



Examining Exams Using Rasch Models and Assessment of Measurement Invariance

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Overview

- Topics
 - Large-scale exams
 - Item response theory with Rasch model
 - Assessment of measurement invariance
- Mathematics 101 exam at Universität Innsbruck
 - Classical tests
 - Anchor methods
 - Score-based tests
 - Model-based recursive partitioning
 - Finite mixture models
- Discussion

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Large-scale exams

Motivation:

- Statisticians often teach large lecture courses for other fields.
- Statistics, probability, or mathematics in curricula such as business and economics, social sciences, psychology, etc.
- At WU Wien and Universität Innsbruck: Some courses are attended by more than 1,000 students per semester.
- Several lecturers teach lectures and tutorials in parallel.

Typical exams:

- Multiple choice or single choice.
- Evaluated and graded automatically.
- Little further examination of results (if any).

Large-scale exams

Potential questions:

- Ability of students.
- Difficulty of exercises (or items).
- Differential item functioning (DIF).
- Unidimensionality.

At WU: Multiple-choice monitor by Ledermüller, Nettekoven, Weiler/Krakovsky.

Here:

- Rasch model for binary single-choice items.
- Assessment of measurement invariance vs. DIF.

IRT with Rasch model

Motivation: Item response theory (IRT) with Rasch model.

- Measure a single latent trait (here: ability in exam).
- Based on binary items y_{ij} (here: solved correctly vs. not).
- Align person's ability θ_i ($i = 1, \dots, n$) and item's difficulty β_j ($j = 1, \dots, m$) on the same scale.

Model:

$$\begin{aligned}\pi_{ij} &= \Pr(\text{person } i \text{ solves item } j) = \Pr(y_{ij} = 1) \\ \text{logit}(\pi_{ij}) &= \theta_i - \beta_j\end{aligned}$$

- Interval scale with arbitrary zero point.
- Fix reference point by zero constraint (e.g., for β_1 or $\sum_j \beta_j$).
- Consistent estimation via conditional maximum likelihood.
- Sufficient statistics for θ_i : Sum of correct items for person i .

Assessment of measurement invariance

Crucial assumption: Measurement invariance (MI). Otherwise observed differences cannot be reliably attributed to the latent variable that the model purports to measure.

Parameter stability: In parametric models, the MI assumption corresponds to stability of parameters across all possible subgroups.

Inference: The typical approach for assessing MI is

- to split the data into reference and focal groups,
- assess the stability of selected parameters (all or only a subset) across these groups
- by means of standard tests: likelihood ratio (LR), Wald, or Lagrange multiplier (LM or score) tests.

Assessment of measurement invariance

Problems:

- Subgroups have to be formed in advance.
- Continuous variables are often categorized into groups in an ad hoc way (e.g., splitting at the median).
- In ordinal variables the category ordering is often not exploited – assessing only if at least one group differs from the others.
- When likelihood ratio or Wald tests are employed, the model has to be fitted to each subgroup which can become numerically challenging and computationally intensive.

Conceivable solutions:

- Score-based tests “along” numerical/ordinal/categorical covariates.
- Recursive partitioning to capture covariate interactions.
- Finite mixture models without covariates.

Mathematics 101 at Universität Innsbruck

Course: Mathematics for first-year business and economics students at Universität Innsbruck.

Format: Biweekly online tests (conducted in OpenOLAT) and two written exams for about 1,000 students per semester.

Here: Individual results from an end-term exam.

- 729 students (out of 941 registered).
- 13 single-choice items with five answer alternatives, covering the basics of analysis, linear algebra, financial mathematics.
- Two groups with partially different item pools (on the same topics). Individual versions of items generated via *exams* in R.
- Correctly solved items yield 100% of associated points. Items without correct solution can either be unanswered (0%) or with an incorrect answer (−25%). Considered as binary here.

Mathematics 101 at Universität Innsbruck

Variables: In MathExam14W.

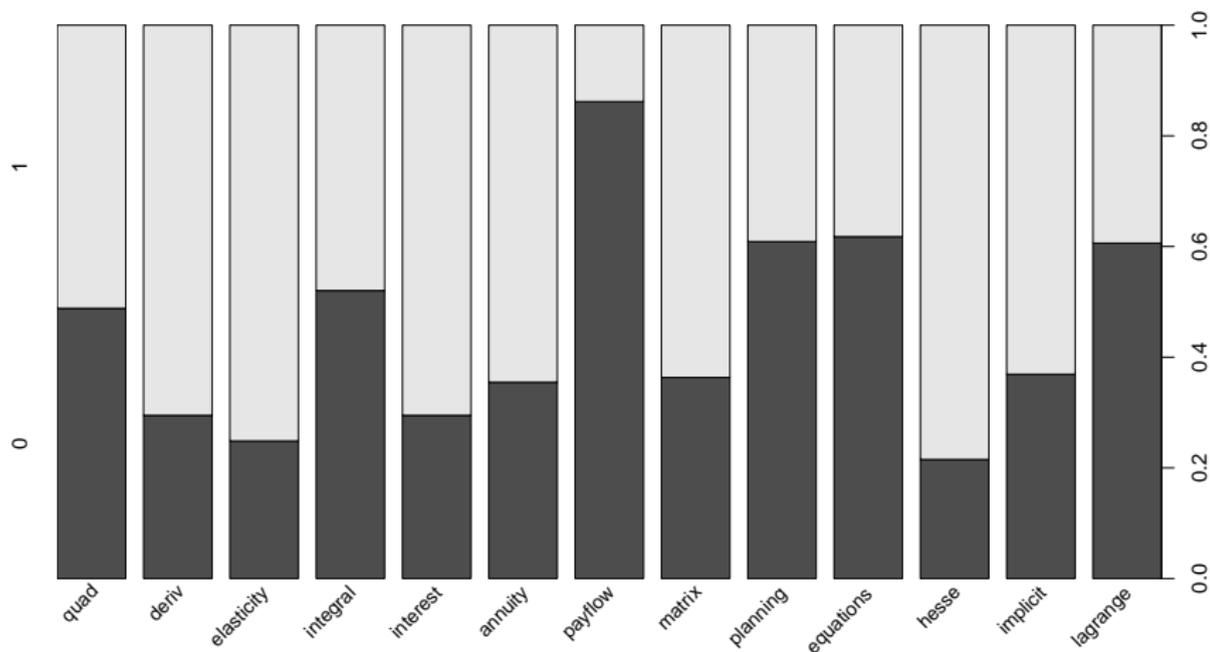
- solved: Item response matrix (1/0 coding).
- group: Factor for group.
- tests: Number of previous online exercises solved (out of 26).
- nsolved: Number of exam items solved (out of 13).
- gender, study, attempt, semester, ...

In R: Load package/data and exclude extreme scorers.

```
R> library("psychotools")  
R> data("MathExam14W", package = "psychotools")  
R> mex <- subset(MathExam14W, nsolved > 0 & nsolved < 13)
```

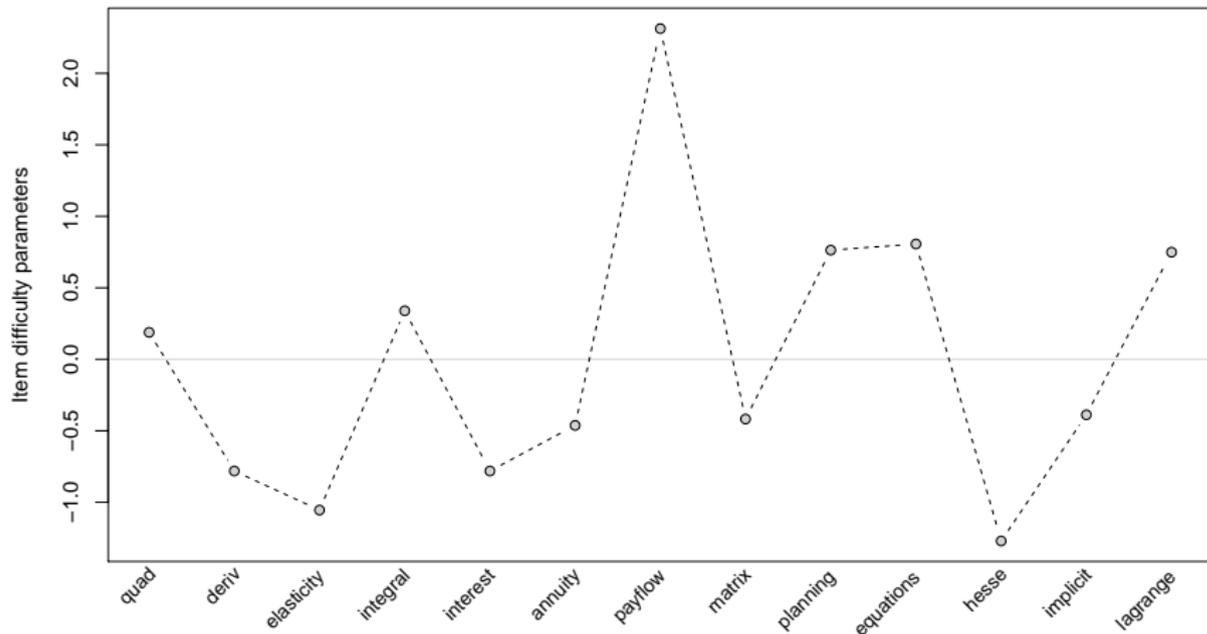
Mathematics 101 at Universität Innsbruck

```
R> plot(mex$solved)
```



Rasch model

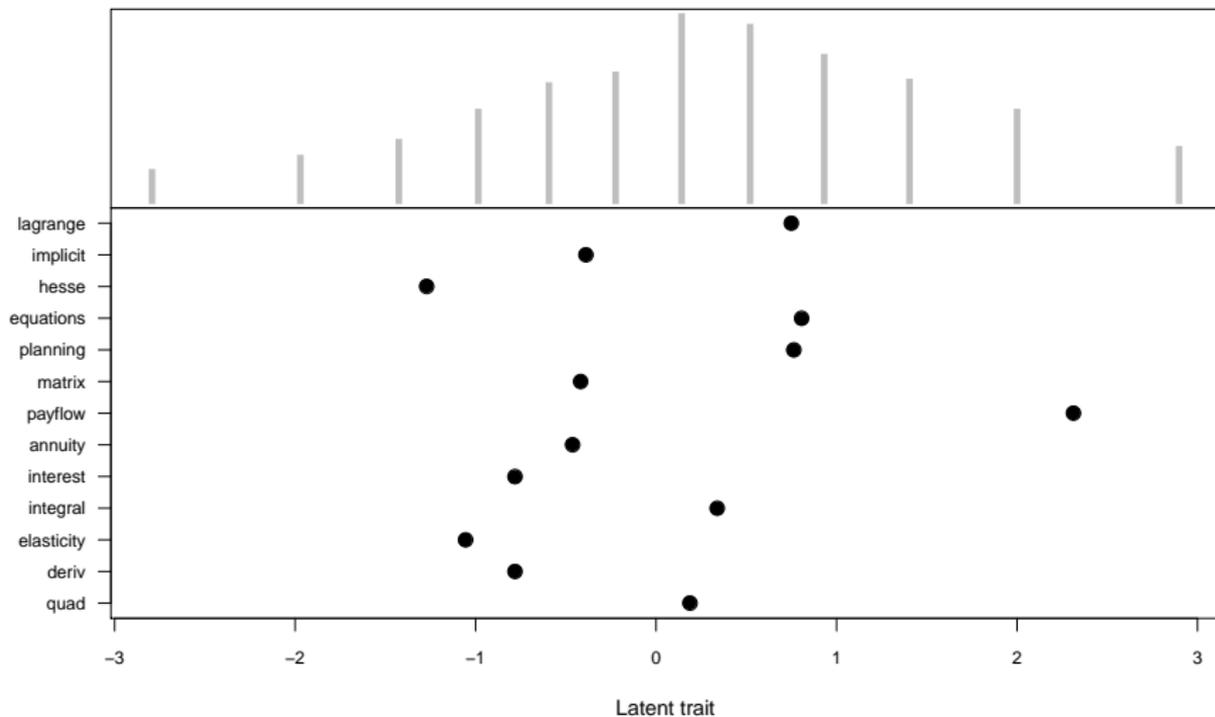
```
R> mr <- raschmodel(mex$solved)
R> plot(mr, type = "profile")
```



Rasch model

```
R> plot(mr, type = "piplot")
```

Person-Item Plot



Classical tests

Of interest: Difference between the two exam groups.

Tests: All χ^2_{12} with 95% critical value 21.0.

- LR: 265.0.
- Wald: 249.4.
- LM/Score: 260.8.

Question: Which items “cause” this DIF?

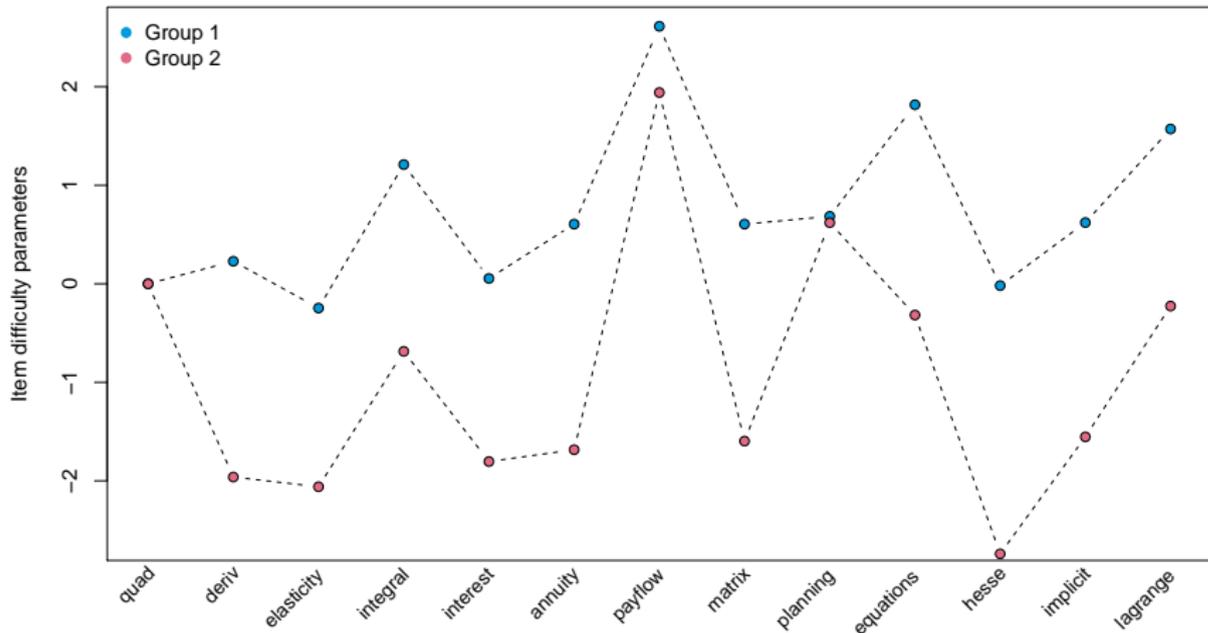
Answer: Use item-wise Wald tests.

$$t_j = \frac{\hat{\beta}_j^{\text{ref}} - \hat{\beta}_j^{\text{toc}}}{\sqrt{\widehat{\text{Var}}(\hat{\beta}^{\text{ref}})_{j,j} + \widehat{\text{Var}}(\hat{\beta}^{\text{toc}})_{j,j}}}.$$

But: “Anchor” items are needed to align the scales from the two groups.

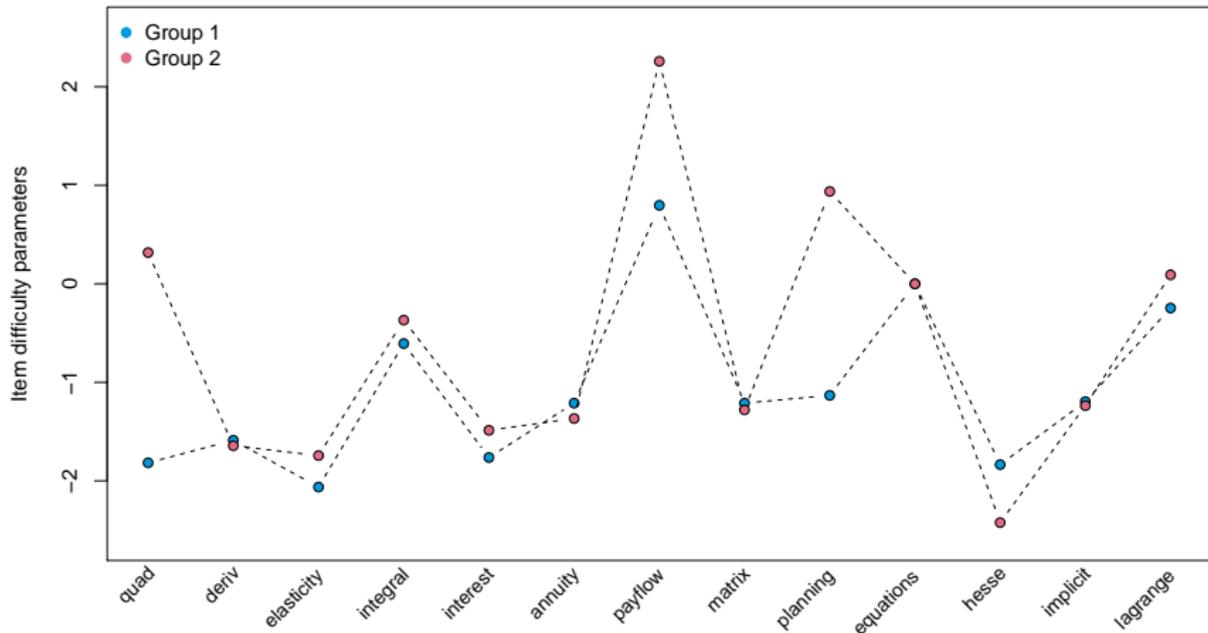
Classical tests

```
R> plot(mr1, parg = list(ref = 1), ...)  
R> plot(mr2, parg = list(ref = 1), ...)
```



Classical tests

```
R> plot(mr1, parg = list(ref = 10), ...)  
R> plot(mr2, parg = list(ref = 10), ...)
```



Anchor methods

Goal: Select DIF-free anchor items to be able to identify items truly associated with DIF (“chicken or the egg” dilemma).

Approaches: Classes of anchors with different characteristics.

- *All other:* All items – except the item currently studied.
- *Constant:* Predefined number of items (e.g., 1 or 4).
- *Forward:* Iteratively add items.

Selection: Rank candidate items based on single-anchor DIF tests.

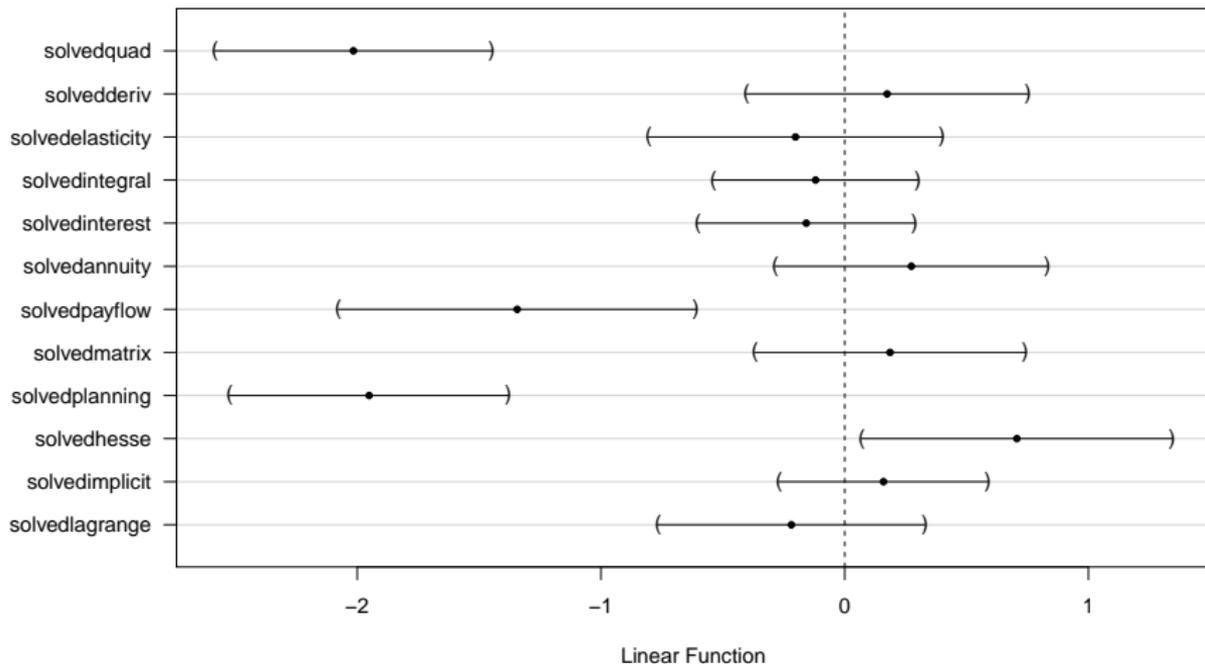
- Number of significant tests.
- Mean test statistic or p -value.
- Mean test statistic or p -value beyond median threshold.

Here: Constant anchor class with 4 items and mean p -value threshold selection. Single-step adjustment of final inference for multiple testing.

Anchor methods

```
R> ma <- anchortest(solved ~ group, data = mex, adjust = "single-step")  
R> plot(ma$final_tests)
```

Anchor items: 10, 4, 12, 5



Score-based tests

Questions:

- Is there further DIF in the two exam groups?
- Is there DIF w.r.t. mathematics ability, e.g., for tests $(0, \dots, 13, \dots, 26)$ or `nsolved` $(1, \dots, 12)$?

Problem: Numeric variables without predefined subgroups. Hence, many possible patterns of deviation from parameter stability.

Idea: Generalize the LM test.

- Model only has to be fitted once under the MI assumption to the full data set.
- Capture model deviations along a variable v that is suspected to cause DIF and violate MI.

Score-based tests

Hypotheses: Under ML parameters β do not depend any variable v_i .
Hence assess for $i = 1, \dots, n$

$$H_0 : \beta_i = \beta,$$

$$H_1 : \beta_i = \beta(v_i).$$

Building block: Casewise model deviations.

- Derivative of the casewise log-likelihood w.r.t. the parameters.
- General measure of model deviation (similar to residuals).

$$\mathbf{s}(\beta; \mathbf{y}_i) = \left(\frac{\partial \ell(\beta; \mathbf{y}_i)}{\partial \beta_2}, \dots, \frac{\partial \ell(\beta; \mathbf{y}_i)}{\partial \beta_m} \right)^\top$$

Score-based tests

Special case: Two subgroups resulting from one split point ν .

$$H_1^* : \beta_i = \begin{cases} \beta^{(A)} & \text{if } v_i \leq \nu \\ \beta^{(B)} & \text{if } v_i > \nu \end{cases}$$

Tests: LR/Wald/LM tests can be easily employed if pattern $\beta(v_i)$ is known, specifically for H_1^* with fixed split point ν .

For unknown split point: Compute LR/Wald/LM tests for each possible split point $v_1 \leq v_2 \leq \dots \leq v_n$ and reject if the maximum statistic is large.

Caution: By maximally selecting the test statistic different critical values are required (not from a χ^2 distribution)!

More generally: Consider a class of tests that assesses whether the model “deviations” $\mathbf{s}(\hat{\beta}; \mathbf{y}_i)$ depend on v_i .

Score-based tests

Fluctuation process: Capture fluctuations in the cumulative sum of the scores ordered by the variable v .

$$\mathbf{B}(t; \hat{\beta}) = \hat{\mathbf{I}}^{-1/2} n^{-1/2} \sum_{i=1}^{\lfloor n \cdot t \rfloor} \mathbf{s}(\hat{\beta}; \mathbf{y}_{(i)}) \quad (0 \leq t \leq 1).$$

- $\hat{\mathbf{I}}$ – estimate of the information matrix.
- t – proportion of data ordered by v .
- $\lfloor n \cdot t \rfloor$ – integer part of $n \cdot t$.
- $x_{(i)}$ – observation with the i -th smallest value of the variable v .

Functional central limit theorem: Under H_0 convergence to a (continuous) Brownian bridge process $\mathbf{B}(\cdot; \hat{\beta}) \xrightarrow{d} \mathbf{B}^0(\cdot)$, from which critical values can be obtained – either analytically or by simulation.

Score-based tests: Continuous variables

Test statistics: The empirical process can be viewed as a matrix $\mathbf{B}(\hat{\beta})_{ij}$ with rows $i = 1, \dots, n$ (observations) and columns $j = 1, \dots, m - 1$ (parameters). This can be aggregated to scalar test statistics along continuous the variable v .

$$\begin{aligned} DM &= \max_{i=1, \dots, n} \max_{j=1, \dots, m-1} |\mathbf{B}(\hat{\beta})_{ij}| \\ CvM &= n^{-1} \sum_{i=1, \dots, n} \sum_{j=1, \dots, m-1} \mathbf{B}(\hat{\beta})_{ij}^2, \\ \max LM &= \max_{i=\underline{i}, \dots, \bar{i}} \left\{ \frac{i}{n} \left(1 - \frac{i}{n} \right) \right\}^{-1} \sum_{j=1, \dots, m-1} \mathbf{B}(\hat{\beta})_{ij}^2. \end{aligned}$$

Critical values: Analytically for DM . Otherwise by direct simulation or further refined simulation techniques.

Score-based tests: Ordinal variables

Test statistics: Aggregation along ordinal variables v with c categories.

$$WDM_o = \max_{i \in \{i_1, \dots, i_{c-1}\}} \left\{ \frac{i}{n} \left(1 - \frac{i}{n} \right) \right\}^{-1/2} \max_{j=1, \dots, m-1} |\mathbf{B}(\hat{\beta})_{ij}|,$$
$$\max LM_o = \max_{i \in \{i_1, \dots, i_{c-1}\}} \left\{ \frac{i}{n} \left(1 - \frac{i}{n} \right) \right\}^{-1} \sum_{j=1, \dots, m-1} \mathbf{B}(\hat{\beta})_{ij}^2,$$

where i_1, \dots, i_{c-1} are the numbers of observations in each category.

Critical values: For WDM_o directly from a multivariate normal distribution. For $\max LM_o$ via simulation.

Score-based tests: Categorical variables

Test statistic: Aggregation within the c (unordered) categories of v .

$$LM_{uo} = \sum_{\ell=1, \dots, c} \sum_{j=1, \dots, m-1} \left(\mathbf{B}(\hat{\beta})_{i_{\ell}j} - \mathbf{B}(\hat{\beta})_{i_{\ell-1}j} \right)^2,$$

Critical values: From a χ^2 distribution (as usual).

Asymptotically equivalent: LR test.

Score-based tests

Here: Test for DIF along tests in group 1 with max *LM* test (continuous vs. ordinal).

Result: Clear evidence for DIF. Students that performed poorly in the previous online tests have a different item profile.

```
R> library("strucchange")
R> mex1 <- subset(mex, group == 1)

R> sctest(mr1, order.by = mex1$tests, vcov = "info",
+         functional = "maxLM")
```

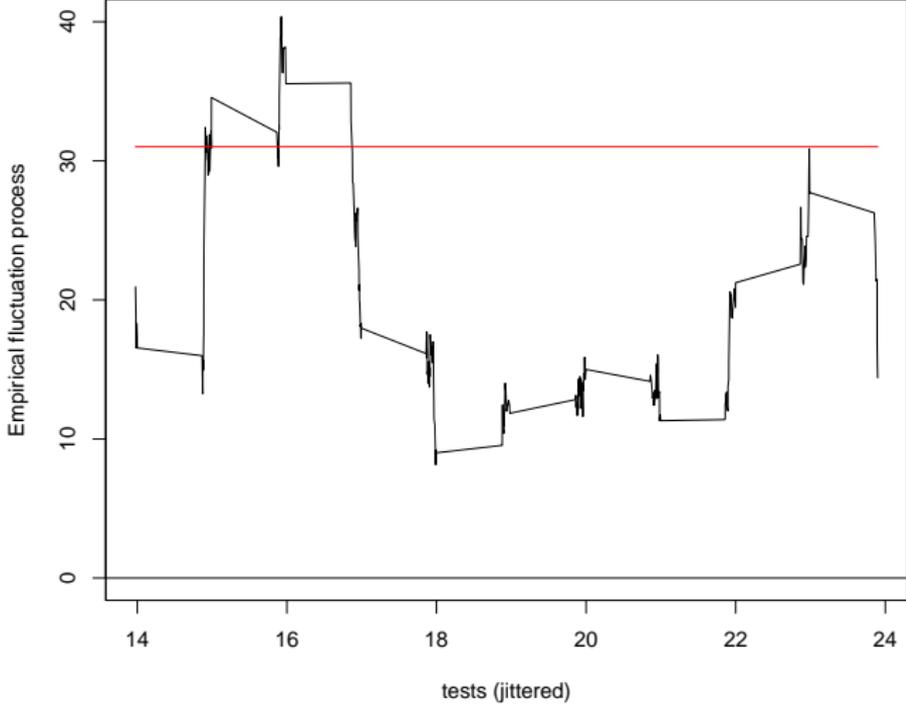
```
           M-fluctuation test
data:  mr1
f(efp) = 40.365, p-value = 0.002508
```

```
R> sctest(mr1, order.by = mex1$tests, vcov = "info",
+         functional = "maxLMo")
```

```
           M-fluctuation test
data:  mr1
f(efp) = 35.543, p-value = 0.003961
```

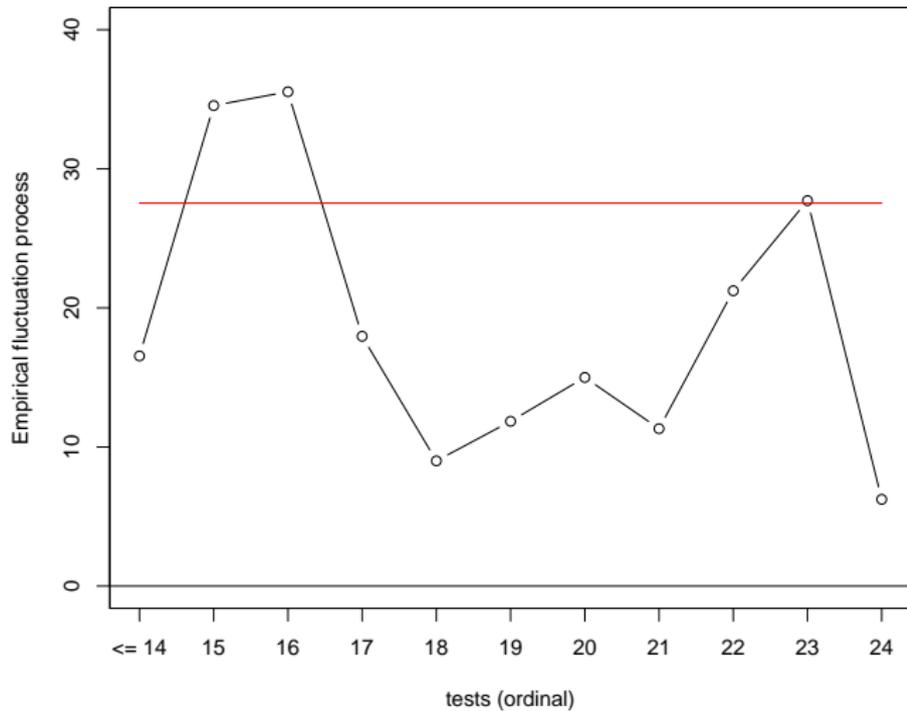
Score-based tests

M-fluctuation test



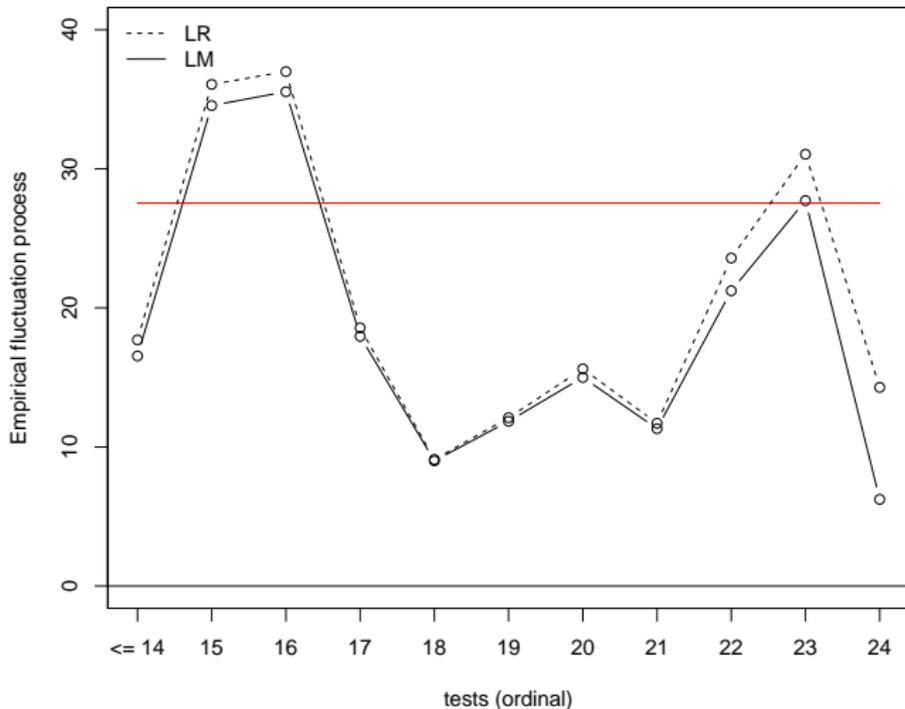
Score-based tests

M-fluctuation test



Score-based tests

M-fluctuation test



Recursive partitioning

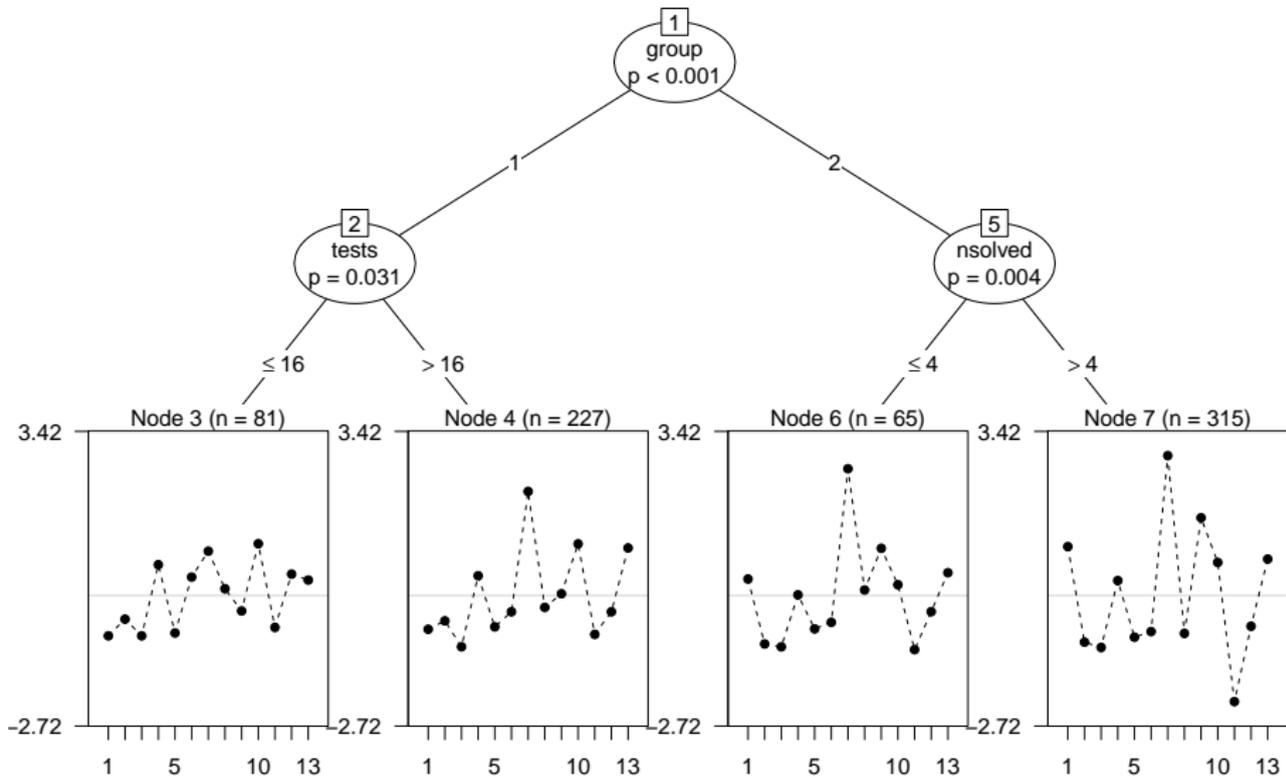
Idea: Apply tests recursively.

- Assess all covariates of interest using Bonferroni adjustment.
- Split w.r.t. covariate with smallest significant p -value.
- Select split point by maximizing the log-likelihood.
- Continue until there are no more significant instabilities (or the sample is too small).

Here: Treat numeric variables with few levels as ordinal. Simulate p -values for max LM_o test.

```
R> library("psychotree")
R> mex$tests <- ordered(mex$tests)
R> mex$nsolved <- ordered(mex$nsolved)
R> mex$attempt <- ordered(mex$attempt)
R> mex$semester <- ordered(mex$semester)
R> mrt <- raschtree(solved ~ group + tests + nsolved + gender +
+   attempt + study + semester, data = mex,
+   vcov = "info", minsize = 50, ordinal = "L2", nrep = 1e5)
```

Recursive partitioning



Finite mixture models

Question: How to detect DIF without covariate information (e.g., in group 1 without tests)?

Answer: Finite mixture of Rasch models with $k = 1, \dots, K$ components. Maximize finite mixture likelihood via EM w.r.t. component-specific weights ω_k and item difficulties $\beta^{(k)}$.

$$\max_{\omega, \beta^{(1)}, \dots, \beta^{(K)}} \prod_{i=1}^n \sum_{k=1}^K \omega_k f(\mathbf{y}_i; \beta^{(k)})$$

Possible extensions:

- Model selection for the number of components K .
- Concomitant variables for the mixture weights ω .
- Component-specific distributions for the raw scores.

Finite mixture models

Here: 2-component mixture with component-specific raw score distribution (mean-variance specification).

```
R> library("psychomix")  
R> mrm <- raschmix(mex1$solved, k = 2, scores = "meanvar")  
R> plot(mrm)
```

Result: The “soft” classification found by the mixture model is rather similar to the “hard” split by the tree.

```
R> print(mrm)
```

Call:

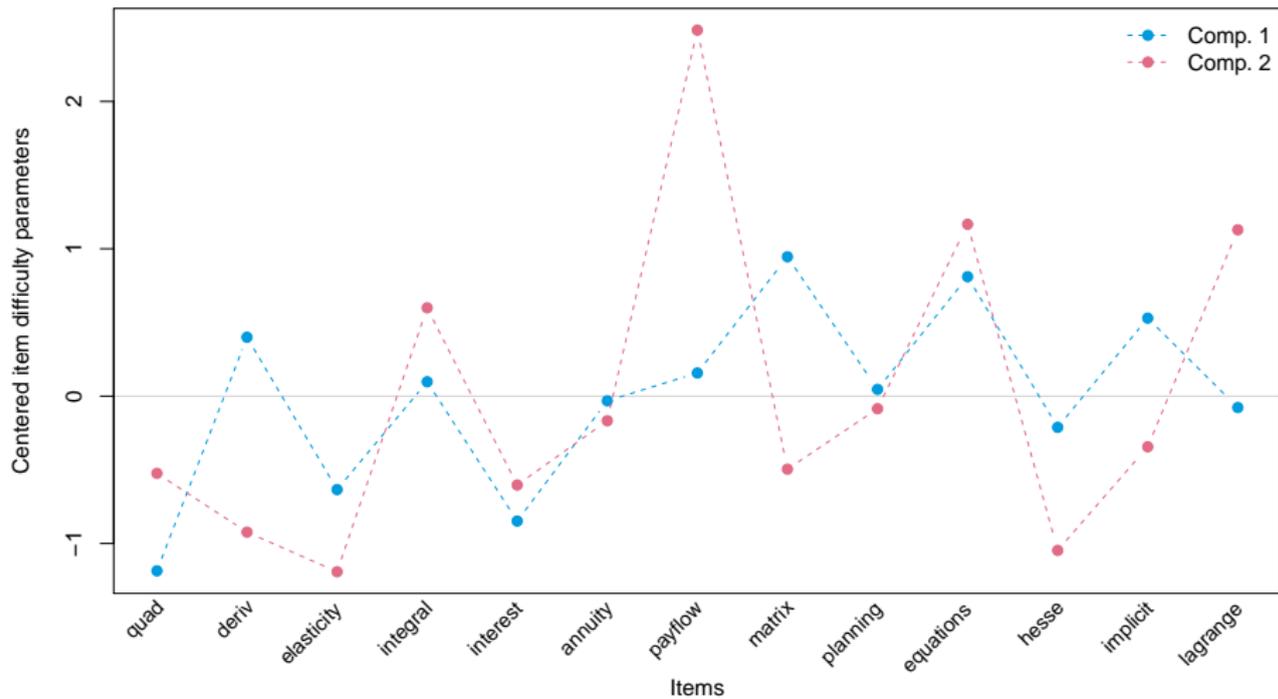
```
raschmix(formula = mex1$solved, k = 2, scores = "meanvar")
```

Cluster sizes:

```
  1  2  
73 235
```

convergence after 79 iterations

Finite mixture models



Discussion

Summary:

- Flexible toolbox for assessing measurement invariance in parametric psychometric models.
- Detecting violations along one (tests), none (mixture), or many (tree) covariates.
- Exploit different scales of the covariates: continuous, ordinal, or categorical.

Here: Probably quickest overview of DIF patterns with Rasch tree.

At UIBK: Resulting “policy” implications.

- Avoid exam groups if at all possible.
- Seemingly equivalent items can function very differently if students focus their learning on well-known parts of the item pool.

Discussion

R packages:

- *strucchange* provides an object-oriented implementation of the score-based parameter instability tests.
- Model-based recursive partitioning available in *partykit*.
- Psychometric models that cooperate with *strucchange* and *partykit* are provided in *psychotools*: IRT models (Rasch, partial credit, rating scale), Bradley-Terry, multinomial processing trees.
- Psychometric trees in *psychotree*.
- Psychometric mixture models in *psychomix* (based on *flexmix* plus *psychotools*).

Discussion

Exams infrastructure: R package *exams*.

- R for random data generation and computations.
- L^AT_EX or Markdown for text formatting
- Answer types: Single/multiple choice, numeric, string, cloze.

Output:

- PDF – either fully customizable or standardized with automatic scanning/evaluation.
- HTML – either fully customizable or embedded into any of the standard formats below.
- Moodle XML.
- QTI XML standard (version 1.2 or 2.1), e.g., for OLAT/OpenOLAT.
- ARSnova, Blackboard, TCEXam, WU-Prüfungsserver, . . .

Discussion



Discussion

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qt12 Finish test

Actual score: 0 / 5

qt12

1. Exercise Still 1 attempt(s)

1.1. Question 0 / 0

2. Exercise 0 / 0

2.1. Question 0 / 0

3. Exercise 0 / 0

3.1. Question 0 / 0

4. Exercise 0 / 0

4.1. Question 0 / 0

5. Exercise 0 / 0

5.1. Question 0 / 0

Question

In Figure the distributions of a variable given by two samples (A and B) are represented by parallel boxplots. Which of the following statements are correct? (Comment: The statements are either about correct or clearly wrong.)

Sample	Min	Q1	Median	Q3	Max
A	-28	-26	-23	-21	-20
B	-35	-29	-23	-19	-15

Figure 1: Parallel boxplots.

- a. The location of both distributions is about the same.
- b. Both distributions contain no outliers.
- c. The spread in sample A is clearly bigger than in B.
- d. The skewness of both samples is similar.
- e. Distribution A is about symmetric.

Save answer

Discussion

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Back R/exams/1

1 2 3

In the following figure the distributions of a variable given by two samples (A and B) are represented by parallel boxplots. Which of the following statements are correct? (Comment: The statements are either about correct or clearly wrong.)

The figure shows two parallel boxplots, A and B, on a vertical axis ranging from -35 to -10. Boxplot A has a median around -22, a box from -28 to -16, and whiskers from -34 to -11. Boxplot B has a median around -22, a box from -24 to -20, and whiskers from -26 to -17.

The location of both distributions is about the same.

Start Questions Feedback System Manual

References

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