-use dataset, *The Longitudinal Survey of American Youth* (**LSAY**): [See documentation here](#)
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Load packages

```
library(naniar)
library(tidyverse)
library(haven)
library(glue)
library(MplusAutomation)
library(here)
library(janitor)
library(gt)
library(poLCA)
```

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Incorporating distal outcome variables with mixture models

**Note:** Prior to adding covariates or distals enumeration must be conducted.

[See Lab 7 for examples of enumeration with MplusAutomation](#)

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## DU3step auxiliary variable integration

- Using the DU3step you can specify distal relations but cannot specify models with covariates & distals
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## Application: Undergraduate Cheating behavior

“Dichotomous self-report responses by 319 undergraduates to four questions about cheating behavior” (poLCA, 2016).

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Prepare data

```
data(cheating)

cheating <- cheating %>% clean_names()

df_cheat <- cheating %>%
  dplyr::select(1:4) %>%
  mutate_all(funs(.-1)) %>%
  mutate(gpa = cheating$gpa)
```

Run the **DU3step** model with gpa as distal outcome

```
m_stepdu <- mplusObject(
  TITLE = "DU3STEP add distal GPA",
  VARIABLE =
    "categorical = lieexam-copyexam;
    usevar = lieexam-copyexam;
    auxiliary = gpa (du3step);
    classes = c(2);",

  ANALYSIS =
    "estimator = mlr;
    type = mixture;
    starts = 500 100;
    processors = 10;",

  OUTPUT = "sampstat patterns tech11 tech14;",

  PLOT =
    "type = plot3;
    series = lieexam-copyexam(*);",

  usevariables = colnames(df_cheat),
  rdata = df_cheat)

m_stepdu_fit <- mplusModeler(m_stepdu,
  dataout=here("20-three-step", "du3step_mplus", "lca_du3step.dat"),
  modelout=here("20-three-step", "du3step_mplus", "c2_lca_du3step.inp") ,
  check=TRUE, run = TRUE, hashfilename = FALSE)
```

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## Application: Longitudinal Study of American Youth, Science Attitudes

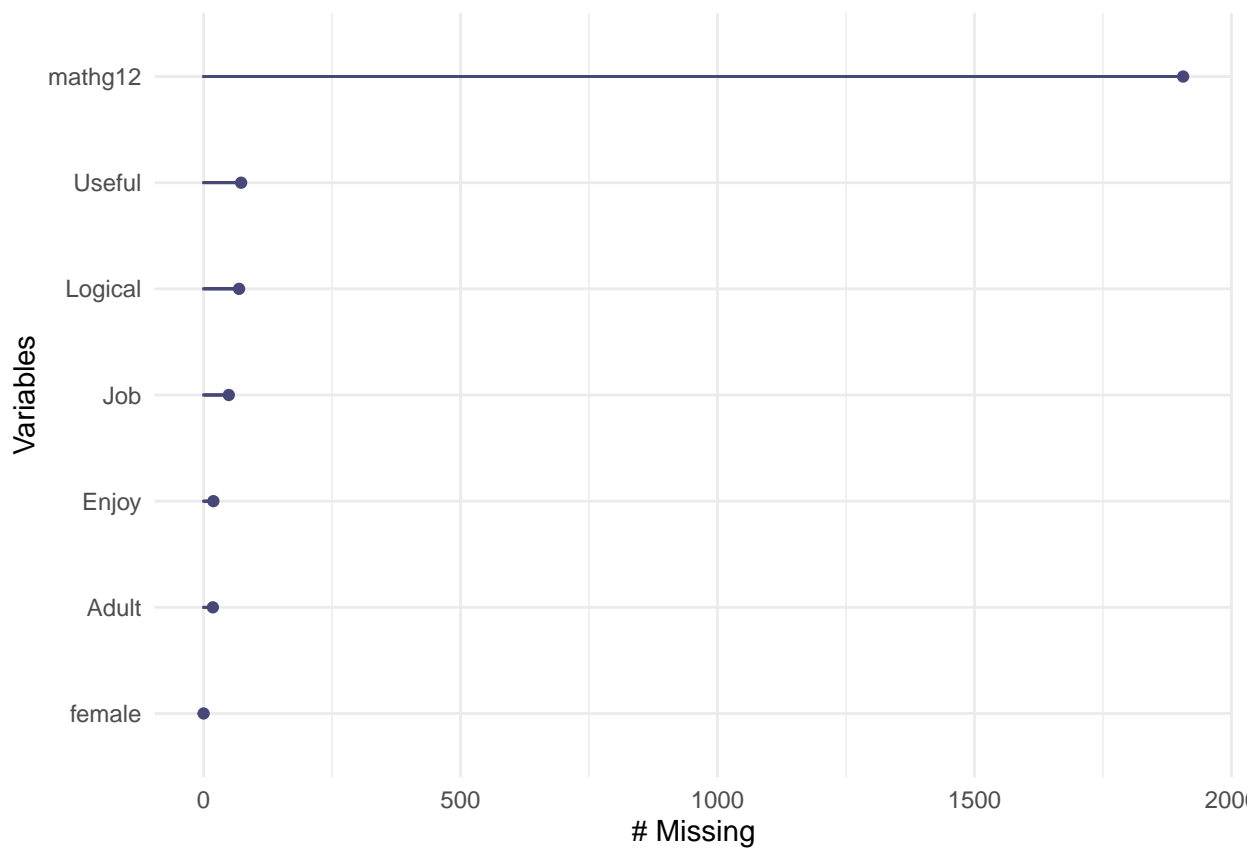
---

Load data

```
lsay_data <- read_csv("https://garberadamc.github.io/project-site/data/lca_lsay_sci.csv",  
  na = c("9999", "9999.00")) %>%  
  clean_names() %>%  
  dplyr::select(1:5, female, mathg12,  
    Enjoy = ab39m, Useful = ab39t,  
    Logical = ab39u, Job = ab39w, Adult = ab39x)
```

Use `{naniar}` to look at missing on covariates and distals

```
naniar::gg_miss_var(lsay_data)
```



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### Manual 3-step

- Adding covariates and distals to a mixture model

- Often called “*auxiliary variable integration*”

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## Step 1

---

```
step1 <- mplusObject(
  TITLE = "Step1 - 3step LSAY",
  VARIABLE =
    "categorical = Enjoy-Adult;
    usevar = Enjoy-Adult;

    classes = c(4);

    auxiliary = ! list all potential covariates and distals here
    female      ! covariate
    mathg12;    ! distal math test score in 12th grade ",

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

  SAVEDATA =
    "File=3step_savedata.dat;
    Save=cprob;
    Missflag= 999;",

  OUTPUT = "sampstat residual tech11 tech14",

  PLOT =
    "type = plot3;
    series = Enjoy-Adult(*);",

  usevariables = colnames(lsay_data),
  rdata = lsay_data)

step1_fit <- mplusModeler(step1,
  dataout=here("20-three-step", "3step_mplus", "Step1_3step_LSAY.dat"),
  modelout=here("20-three-step", "3step_mplus", "Step1_3step_LSAY.inp") ,
  check=TRUE, run = TRUE, hashfilename = FALSE)
```

---

## Step 2

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Extract logits for the classification probabilities for the most likely latent class

```
logit_cprobs <- as.data.frame(step1_fit[["results"]]
                                [["class_counts"]]
                                [["logitProbs.mostLikely"]])
```

Extract saved dataset which is part of the mplusObject “step1\_10\_fit”

```
savedata <- as.data.frame(step1_fit[["results"]]
                            [["savedata"]])
```

Rename the column in savedata named “C” and change to “N”

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

Run step 2

```
step2 <- mplusObject(
  TITLE = "Step2 - 3step LSAY",

  VARIABLE =
    "nominal=N;
    USEVAR = n;
    missing are all (999);
    classes = c(4); ",

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

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

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

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

      %C#4%
      [n#1@{logit_cprobs[4,1]}};
      [n#2@{logit_cprobs[4,2]}};
      [n#3@{logit_cprobs[4,3]}};"),

  usevariables = colnames(savedata),
```

```

rdata = savedata)

step2_fit <- mplusModeler(step2,
                          dataout=here("20-three-step", "3step_mplus", "Step2_3step_LSAY.dat"),
                          modelout=here("20-three-step", "3step_mplus", "Step2_3step_LSAY.inp"),
                          check=TRUE, run = TRUE, hashfilename = FALSE)

```

---

## Step 3

---

Model with 1 covariate and 1 distal outcome

```

step3 <- mplusObject(
  TITLE = "Step3 - 3step LSAY",

  VARIABLE =
  "nominal=N;
  usevar = n;
  missing are all (999);
  classes = c(4);

  usevar = female mathg12;" ,

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

  MODEL =
  glue(
  " %OVERALL%

  C on female;      ! covariate as predictor of C

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

  [mathg12](m1);      ! conditional distal mean
  mathg12;            ! conditional distal variance (freely estimated)

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

  [mathg12](m2);
  mathg12;

```

```

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

[mathg12](m3);
mathg12;

%C#4%
[n#1@{logit_cprobs[4,1]}};
[n#2@{logit_cprobs[4,2]}};
[n#3@{logit_cprobs[4,3]}};

[mathg12](m4);
mathg12; "),

MODELCONSTRAINT =
  "New (diff12 diff13 diff23
    diff14 diff24 diff34);

  diff12 = m1-m2; ! test pairwise distal mean differences
  diff13 = m1-m3;
  diff23 = m2-m3;
  diff14 = m1-m4;
  diff24 = m2-m4;
  diff34 = m3-m4; ",

MODELTEST = "      ! omnibus test of distal means
  m1=m2;
  m2=m3;
  m3=m4; ",

usevariables = colnames(savedata),
rdata = savedata)

step3_fit <- mplusModeler(step3,
  dataout=here("20-three-step", "3step_mplus", "Step3_3step_L SAY.dat"),
  modelout=here("20-three-step", "3step_mplus", "Step3_3step_L SAY.inp"),
  check=TRUE, run = TRUE, hashfilename = FALSE)

```

End of manual 3-step

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Model with latent categorical variable ( $C_k$ ) as moderator

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```

step3mod <- mplusObject(
  TITLE = "Step3 - 3step LSAY",

```

```

VARIABLE =
"nominal=N;
usevar = n;
missing are all (999);
classes = c(4);

usevar = female mathg12;" ,

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

MODEL =
glue(
"!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
!!!!DISTAL = mathg12, COVARIATE = female, MODERATOR = C!!!!
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
%OVERALL%
mathg12 on female;
mathg12;

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

mathg12 on female(s1); ! conditional slope (class 1)
[mathg12] (m1); ! conditional distal mean
mathg12; ! conditional distal variance (freely estimated)

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

mathg12 on female(s2);
[mathg12] (m2);
mathg12;

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

mathg12 on female(s3);
[mathg12] (m3);
mathg12;

%C#4%
[n#1@{logit_cprobs[4,1]}};
[n#2@{logit_cprobs[4,2]}};
[n#3@{logit_cprobs[4,3]}};

```



```

mathg12 on female(s4);
[mathg12](m4);
mathg12; "),

MODELCONSTRAINT =
  "New (slope12 slope13 slope23
    slope14 slope24 slope34);

    slope12 = s1-s2; ! test pairwise slope differences
    slope13 = s1-s3;
    slope23 = s2-s3;
    slope14 = s1-s4;
    slope24 = s2-s4;
    slope34 = s3-s4;",

MODELTEST = " ! can run only a single Omnibus test per model
  s1=s2;
  s2=s3;
  s3=s4;",

usevariables = colnames(savedata),
rdata = savedata)

step3mod_fit <- mplusModeler(step3mod,
  dataout=here("20-three-step", "3step_mplus", "Step3_moderation_L SAY.dat"),
  modelout=here("20-three-step", "3step_mplus", "Step3_moderation_L SAY.inp"),
  check=TRUE, run = TRUE, hashfilename = FALSE)

```

## References

- Drew A. Linzer, Jeffrey B. Lewis (2011). poLCA: An R Package for Polytomous Variable Latent Class Analysis. *Journal of Statistical Software*, 42(10), 1-29. URL <http://www.jstatsoft.org/v42/i10/>.
- Hallquist, M. N., & Wiley, J. F. (2018). MplusAutomation: An R Package for Facilitating Large-Scale Latent Variable Analyses in Mplus. *Structural equation modeling: a multidisciplinary journal*, 25(4), 621-638.
- Miller, J. D., Hoffer, T., Suchner, R., Brown, K., & Nelson, C. (1992). LSAY codebook. Northern Illinois University.
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- Wickham et al., (2019). Welcome to the tidyverse. *Journal of Open Source Software*, 4(43), 1686, <https://doi.org/10.21105/joss.01686>