

Introduction to Item Response Theory

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IRT TO THE PEOPLE



Outline

- Review Measurement and Classical Test Theory
- Establish Item Response Theory Fundamentals
- Compare Classical Test Theory and Item Response Theory
- Assumptions
- Dichotomous Models
- Example Using R
- Interpretation

Measurement

- Definitions for consideration
 - Assigning numbers to individuals in a systematic way
 - Applied to the properties of objects rather than to the objects themselves
- Conceptualize the latent variable or construct
 - Products of the informed scientific imagination of social scientists who attempt to develop theories for explaining human behavior
 - Represents what is common across manifested behaviors
 - The hidden source of common variability or covariability of a set of similar observable behaviors
 - Psychological theory is a statement about a possible relationship between two or more psychological constructs or between a construct and an observable behavior
- Establish the operational definition
 - What are the behavioral manifestations of the construct?

Classical Test Theory

- Evaluates the extent to which observed scores are affected by measurement error
- Classical Test Theory decomposition

$$X = T + E$$

- True score: average of all measurements obtained after a test is administered many times to the same individual ensuring measurements are i.i.d (independent and identically distributed).

$$T_{ki} = \bar{X}_{ki}$$

- Test and subject specific
- Error score: the complement of an individual's true score to the observed score on a given test

$$E = X - T$$

IRT Origins

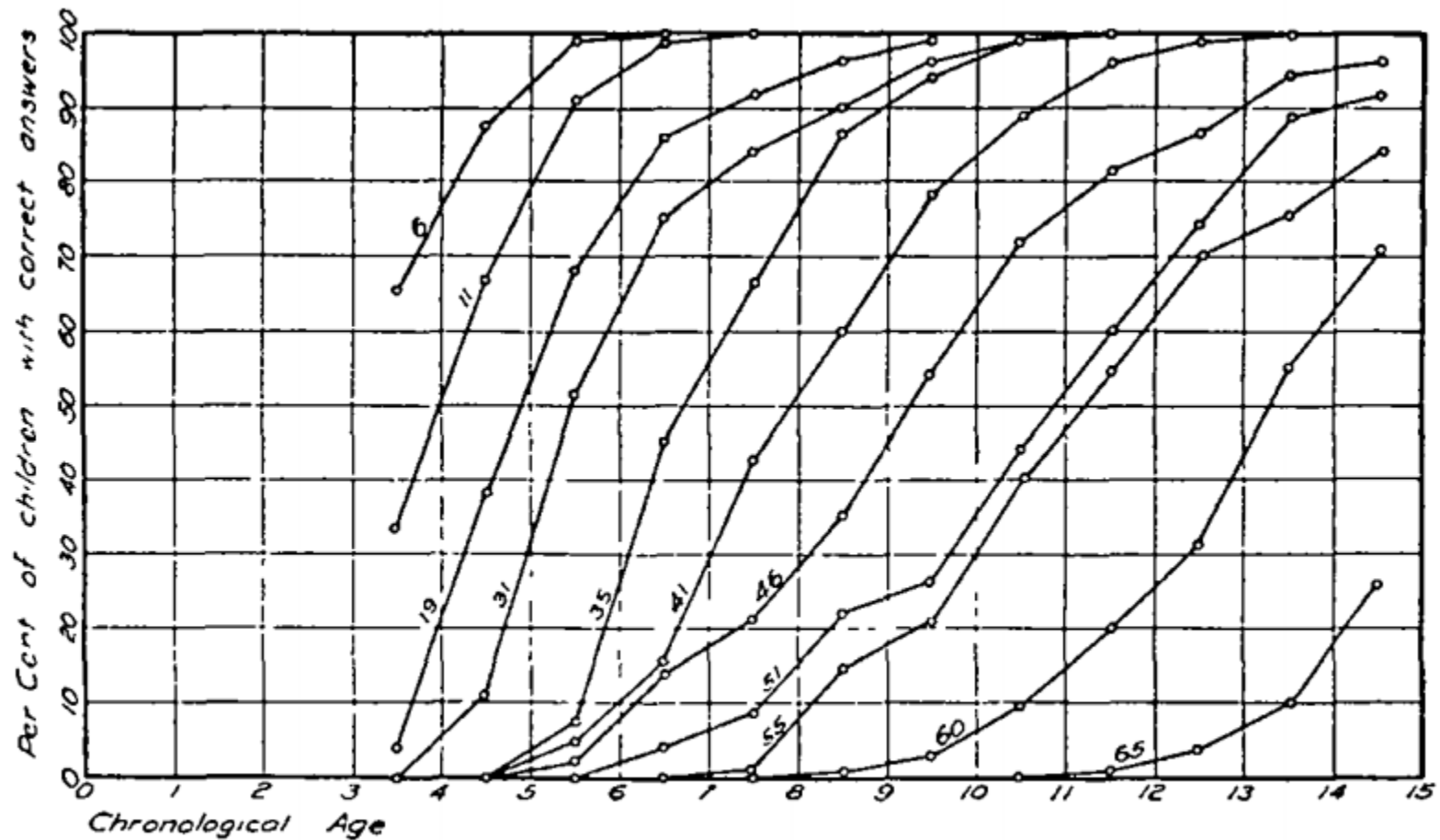


FIG. 5.

Proportions of correct response to selected items from the Binet-Simon test among children in successive age groups (Thurstone, 1925)

IRT Origins

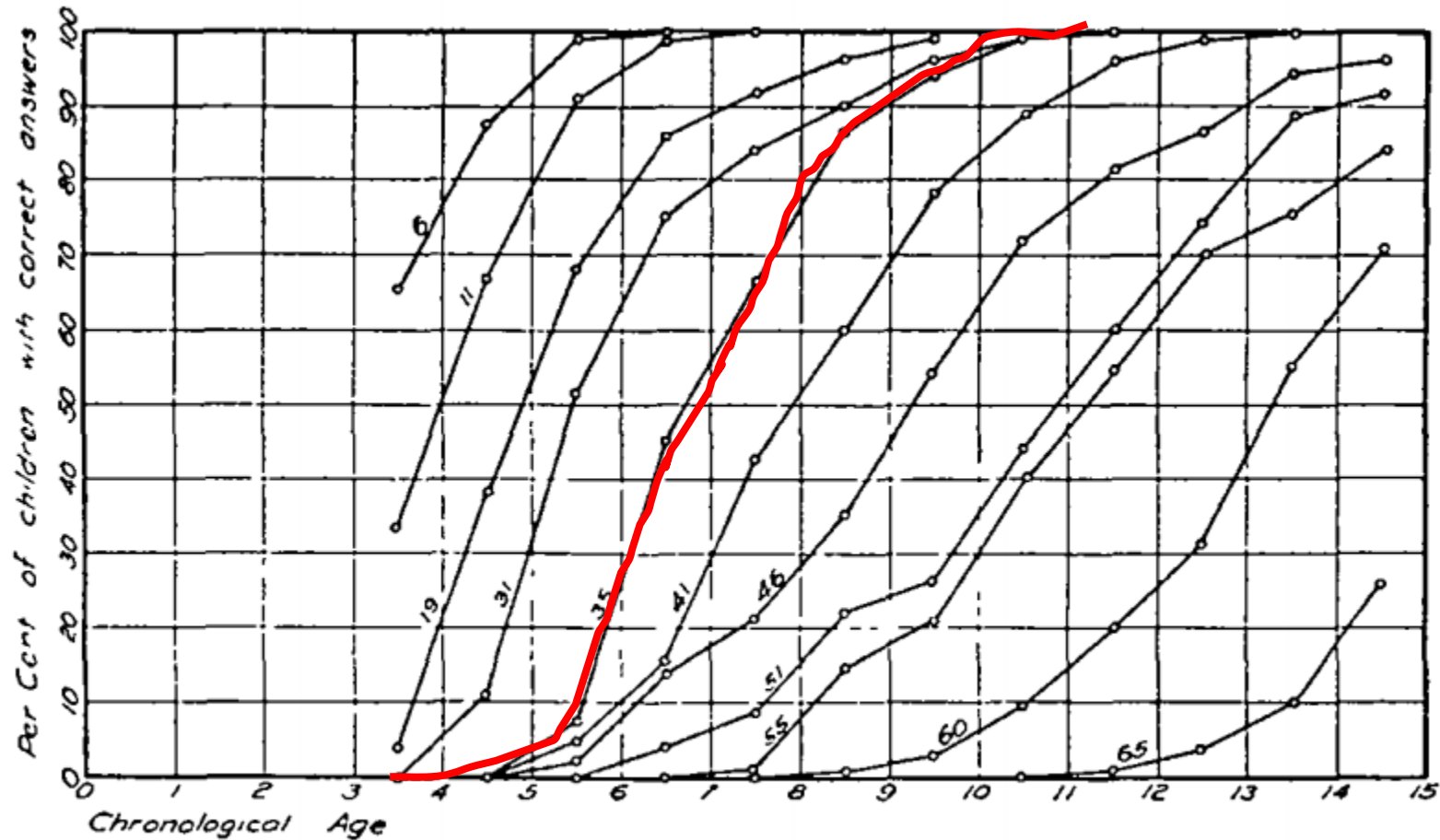
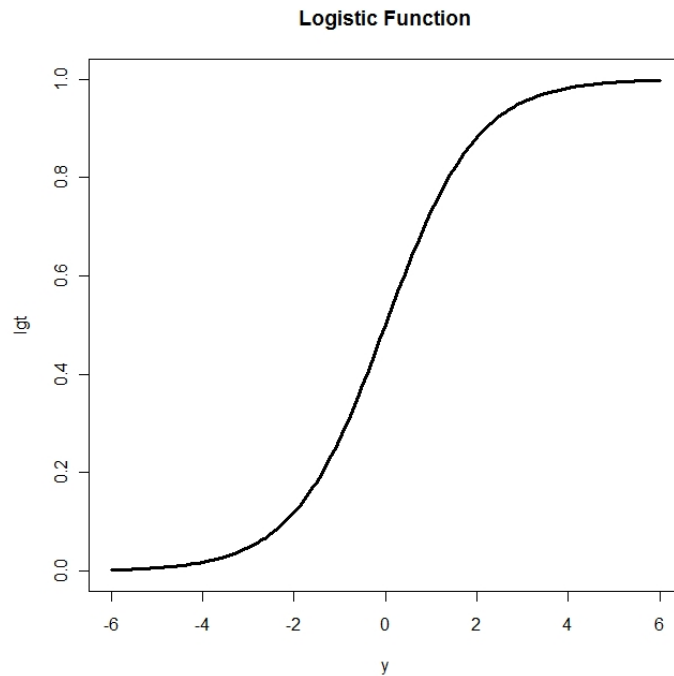


FIG. 5.

Proportions of correct response to selected items from the Binet-Simon test among children in successive age groups (Thurstone, 1925)

Logistic Curve

$$f(z) = \frac{1}{1 + e^{-z}}$$

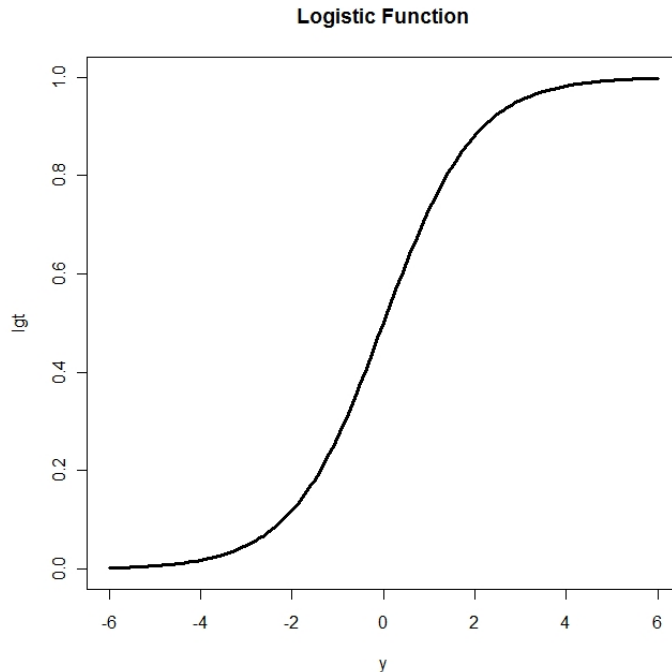


Logistic Curve

Logistic Regression

$$z = \beta_0 + \beta_1 x_1$$

$$f(z) = \frac{1}{1 + e^{-z}}$$



Logistic Curve

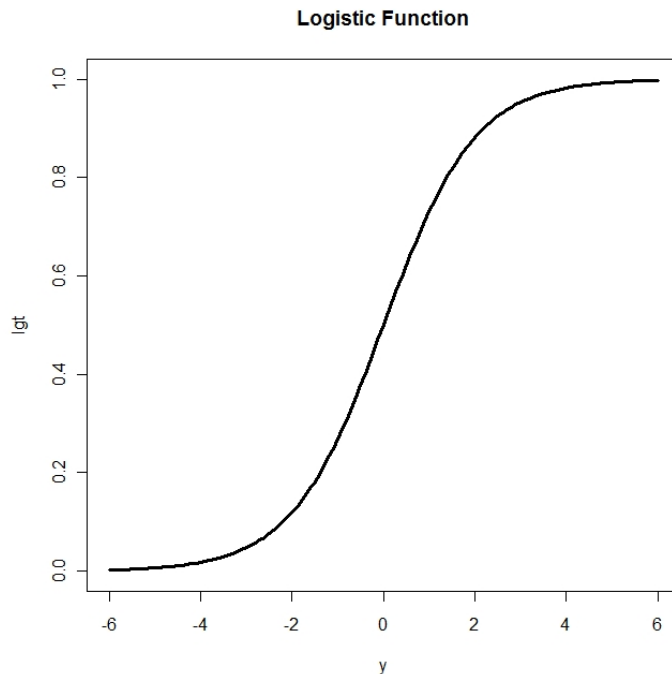
$$f(z) = \frac{1}{1 + e^{-z}}$$

Logistic Regression

$$z = \beta_0 + \beta_1 x_1$$

Item Response Theory

$$z = a\theta + b$$



Logistic Curve

$$f(z) = \frac{1}{1 + e^{-z}}$$

Logistic Regression

$$z = \beta_0 + \beta_1 x_1$$

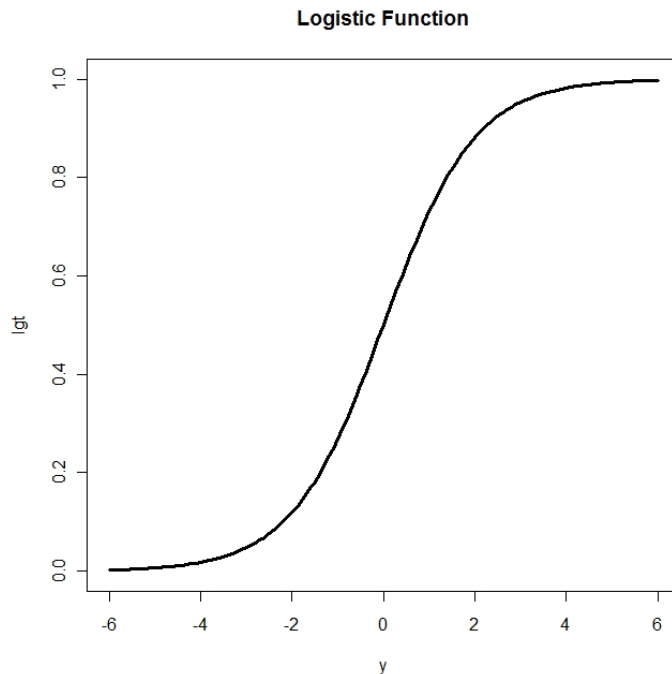
Item Response Theory

$$z = a\theta + b$$

$$P_j(\theta) = \frac{1}{1 + e^{-(a_j\theta + b_j)}}$$

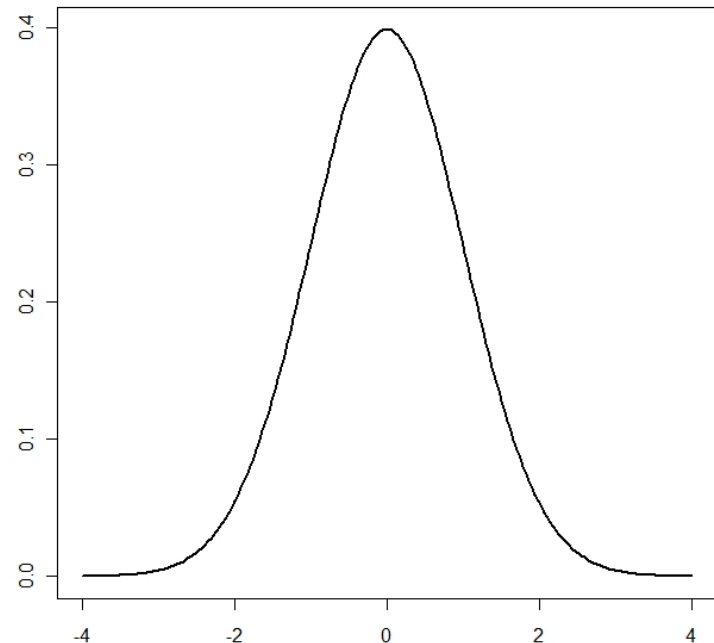
a_j = item discrimination

b_j = item location



θ

- Represents individual differences on construct or “latent trait” being measured
- The human capacity measured by the test
 - i.e. knowledge level, cognitive ability, personality characteristic
- Z-score interpretation



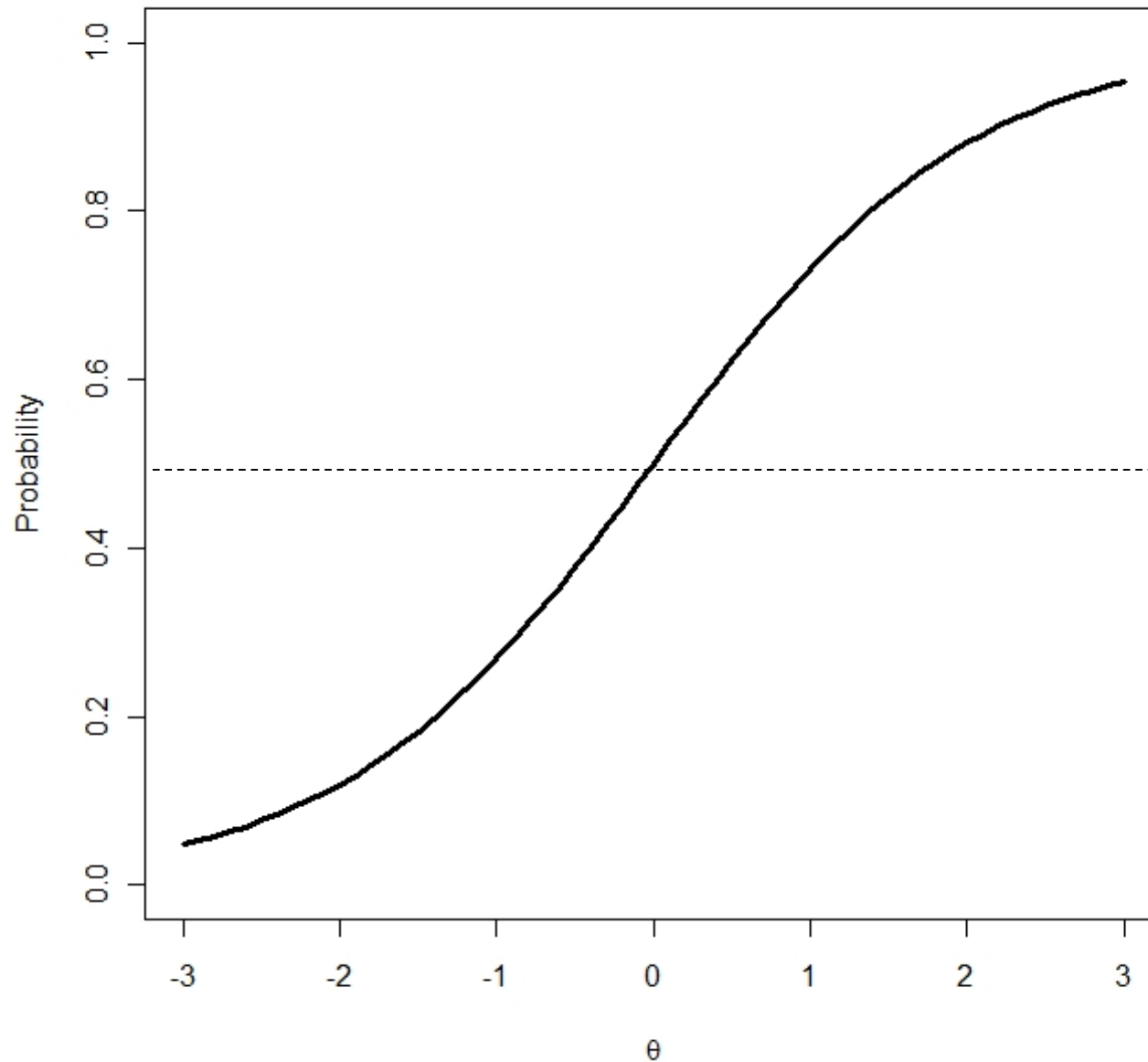
IRT Terms

- **Assumption of local independence:** a response to a question is independent of responses to other questions in a scale after controlling for the latent trait measured by the scale
- **Assumption of unidimensionality:** the set of questions are measuring a single continuous latent variable
- **Information function (IIF/TIF):** an index indicating the range of trait level over which an item or test is most useful for distinguishing among individuals
 - Characterizes the precision of measurement for persons at different levels of the underlying latent construct, with higher information denoting more precision
- **Item:** question on a scale

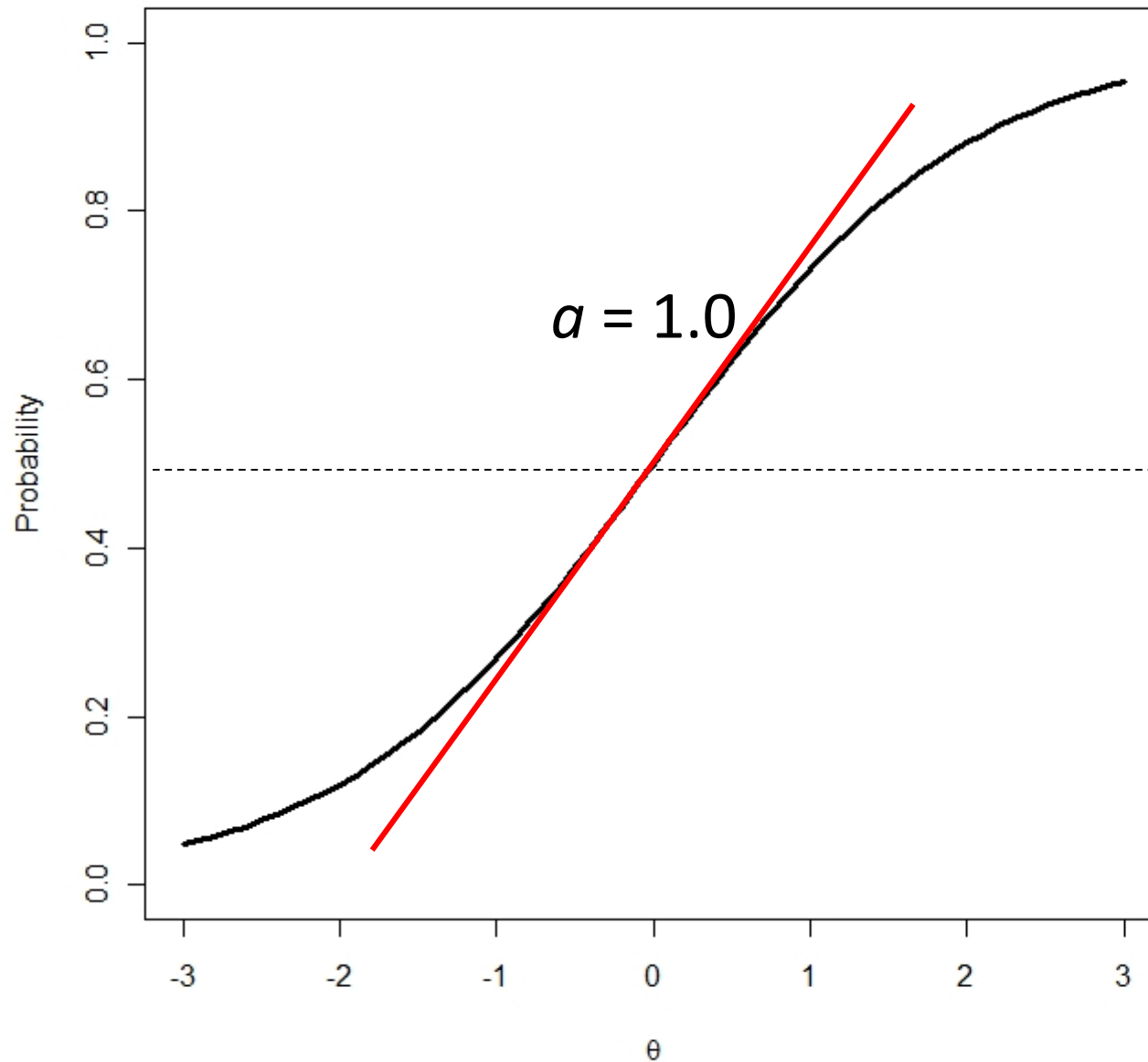
More IRT Terms

- **Item Characteristic Curve (ICC):** The ICC models the relationship between a person's probability for endorsing an item category and the level on the construct measured by the scale
- **Item Difficulty (threshold) parameter b :** the point on the latent scale where a person has a 50% chance of responding positively to the scale item
- **Item Discrimination (slope) parameter a :** describes the strength of an item's discrimination between people with trait levels below and above the threshold b .
 - The a parameter may also be interpreted as describing how an item may be related to the trait measured by the scale
- **Test Characteristic Curve (TCC):** describes the expected number of scale items endorsed as a function of the underlying latent variable
- **Theta (θ):** the unobservable construct being measured by the questionnaire

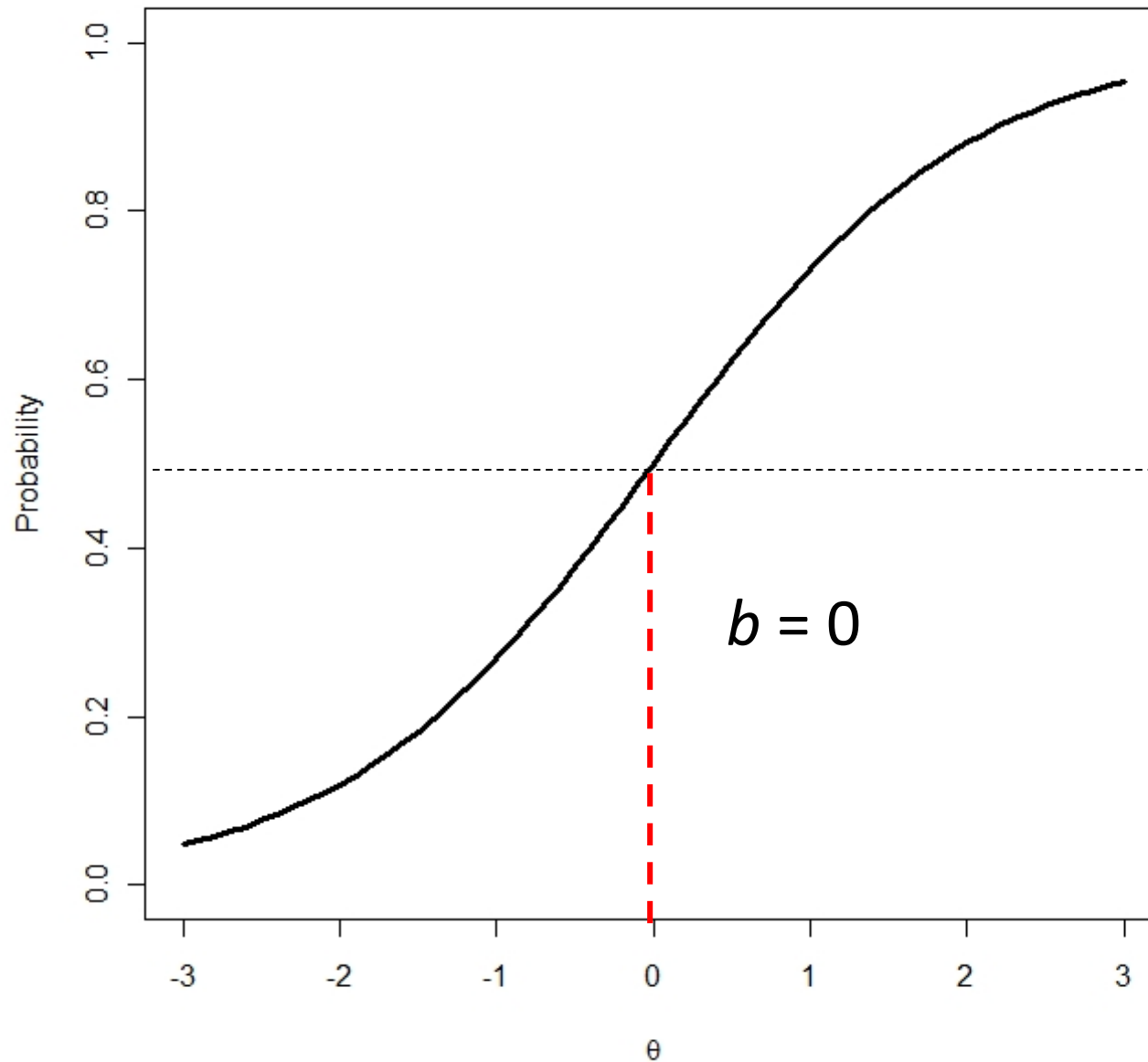
Item Characteristic Curve



Item Characteristic Curve

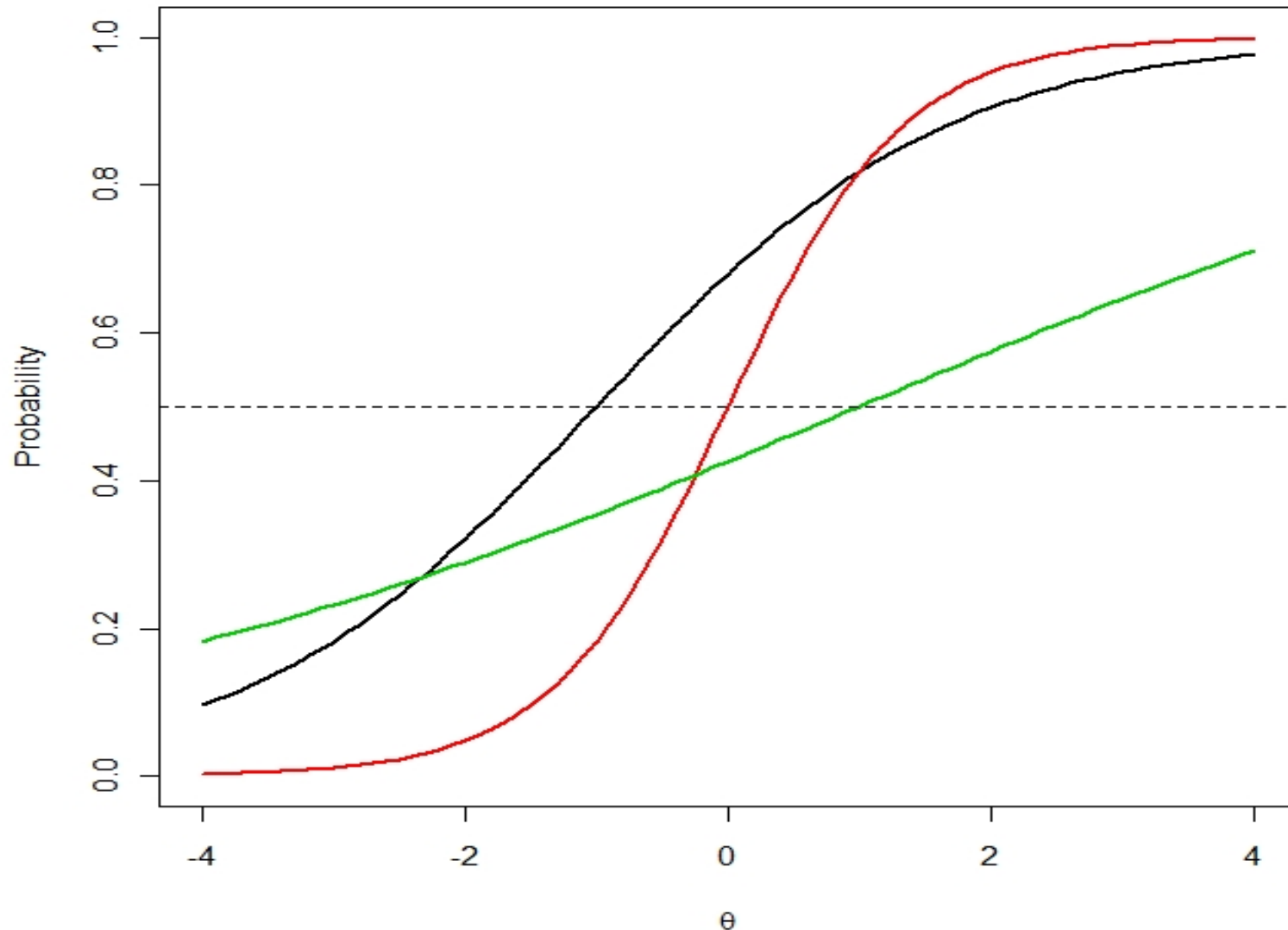


Item Characteristic Curve



Item Characteristic Curves

$a = 0.75, 1.5, 0.3$ $b = -1, 0, 1$



CTT versus IRT

Differences between CTT and IRT

- Test Precision

- Reliability

$$\alpha = \frac{p}{p-1} \left[1 - \frac{\sum_{k=1}^p \sigma_{X_k}^2}{\sigma_X^2} \right]$$

- $\alpha = .97$

- Information

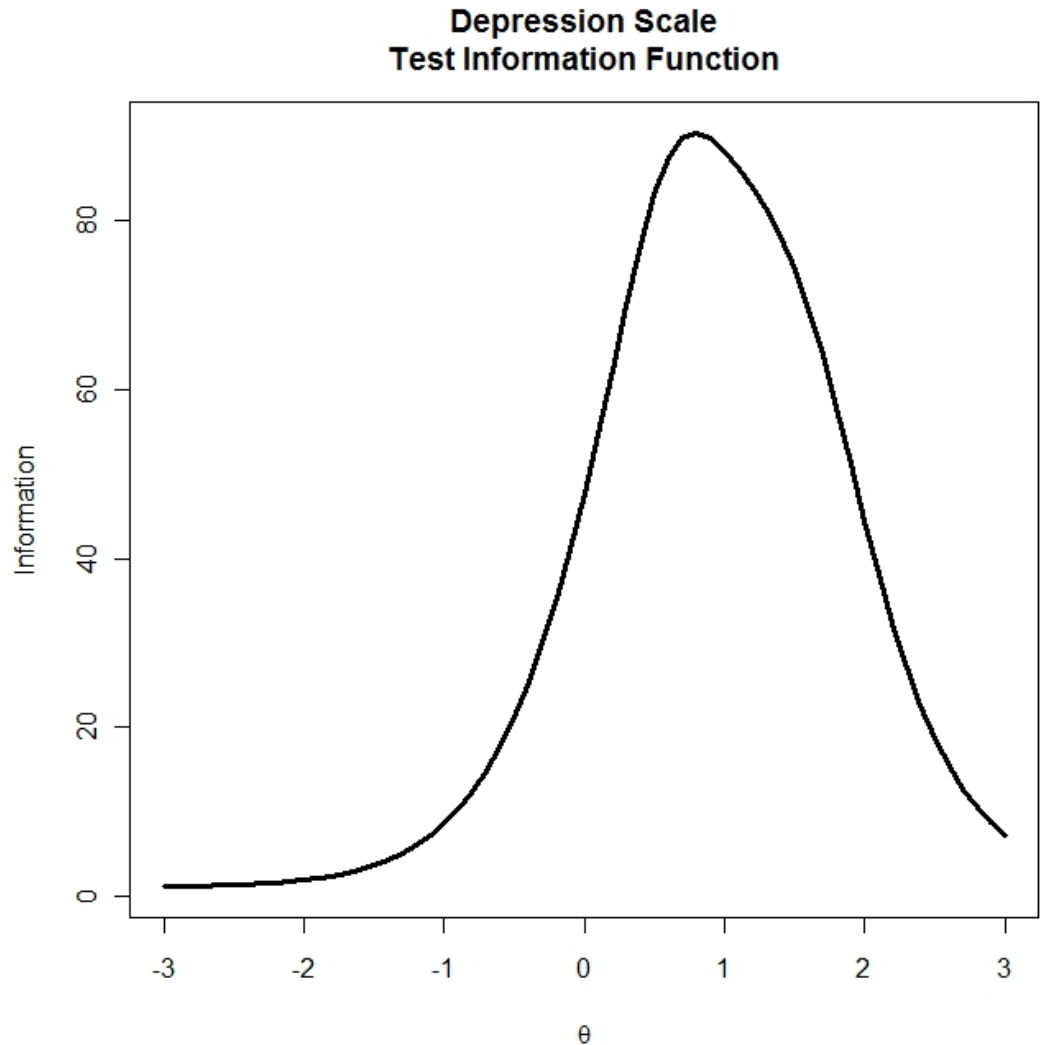
- For 1- and 2-PL

$$I(\theta) = a_i^2 P_i(\theta)(1 - P_i(\theta))$$

$$TIF(\theta) = \sum_{i=1}^I I(\theta)$$

- Composite reliability

$$r_{tt} = \frac{\sigma_{\theta}^2}{\sigma^2}$$



Differences between CTT and IRT

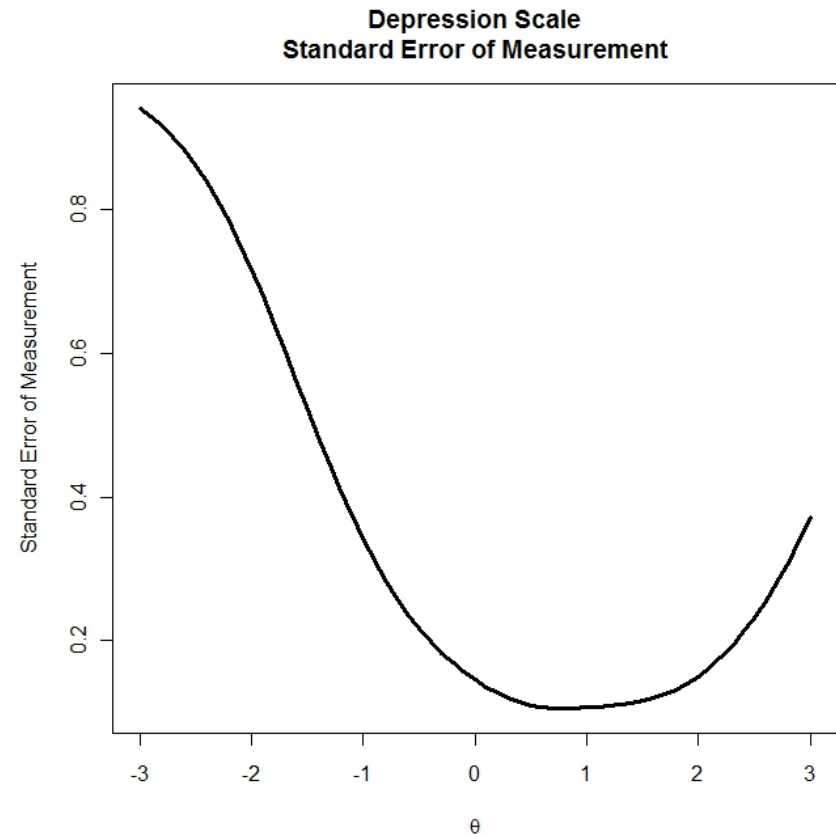
- Errors of Measurement

- Constant for all scores

- $SEM = \sqrt{(1 - r_{tt})}\sigma$
 - $SEM = 3.08$

- Depends on trait level

- $SEM|\theta = \frac{1}{\sqrt{TIF|\theta}}$
 - Composite error
 - $\sigma_\theta = \frac{SEM|\theta}{\theta}$



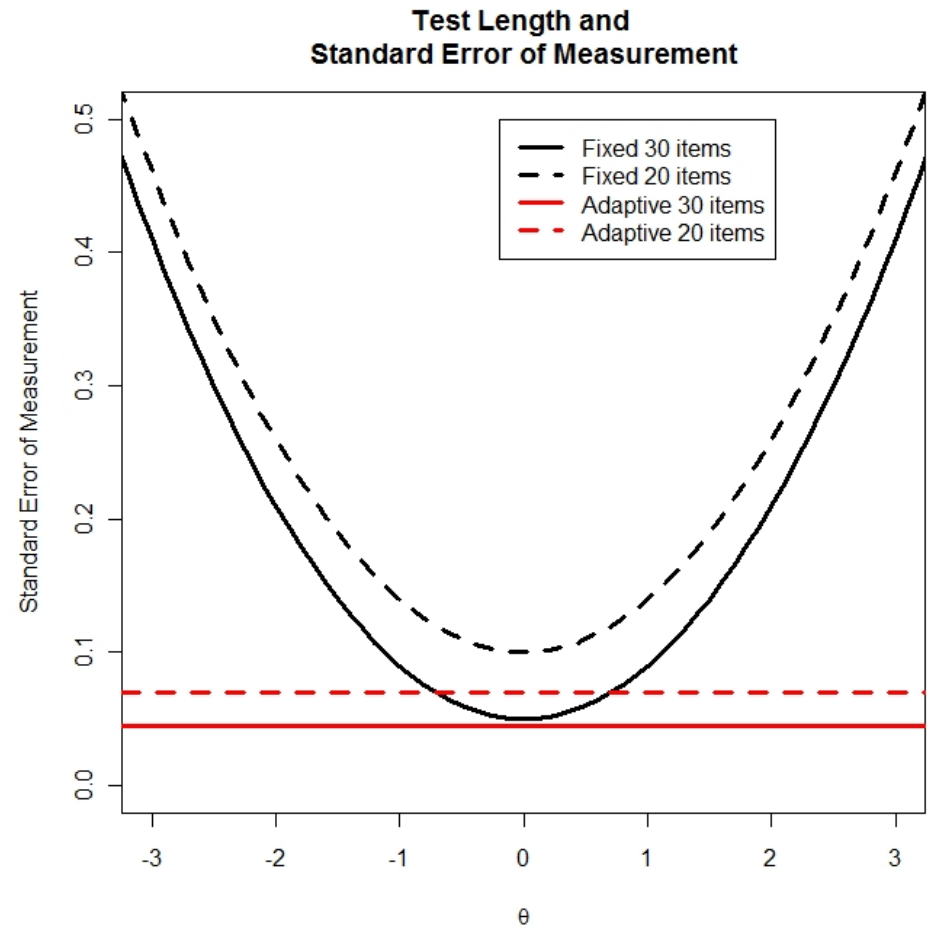
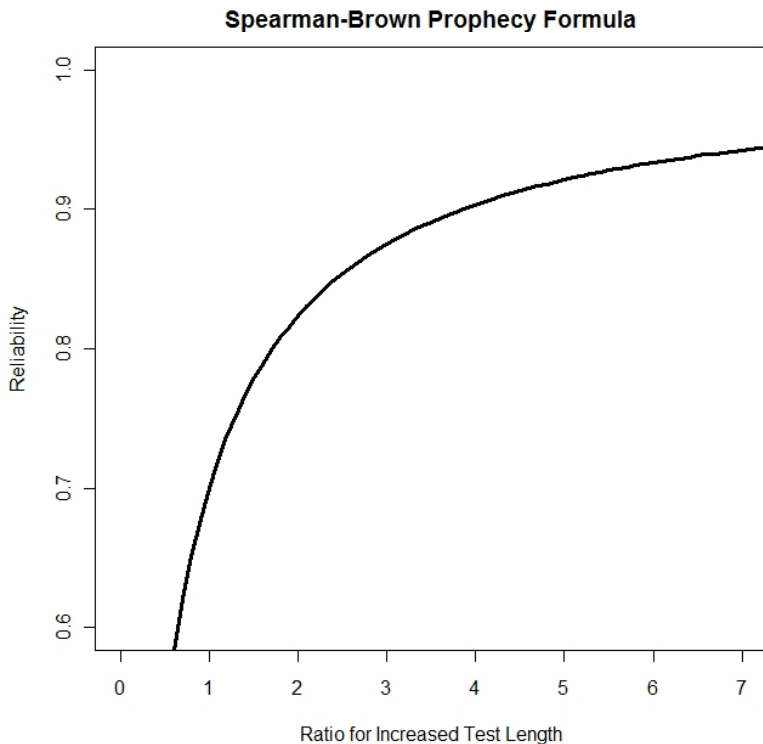
Differences between CTT and IRT

- Test Length

- Longer tests are more reliable

- $\rho = \frac{k\rho}{1+(k-1)\rho}$

- Shorter tests can be more reliable

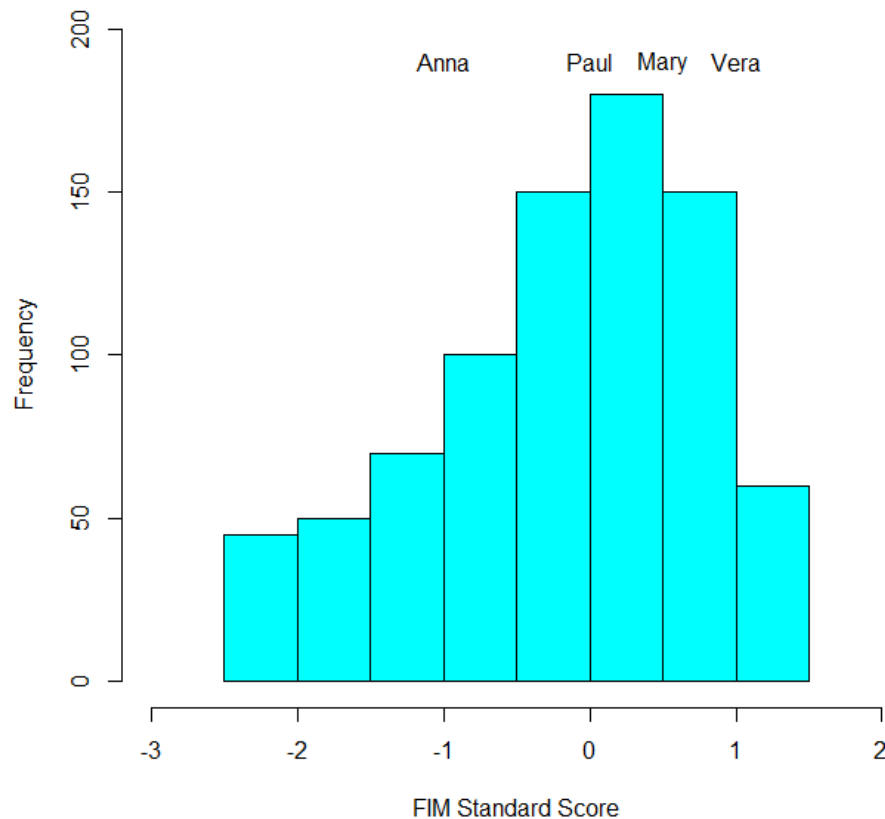


Differences between CTT and IRT

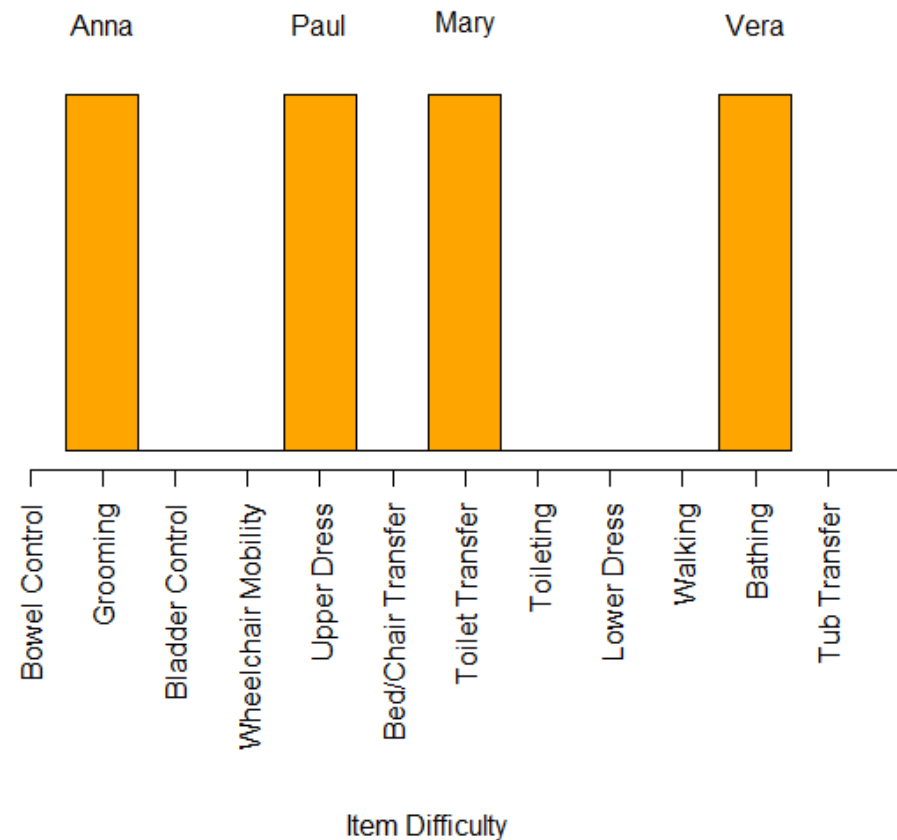
- Interchangeable test forms
 - Tests must be parallel to compare across multiple forms
 - Comparing multiple forms is optimal when difficult levels vary between persons
- Unbiased assessment of item properties
 - Requires representative sample
 - May be obtained from unrepresentative samples
- Change scores
 - Cannot be meaningfully compared when initial score levels differ
 - Can be meaningfully compared when initial score levels differ

Differences between CTT and IRT

- Compare individuals' position in a norm group



- Compare individuals' distance from items



Differences between CTT and IRT

- Different Response Formats
 - Problem
 - No Problem
- Test properties
 - Sample dependent
 - Sample free – given that the model fits that data
- Scoring Individuals
 - Raw Scores – summed scores are on an ordinal scale
 - Trait Level Estimates – theta scores are on an interval scale
- Factor analysis of dichotomous items
 - Produces artifacts rather than factors – creates difficult factors from phi correlations
 - Yields full information factor analysis

IRT Assumptions

- Unidimensionality: items are measuring a single continuous latent variable θ ranging from $-\infty$ to $+\infty$

- Evaluate dimensionality using ECV index (Reise, 2012)

1. Estimate bifactor model from poly/tetrachoric correlations
2. Apply a Schmid-Leiman (SL) transformation to the factor loadings to obtain a semi-restricted exploratory hierarchical solution
3. ECV_2 - Ratio of variance explained by the general factor to variance explained by general and group factors

$$ECV_2 = \frac{\sum(\lambda_G)^2}{\sum(\lambda_G)^2 + \sum(\lambda_{g1})^2 + \sum(\lambda_{g2})^2 + \sum(\lambda_{g3})^2}$$

- Violation

- Multidimensional IRT
- Separate IRT models for each subscale (dimension)

- Local Independence: item responses are independent of one another

- Evaluate using Yen's (1984) Q3 correlations

- Evaluate further if Yen's Q3 correlations diverge from the expected value of $-1/(n - 1)$, where n is the number of items

- Violation

- Testlet: combine two dependent items into an item with ordered levels

Dichotomous IRT Models

- Models the relationship between an unobserved variable and the probability of the examinee correctly responding to any particular item
- Common dichotomous IRT models
 - 1PL or Rasch model
 - 2PL
 - 3PL

2 Theories toward measurement

1PL

- Goal is to develop a well-fitting model to reflect the data by parameterizing the ability of interest as well as the properties of the items
- The model should reflect the properties of the data
- Items are assumed to measure as they do, not as they should
- Measurement is to explain the data
- Only requires the slope to be equal for all items

Rasch

- The data must fit the specific measurement properties defined by the model
- If the item or person does not fit, then they are discarded
- Believes optimal measurement is defined mathematically
- *Specific objectivity*: comparison of two items' difficulty are assumed to be independent; and subjects' trait levels are not dependent on any subset of items administered
- Slope is fixed at 1 for all items

Common IRT models

1PL – items can vary in difficulty

$$P(\theta) = \frac{1}{1 + \exp[-a(\theta - b_i)]}$$

2PL – items can vary in discrimination and difficulty

$$P(\theta) = \frac{1}{1 + \exp[-a_i(\theta - b_i)]}$$

3PL – items can vary in discrimination, difficulty, and **guessing** (respondents may over report desirable, or underreport embarrassing behaviors)

$$P(\theta) = c_i + \frac{1 - c_i}{1 + \exp[-a_i(\theta - b_i)]}$$

Example:

Rosenberg Self-Esteem Scale (1965)

1. On the whole, I am satisfied with myself.
2. At times, I think I am not good at all (R)
3. I feel that I have a number of good qualities
4. I am able to do things as well as most other people.
5. I feel I do not have much to be proud of. (R)
6. I certainly feel useless at times. (R)
7. I feel that I'm a person of worth, at least on an equal plane with others.
8. I wish I could have more respect for myself (R)
9. All in all, I am inclined to feel that I am a failure (R)
10. I take a positive attitude toward myself.

Response format: 0 = Strongly Disagree, 1 = Disagree, 2 = Agree, 3 = Strongly Agree

Recoded into dichotomies: 0 = Disagree, 1 = Agree

N = 441

Very Brief Introduction to R

Download R for your system

<http://cran.stat.ucla.edu>

The Comprehensive R Archive Network

Download and Install R

Precompiled binary distributions of the base system and contributed packages, **Windows and Mac** users most likely want one of these versions of R:

- [Download R for Linux](#)
- [Download R for MacOS X](#)
- [Download R for Windows](#)

Choose 'base'

R for Windows

Subdirectories:

[base](#)

[contrib](#)

[old contrib](#)

[Rtools](#)

Binaries for base distribution (managed by Duncan Murdoch). This is what you want to [install R for the first time](#).

Binaries of contributed CRAN packages (for R \geq 2.11.x; managed by Uwe Ligges). There is also information on [third party software](#) available for CRAN Windows services and corresponding environment and make variables.

Binaries of contributed CRAN packages for outdated versions of R (for R $<$ 2.11.x; managed by Uwe Ligges).

Tools to build R and R packages (managed by Duncan Murdoch). This is what you want to build your own packages on Windows, or to build R itself.

Please do not submit binaries to CRAN. Package developers might want to contact Duncan Murdoch or Uwe Ligges directly in case of questions / suggestions related to Windows binaries.

You may also want to read the [R FAQ](#) and [R for Windows FAQ](#).

Note: CRAN does some checks on these binaries for viruses, but cannot give guarantees. Use the normal precautions with downloaded executables.

Download R



R-3.3.2 for Windows (32/64 bit)

[Download R 3.3.2 for Windows](#) (62 megabytes, 32/64 bit)

[Installation and other instructions](#)

[New features in this version](#)

[CRAN](#)

[Mirrors](#)

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[The R Journal](#)

If you want to double-check that the package you have downloaded matches the package distributed by CRAN, you can compare the [md5sum](#) of the .exe to the [fingerprint](#) on the master server. You will need a version of md5sum for windows: both [graphical](#) and [command line versions](#) are available.

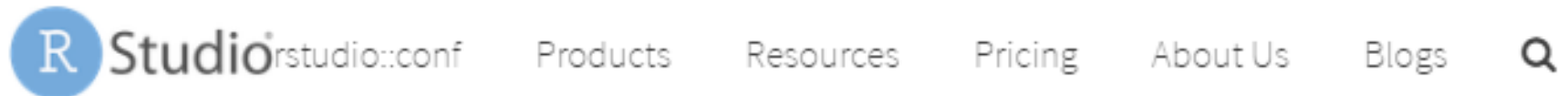
Frequently asked questions

- [Does R run under my version of Windows?](#)
- [How do I update packages in my previous version of R?](#)

Downloading RStudio

<https://www.rstudio.com/products/rstudio/download/>

- Choose the FREE Open Source Edition



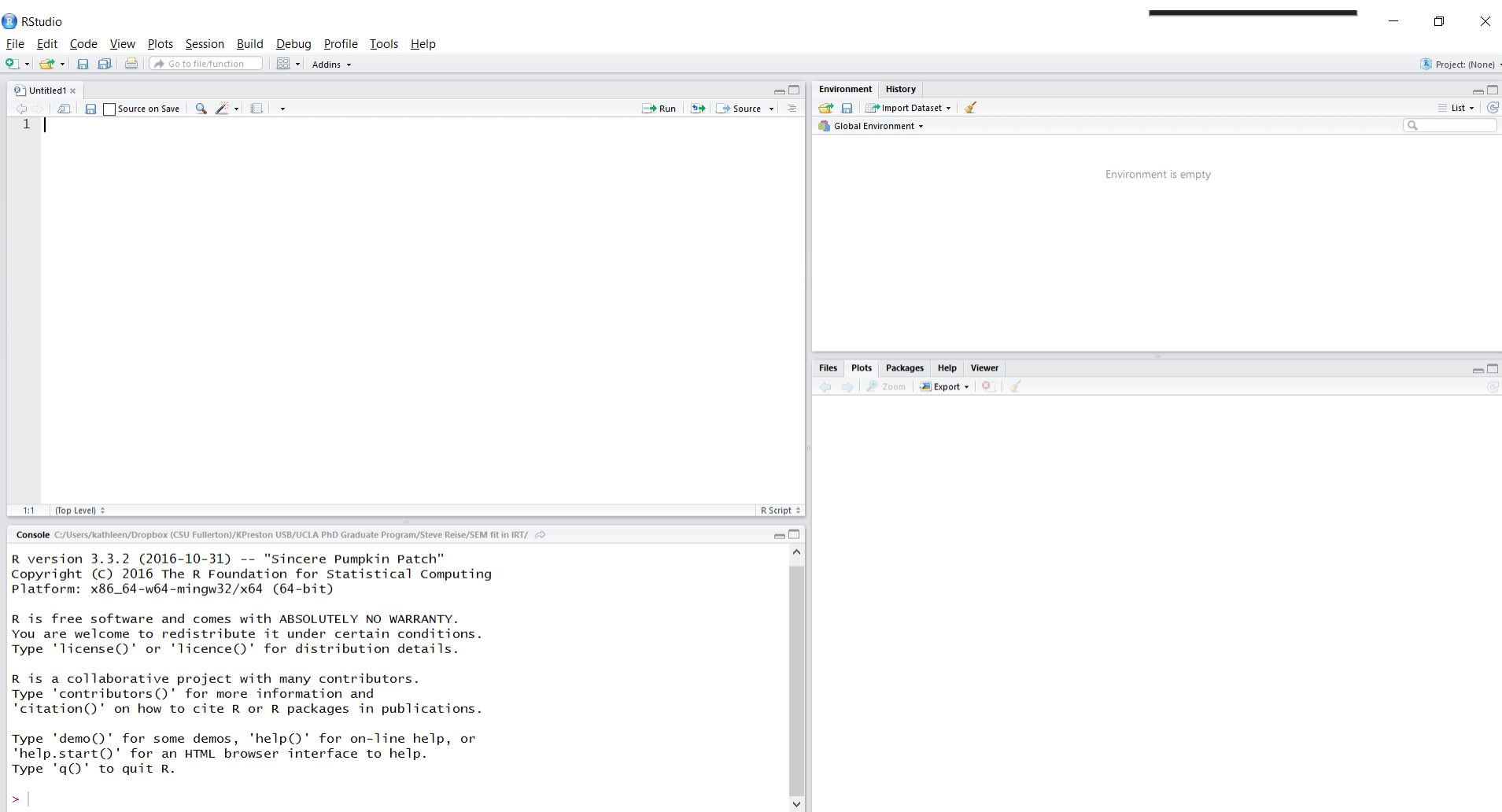
RStudio Desktop 1.0.136 — Release Notes

RStudio requires R 2.11.1+. If you don't already have R, download it [here](#).

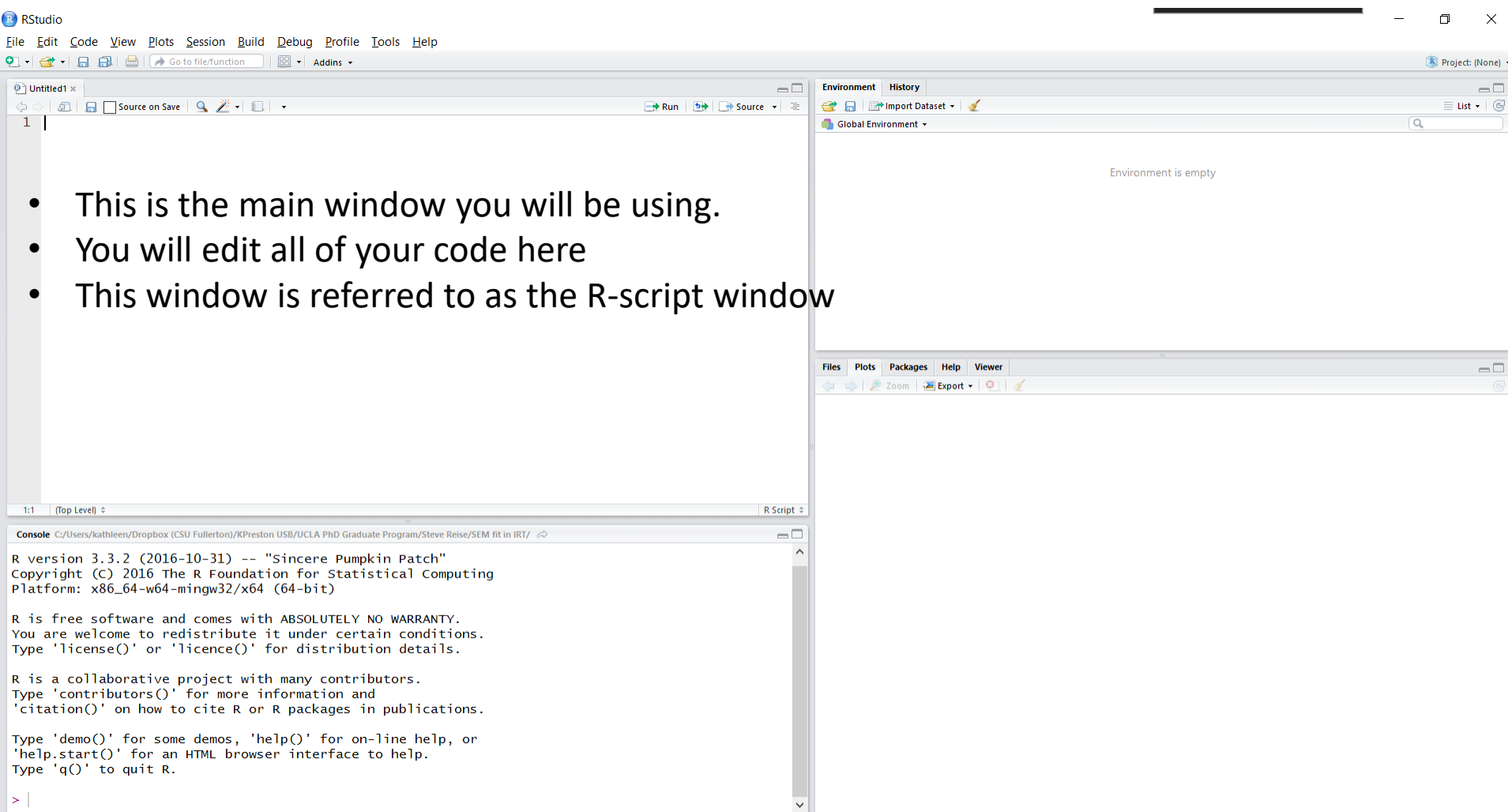
Installers for Supported Platforms

Installers	Size	Date	MD5
RStudio 1.0.136 - Windows Vista/7/8/10	81.9 MB	2016-12-21	93b3f307f567c33f7a4db4c114099b3e
RStudio 1.0.136 - Mac OS X 10.6+ (64-bit)	71.2 MB	2016-12-21	12d6d6ade0203a2fce6fe3dea65c1ae
RStudio 1.0.136 - Ubuntu 12.04+/Debian 8+ (32-bit)	85.5 MB	2016-12-21	0a20fb89d8aaeb39b329a640ddadd2c5
RStudio 1.0.136 - Ubuntu 12.04+/Debian 8+ (64-bit)	92.1 MB	2016-12-21	2a73b88a12a9fbaf96251cecf8b41340
RStudio 1.0.136 - Fedora 19+/RedHat 7+/openSUSE 13.1+ (32-bit)	84.7 MB	2016-12-21	fa6179a7855bff0f939a34c169da45fd
RStudio 1.0.136 - Fedora 19+/RedHat 7+/openSUSE 13.1+ (64-bit)	85.7 MB	2016-12-21	2b3a148ded380b704e58496befb55545

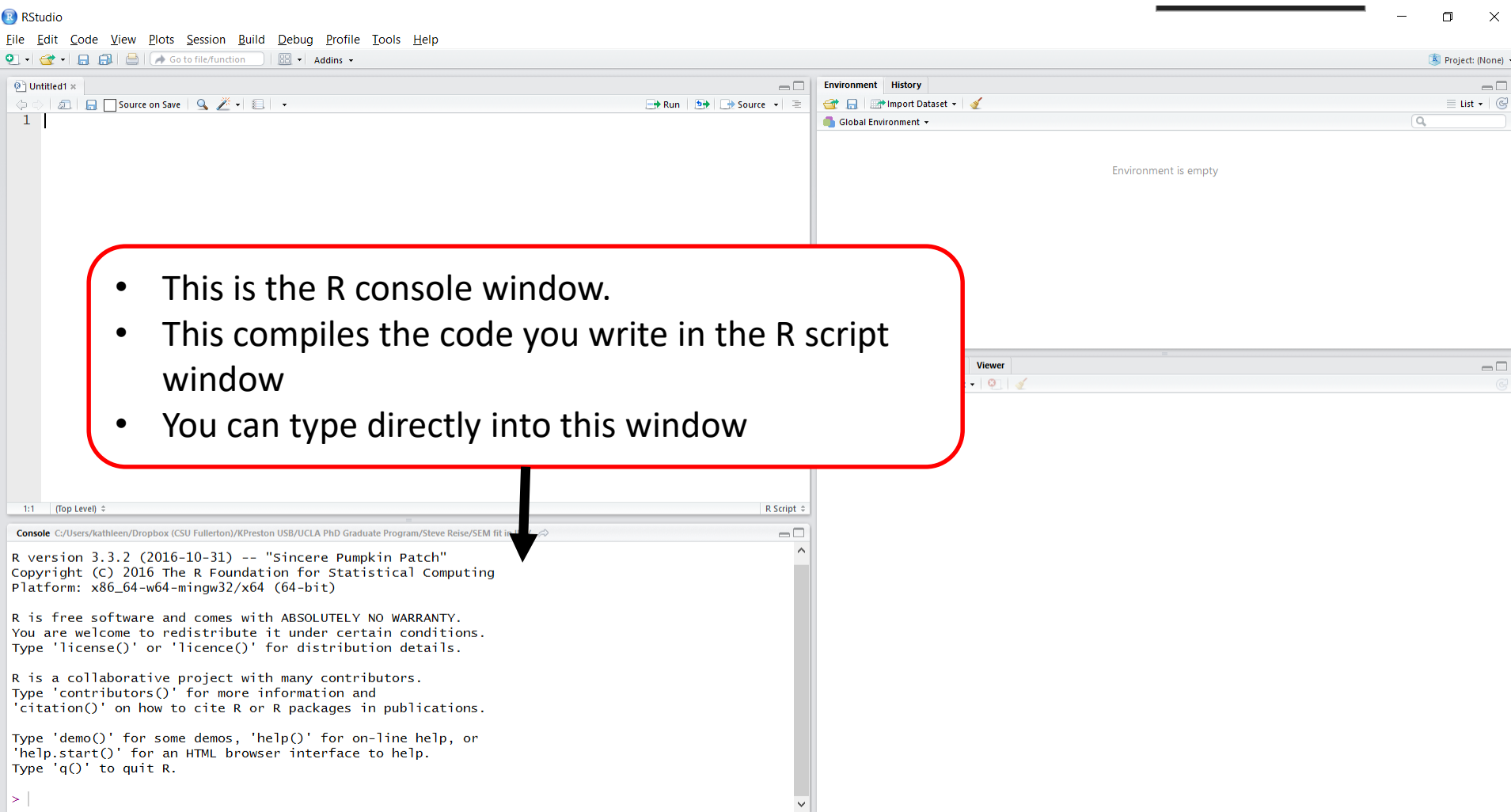
Navigating RStudio



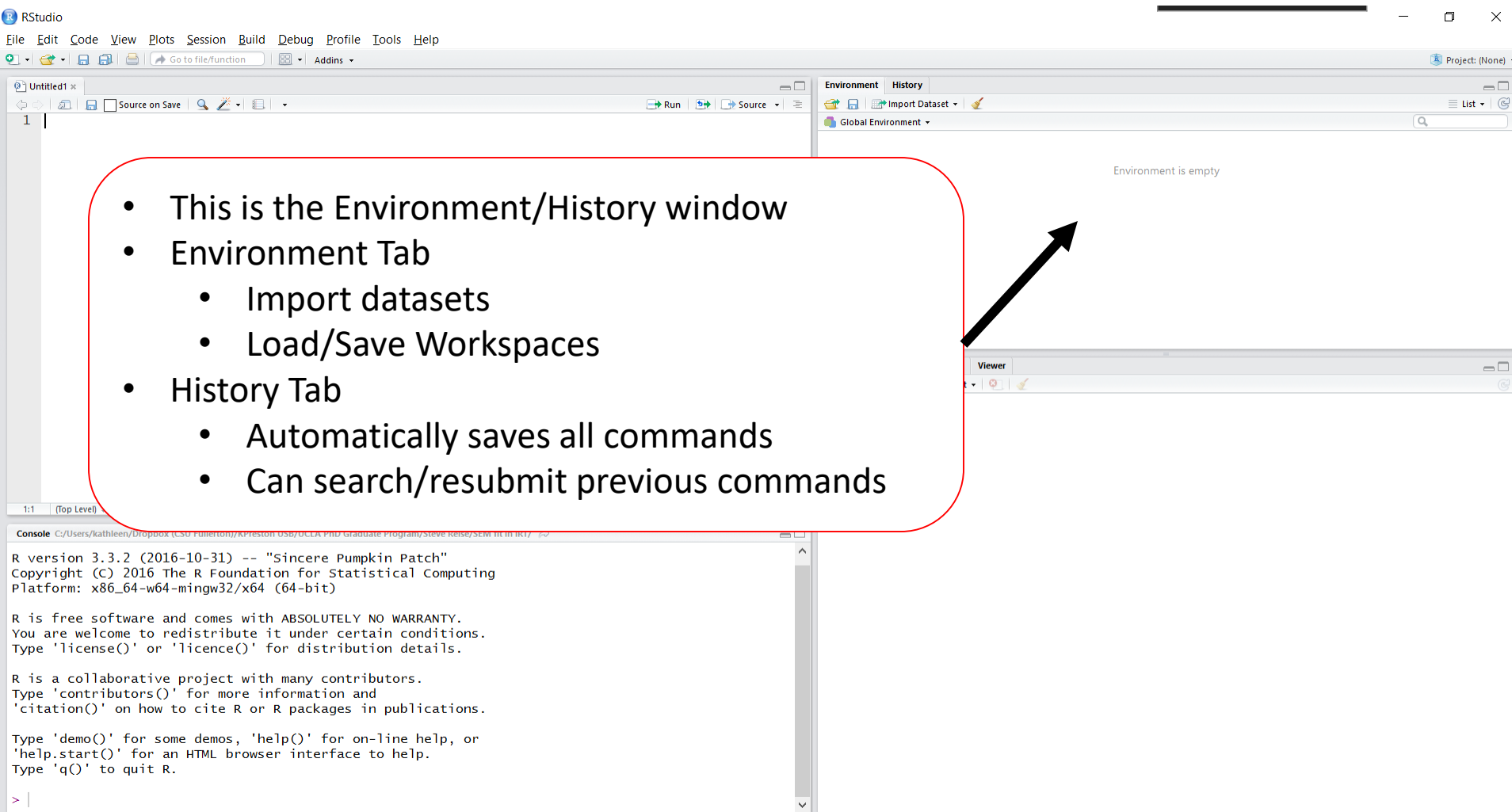
RStudio: R-script



RStudio: R console



RStudio: Workspace/History



The screenshot shows the RStudio interface. The top menu bar includes File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, and Help. The main editor window shows a file named 'Untitled1.R' with a single line of code: '1 |'. The Environment pane on the right is active, showing 'Global Environment' and 'Environment is empty'. The History pane is visible below it. A red rounded rectangle highlights the Environment pane, and a black arrow points from the History pane to the Environment pane.

- This is the Environment/History window
- Environment Tab
 - Import datasets
 - Load/Save Workspaces
- History Tab
 - Automatically saves all commands
 - Can search/resubmit previous commands

Console output:

```
R version 3.3.2 (2016-10-31) -- "Sincere Pumpkin Patch"
Copyright (C) 2016 The R Foundation for Statistical Computing
Platform: x86_64-w64-mingw32/x64 (64-bit)

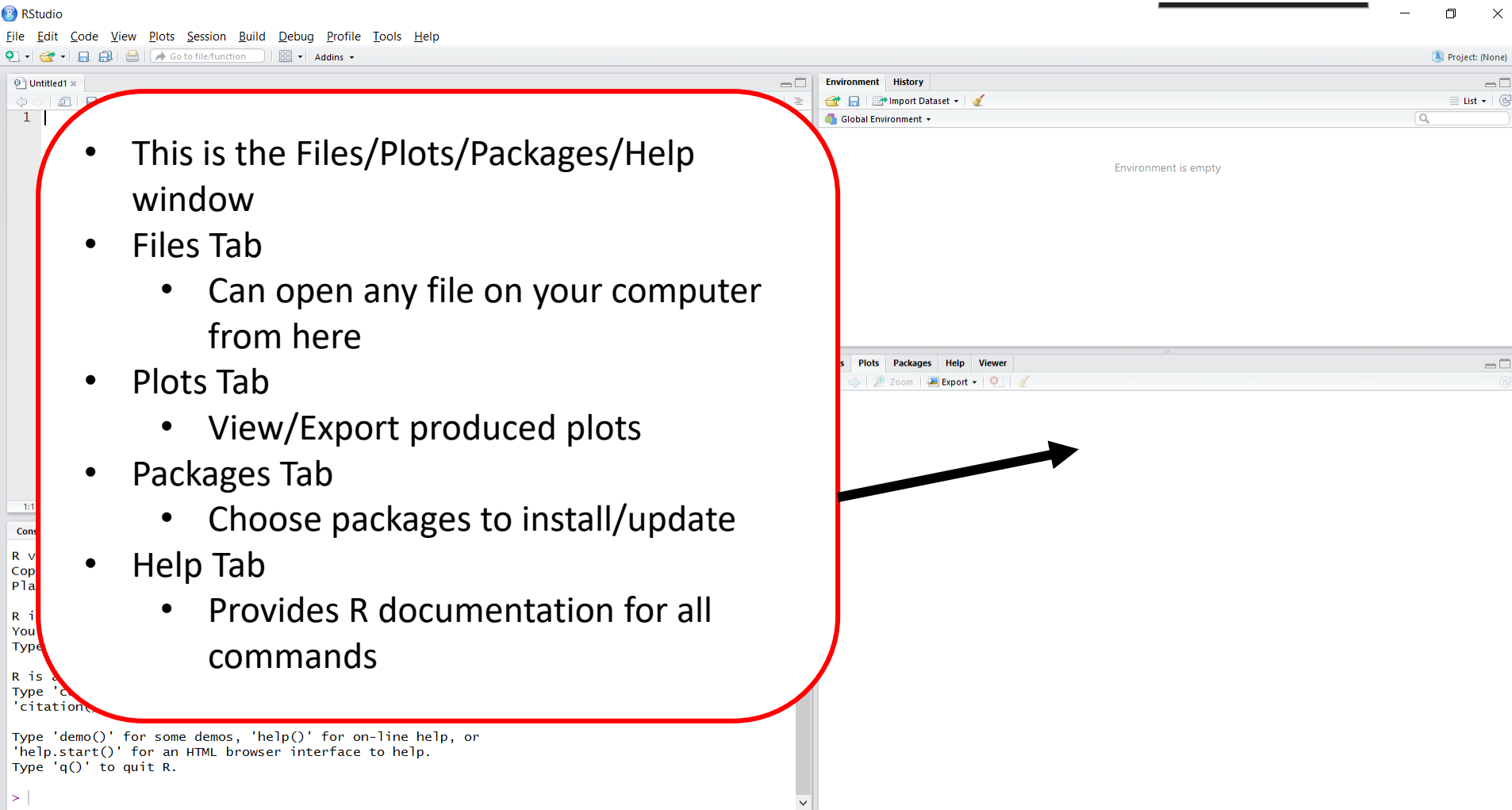
R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.

R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

> |
```


RStudio: Files/Plots/Packages/Help



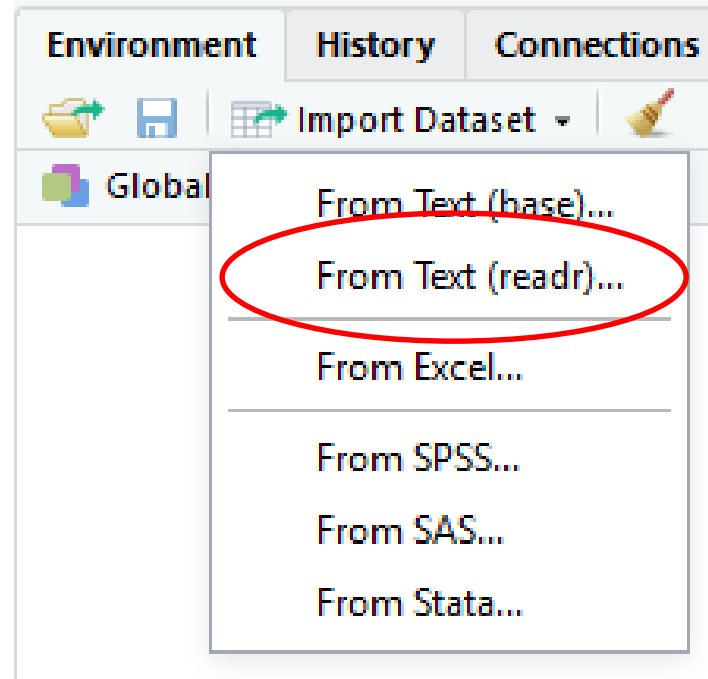
The screenshot shows the RStudio application window. The top menu bar includes File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, and Help. Below the menu bar is a toolbar with icons for file operations and a 'Go to file/function' search bar. The main workspace is divided into four panes: Environment, History, Plots, and Packages/Help. The Environment pane shows 'Global Environment' and 'Environment is empty'. The History pane is empty. The Plots pane is empty. The Packages/Help pane is empty. A red rounded rectangle highlights the Files, Plots, Packages, and Help tabs. An arrow points from the text box to the Plots tab.

- This is the Files/Plots/Packages/Help window
- Files Tab
 - Can open any file on your computer from here
- Plots Tab
 - View/Export produced plots
- Packages Tab
 - Choose packages to install/update
- Help Tab
 - Provides R documentation for all commands

RStudio: Workspace

Importing a Dataset

- Import Dataset
 - Can import from the following formats
 - .csv or .dat or .txt
 - .xls or .xlsx
 - .sav
 - .sas7bdat
 - .dta
- Choose ChildSleepData.dat file from computer



RStudio: Workspace

Importing a Dataset

Import Text Data

File/Url:

C:/Users/kpreston/Dropbox (Personal)/Research Projects/Talks/WPA Stats Talk/2017/ChildSleepData.dat Browse...

Data Preview:

Child WakeUp AcadPerf (character)
A 11 2.4
B 9 3.6
C 9 3.2
D 12 2.2
E 7 3.8
F 10 2.2
G 10 3
H 8 3

Previewing first 50 entries.

Import Options:

Name: ☒ First Row as Names

Skip: ☒ Trim Spaces

☒ Open Data Viewer

Delimiter: Escape:

Quotes: Comment:

Locale: NA:

Code Preview:

```
library(readr)
chidsleepData <- read_csv("C:/Users/kpreston/Dropbox (Personal)/Research Projects/Talks/WPA Stats Talk/2017/childsleepData.dat")
view(chidsleepData)
```

Import Cancel

[? Reading rectangular data using readr](#)

RStudio: Workspace

Change Delimiter to 'Tab'

Import Text Data

File/Url:

Data Preview:

Child (character)	WakeUp (integer)	AcadPerf (double)
A	11	2.4
B	9	3.6
C	9	3.2
D	12	2.2
E	7	3.8
F	10	2.2
G	10	3.0
H	8	3.0

Previewing first 50 entries.

Import Options:

Name: Skip:

☒ First Row as Names ☒ Trim Spaces ☒ Open Data Viewer

Delimiter: **Tab** Quotes: **Default** Escape: **None** Comment: **Default** NA: **Default**

Locale:

Code Preview:

```
library(readr)
ChildSleepData <- read_delim("C:/Users/kpreston/Dropbox (Personal)/Research Projects/Talks/WPA Stats Talk/2017/ChildSleepData.dat",
                             "\t", escape_double = FALSE, trim_ws = TRUE)
view(ChildSleepData)
```

? Reading rectangular data using readr

RStudio: Dataset Successfully Imported

The screenshot shows the RStudio interface with the following components:

- Source Editor:** Displays a table with 8 rows and 3 columns: Child, WakeUp, and AcadPerf.
- Environment Pane:** Shows the 'Global Environment' with a data object 'ChildSleepData' containing 8 observations of 3 variables.
- Console:** Shows the R version (3.4.4) and the executed code for importing the dataset.

Arrows point from the text boxes to the corresponding elements in the RStudio interface:

- Arrow 1 points from the 'New 'ChildSleepData' Tab shows the format of the imported dataset' box to the Source Editor table.
- Arrow 2 points from the 'Environment Tab provides information about the imported dataset' box to the Environment Pane.
- Arrow 3 points from the 'R console shows the executed code' box to the Console.

Table Data:

	Child	WakeUp	AcadPerf
1	A	11	2.4
2	B	9	3.6
3	C	9	3.2
4	D	12	2.2
5	E	7	3.8
6	F	10	2.2
7	G	10	3.0
8	H	8	3.0

Console Output:

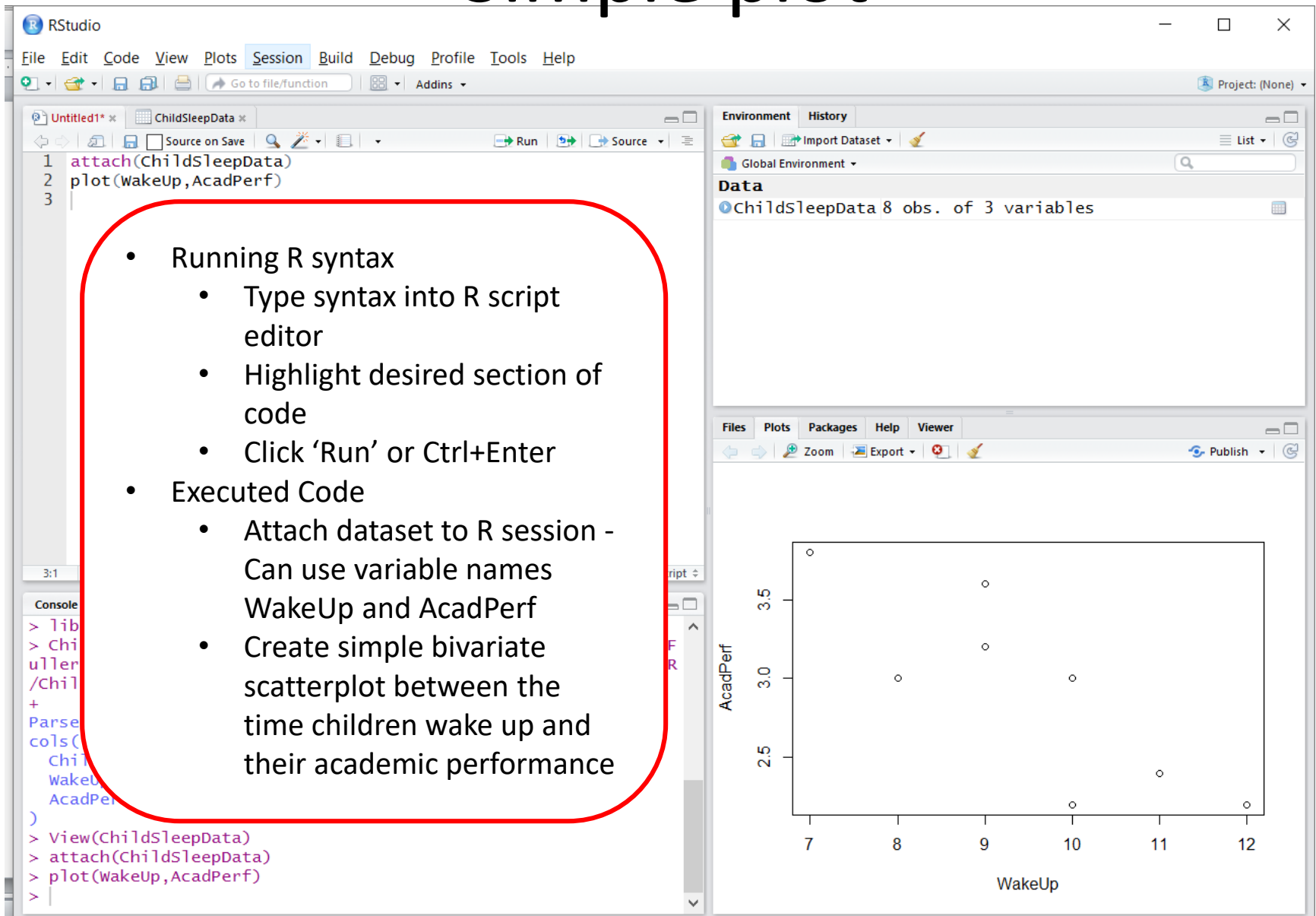
```
R version 3.4.4 (2018-03-15) -- "Someone to Lean On"  
Copyright (C) 2018 The R Foundation for Statistical Computing  
Platform: x86_64-w64-mingw32/x64 (64-bit)  
  
R is free software and comes with ABSOLUTELY NO WARRANTY.  
You are welcome to redistribute it under certain conditions.  
Type 'license()' or 'licence()' for distribution details.  
  
R is a collaborative project with many contributors.  
Type 'contributors()' for more information and  
'citation()' on how to cite R or R packages in publications.  
  
Type 'demo()' for some demos, 'help()' for on-line help, or  
'help.start()' for an HTML browser interface to help.  
Type 'q()' to quit R.  
  
> library(readr)  
> ChildSleepData <- read_delim("c:/Users/kpreston/Dropbox (Personal)/Research Projects/Talks/WPA Stats Talk/2017/ChildSleepData.dat",  
+                             "\t", escape_double = FALSE, trim_ws = TRUE)  
Parsed with column specification:  
cols(  
  child = col_character(),  
  wakeup = col_integer(),  
  AcadPerf = col_double()  
)  
> view(ChildSleepData)  
>
```

- New 'ChildSleepData' Tab shows the format of the imported dataset

- Environment Tab provides information about the imported dataset

- R console shows the executed code

Simple plot



RStudio: Plots

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Untitled1* ChildSleepData*

Source on Save Run Source

```
1 attach(ChildSleepData)
2 plot(WakeUp, AcadPerf)
3
```

- All executed plot commands are produced here
- **Zoom**
 - Opens the plot in a new window
- **Export**
 - Can save plot as:
 - Image
 - PDF
 - Copy to clipboard

Environment History

Global Environment

Data

ChildSleepData 8 obs. of 3 variables

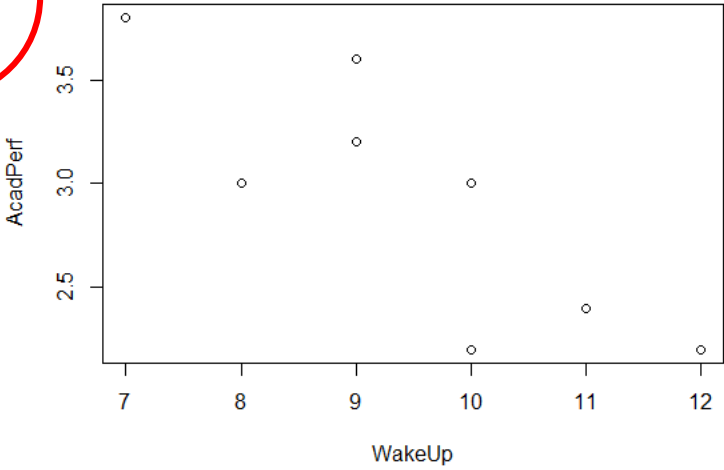
Files Plots Packages Help Viewer

Zoom Export

3:1 (Top Level) R Script

```
> library(readr)
> ChildSleepData <- read_delim("C:/Users/kathleen/Dropbox (CSU Fullerton)/CGU - Psychology 315FF/Lectures/1b - Introduction to R/ChildSleepData.dat",
+   "\t", escape_double = FALSE, trim_ws = TRUE)
Parsed with column specification:
cols(
  child = col_character(),
  WakeUp = col_integer(),
  AcadPerf = col_double()
)
> View(ChildSleepData)
> attach(ChildSleepData)
> plot(WakeUp, AcadPerf)
>
```

AcadPerf



WakeUp

The scatter plot shows the relationship between WakeUp (x-axis, ranging from 7 to 12) and AcadPerf (y-axis, ranging from 2.5 to 3.5). The data points are as follows:

WakeUp	AcadPerf
7	3.7
8	3.0
9	3.2
9	3.6
10	3.0
10	2.6
11	2.4
12	2.2

R Language Basics

Packages

- Base R and most Rpackages are available for download from the Comprehensive R Archive Network (CRAN)
 - `cran.r-project.org`
 - base R comes with a number of basic data management, analysis, and graphical tools
- We will start by installing a package, `car`, a package that is frequently used in R tutorials
 - Packages only need to be installed once
 - `install.packages("car")`
- Once the package is installed, it must be loaded
 - Packages must be loaded every time you start an R session
 - `library(car)`

R Programming Basics

- R is case sensitive.
- The # character at the beginning of a line signifies a comment, which is not executed. Each commented line must start with a #
- Help for function keywords are accessed by the help() function with the keyword inside the parentheses
- stores both data and output from data analysis (as well as everything else) in objects
- Things are assigned to and stored in objects using the <- or = operator
- Note: do NOT separate < and - with a space to create <-

DataFrames

```
# Referencing within Data Frames
# single cell value
ChildSleepData[2, 3]
[1] 3.6
# omitting row value implies all rows; here all rows in column 3
ChildSleepData[, 3]
[1] 2.4 3.6 3.2 2.2 3.8 2.2 3.0 3.0
# omitting column values implies all columns; here all columns in row 2
ChildSleepData[2, ]
  Child WakeUp AcadPerf
2      B          9    3.6
# can also use ranges - rows 2 and 3, columns 2 and 3
ChildSleepData[2:3, 2:3]
  WakeUp AcadPerf
2      9    3.6
3      9    3.2
```

Variable Indexing

- We can determine the variable names in the data set with the `names` function

```
#prints the variable names in the dataset
names(ChildSleepData)
[1] "Child"      "WakeUp"     "AcadPerf"
```

- We can also access variables directly by using their names, either with `object[, "variable"]` notation or `object$variable` notation

```
# get first 3 rows of variable female using two methods
ChildSleepData[1:3, "WakeUp"]
[1] 11  9  9
ChildSleepData$WakeUp[1:3]
[1] 11  9  9
```

Exploring the data set

- Using `dim()`, we get the number of observations (rows) and variables (columns) in the data set.

```
dim(ChildSleepData)
[1] 8 3
```

- `summary()` is a generic function to summarize many types of R objects, including data sets.
- When used on a data set, `summary()` returns distributional summaries of variables in the data set.

```
summary(ChildSleepData)
```

	Child	WakeUp	AcadPerf
A	:1	Min. : 7.00	Min. :2.200
B	:1	1st Qu.: 8.75	1st Qu.:2.350
C	:1	Median : 9.50	Median :3.000
D	:1	Mean : 9.50	Mean :2.925
E	:1	3rd Qu.:10.25	3rd Qu.:3.300
F	:1	Max. :12.00	Max. :3.800
(Other)	:2		

Preparing R Session

```
install.packages("psych")  
install.packages("GPArotation")  
install.packages("sirt")  
install.packages("ltm")  
install.packages("irtoys")  
install.packages("mirt")  
install.packages("latticeExtra")
```

```
library(psych)  
library(GPArotation)  
library(sirt)  
library(msm)  
library(ltm)  
library(irtoys)  
library(mirt)  
library(latticeExtra)
```

Data Entry and Recoding

Read in Raw Data

```
data<- read.delim("C:/Users/kpreston/Self-esteem  
Dichotomous.dat", header=FALSE)  
data[data==9] <-NA #Replace missing code 9 with NA  
nitems<-ncol(data)
```

#recode data

```
keys <- c(1,-1,1,1,-1,-1,1,-1,-1,1)  
data <-  
reverse.code(keys,data,mini=rep(0,10),maxi=rep(1,10))
```

Checking Response Frequencies

Check response frequencies

response.frequencies(data)

##		0	1	miss
##	V1	0.7573696	0.24263039	0.0000000000
##	V2-	0.5124717	0.48752834	0.0000000000
##	V3	0.9183673	0.08163265	0.0000000000
##	V4	0.9138322	0.08616780	0.0000000000
##	V5-	0.7977273	0.20227273	0.002267574
##	V6-	0.5419501	0.45804989	0.0000000000
##	V7	0.8820862	0.11791383	0.0000000000
##	V8-	0.4454545	0.55454545	0.002267574
##	V9-	0.8707483	0.12925170	0.0000000000
##	V10	0.8049887	0.19501134	0.0000000000

Assumption Checking: Unidimensionality

1. Estimate bifactor model from poly/tetrachoric correlations

```
(rfac <- tetrachoric(data,na.rm=TRUE))
```

```
## Call: tetrachoric(x = data, na.rm = TRUE)
## tetrachoric correlation
##      V1    V2-   V3    V4    V5-   V6-   V7    V8-   V9-   V10
## V1    1.00
## V2-   0.54 1.00
## V3    0.67 0.63 1.00
## V4    0.57 0.46 0.77 1.00
## V5-   0.49 0.60 0.52 0.52 1.00
## V6-   0.44 0.80 0.65 0.38 0.62 1.00
## V7    0.50 0.48 0.76 0.68 0.50 0.46 1.00
## V8-   0.59 0.63 0.56 0.39 0.45 0.60 0.47 1.00
## V9-   0.47 0.58 0.39 0.41 0.62 0.61 0.49 0.65 1.00
## V10  0.81 0.59 0.76 0.62 0.56 0.61 0.70 0.66 0.56 1.00
##
## with tau of
##      V1    V2-   V3    V4    V5-   V6-   V7    V8-   V9-   V10
## 0.698  0.031  1.394  1.365  0.834  0.105  1.185 -0.137  1.130  0.860
```


Assumption Checking: Unidimensionality

2. Apply a Schmid-Leiman (SL) transformation to the factor loadings to obtain a semi-restricted exploratory hierarchical solution

```
rfac <- tetrachoric(data,na.rm=TRUE)$rho
(g3 <- schmid(rfac, nfactors = 3, fm = "minres",digits=3,rotate="oblimin"))

## Schmid-Leiman analysis
## Call: schmid(model = rfac, nfactors = 3, fm = "minres", digits = 3,
##      rotate = "oblimin")
## Schmid Leiman Factor loadings greater than 0.2
##      g      F1*    F2*    F3*    h2    u2    p2
## V1  0.75          0.36 0.71 0.29 0.80
## V2- 0.67 0.52          0.73 0.27 0.62
## V3  0.77          0.52    0.88 0.12 0.68
## V4  0.66          0.53    0.72 0.28 0.61
## V5- 0.61 0.36          0.52 0.48 0.71
## V6- 0.65 0.65          0.84 0.16 0.50
## V7  0.67          0.43    0.64 0.36 0.71
## V8- 0.69 0.28          0.28 0.64 0.36 0.74
## V9- 0.62 0.39          0.58 0.42 0.67
## V10 0.85          0.34 0.87 0.13 0.83
```

Assumption Checking:

Unidimensionality

3. ECV_2 - Ratio of variance explained by the general factor to variance explained by general and group factors

$$ECV_2 = \frac{\sum(\lambda_G)^2}{\sum(\lambda_G)^2 + \sum(\lambda_{g1})^2 + \sum(\lambda_{g2})^2 + \sum(\lambda_{g3})^2}$$

```
(sumout <- colSums(g3$sl[,1:(ncol(g3$sl)-3)])^2)
##           g           F1*           F2*           F3*
## 48.265261  5.437458  3.444150  1.619778
(ECV2 <- sumout[1] / sum(sumout))
##           g
## 0.8213036
```

- 82.1% of the variance is explained by the general factor

Rasch Model

$$P(\theta) = \frac{1}{1 + \exp[-(\theta - b_i)]}$$

Rasch Model

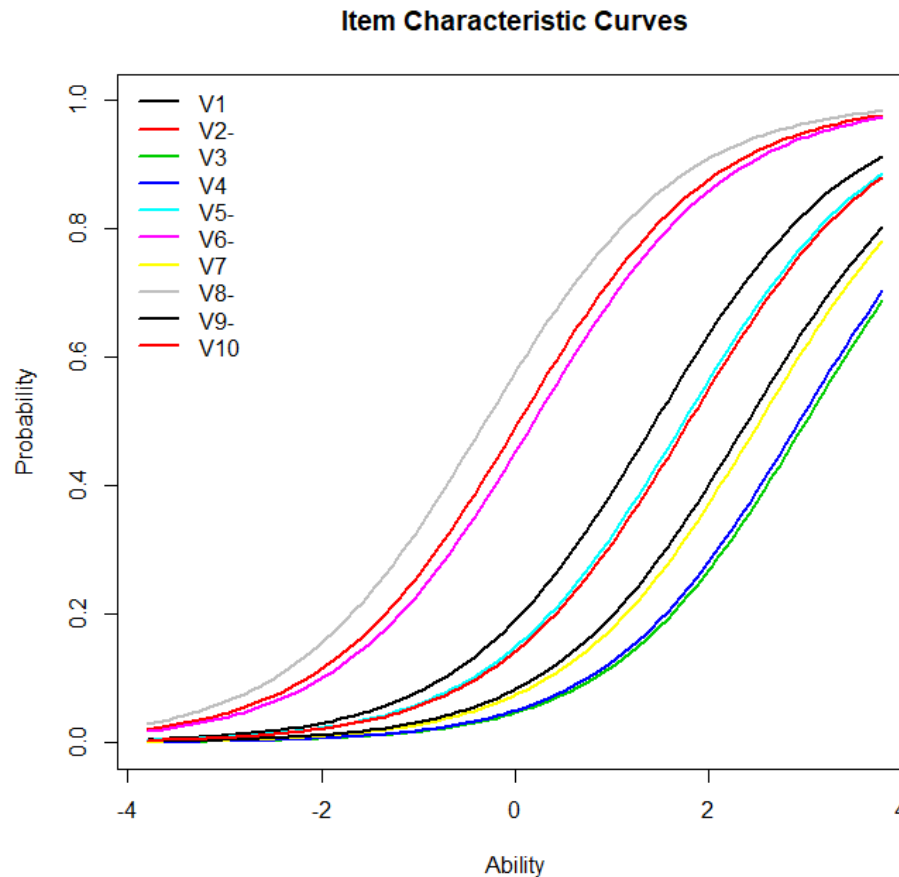
```
modRasch <- rasch(data, IRT.param =  
TRUE, constraint=cbind(ncol(data)+1,1))
```

```
round(coef(modRasch),2) # Obtain difficulty and discrimi  
nation parameter estimates
```

##	Dffclt	Dscrmn
## V1	1.45	1
## V2-	0.04	1
## V3	3.01	1
## V4	2.94	1
## V5-	1.74	1
## V6-	0.19	1
## V7	2.53	1
## V8-	-0.31	1
## V9-	2.40	1
## V10	1.80	1

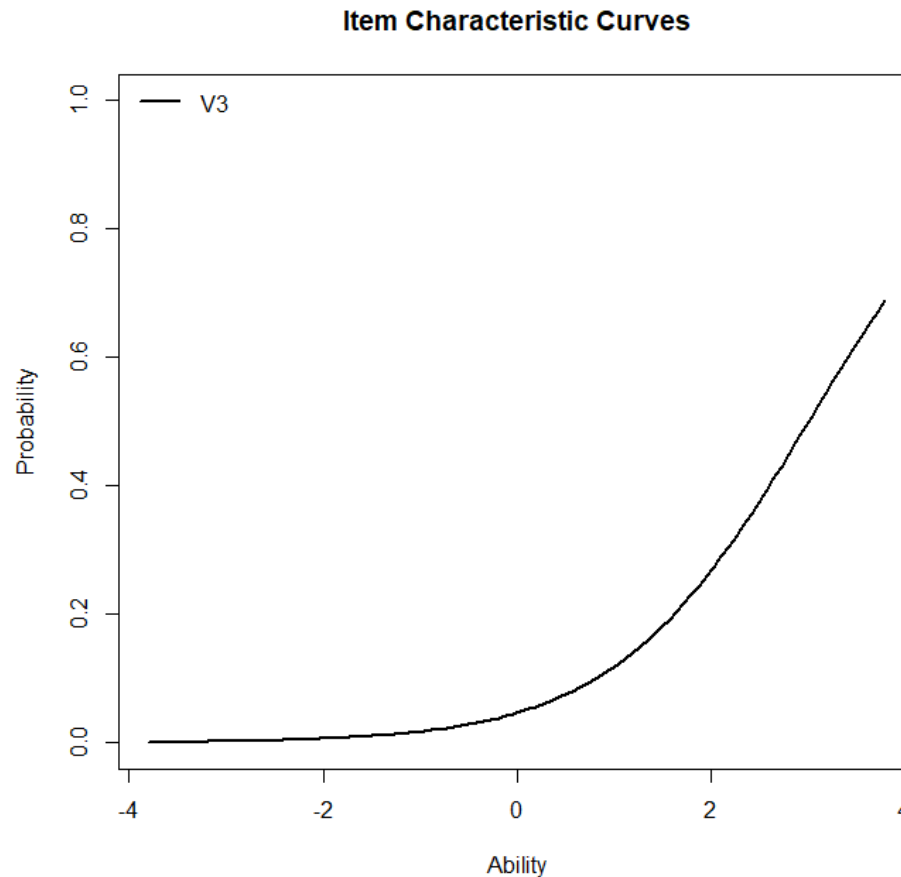
Rasch Model

```
plot(modRasch, type = "ICC", lwd = 2, legend=TRUE)  
#Item Characteristic Curves
```



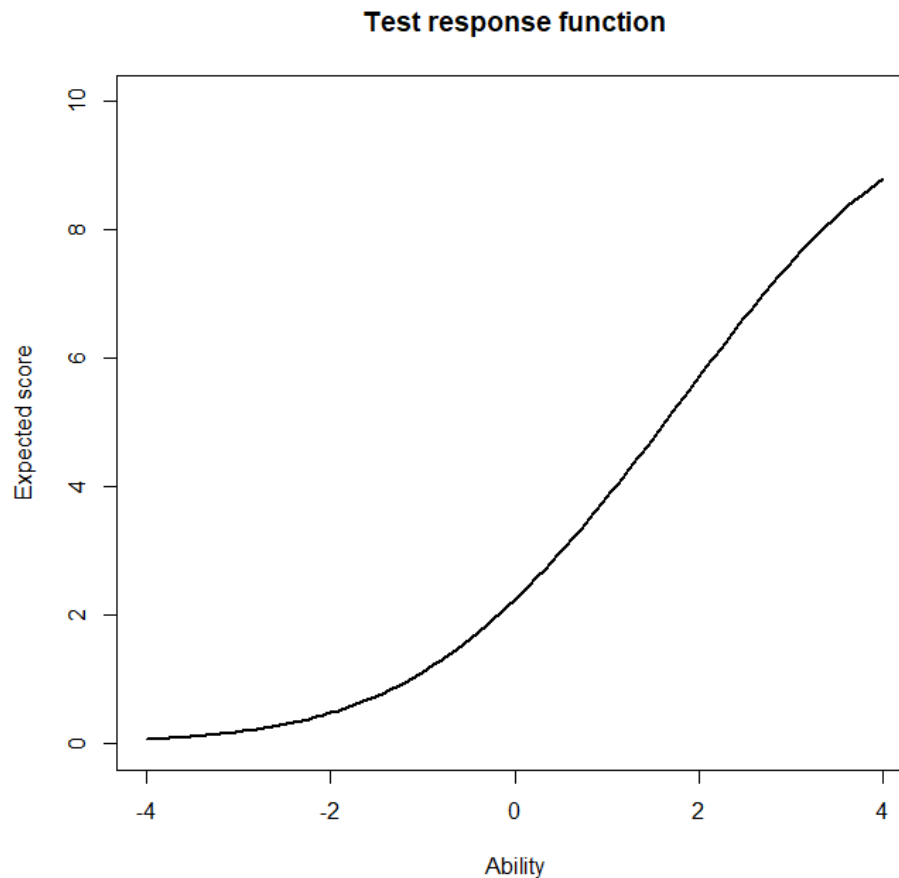
Rasch Model

```
plot(modRasch, type = "ICC", lwd = 2, legend=TRUE,  
item=3) #Only Item 3 ICC
```



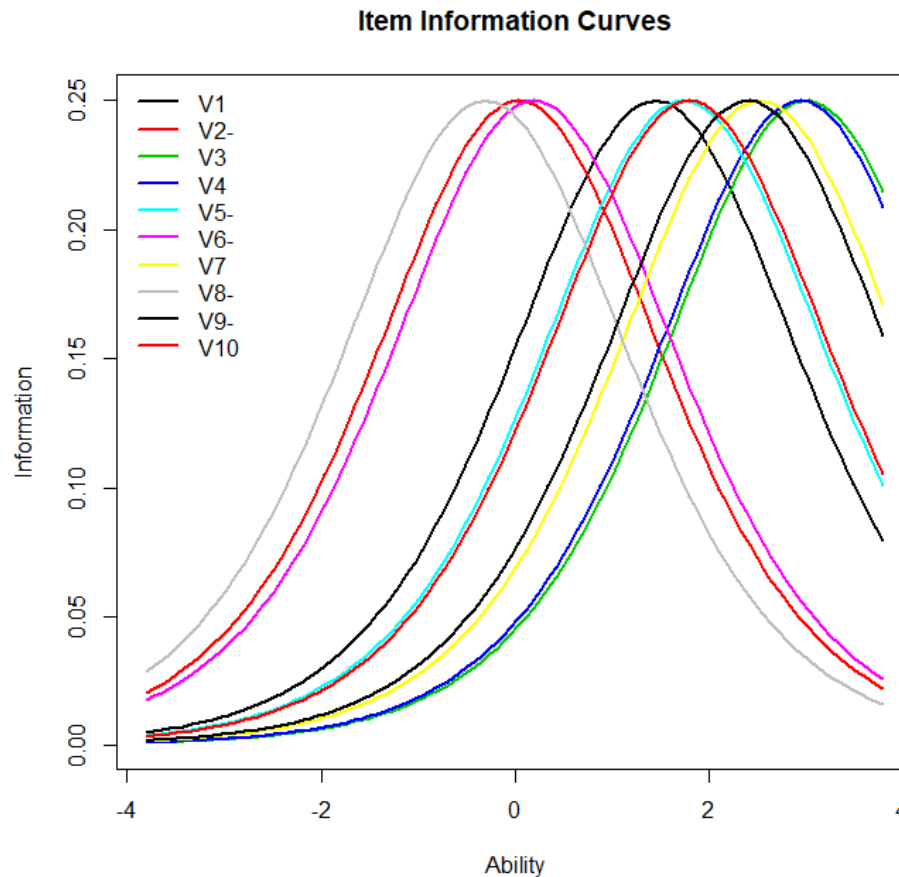
Rasch Model

```
plot(trf(est(data, model = "1PL", rasch="T", engine  
= "ltm"))) #Test Response Curve
```



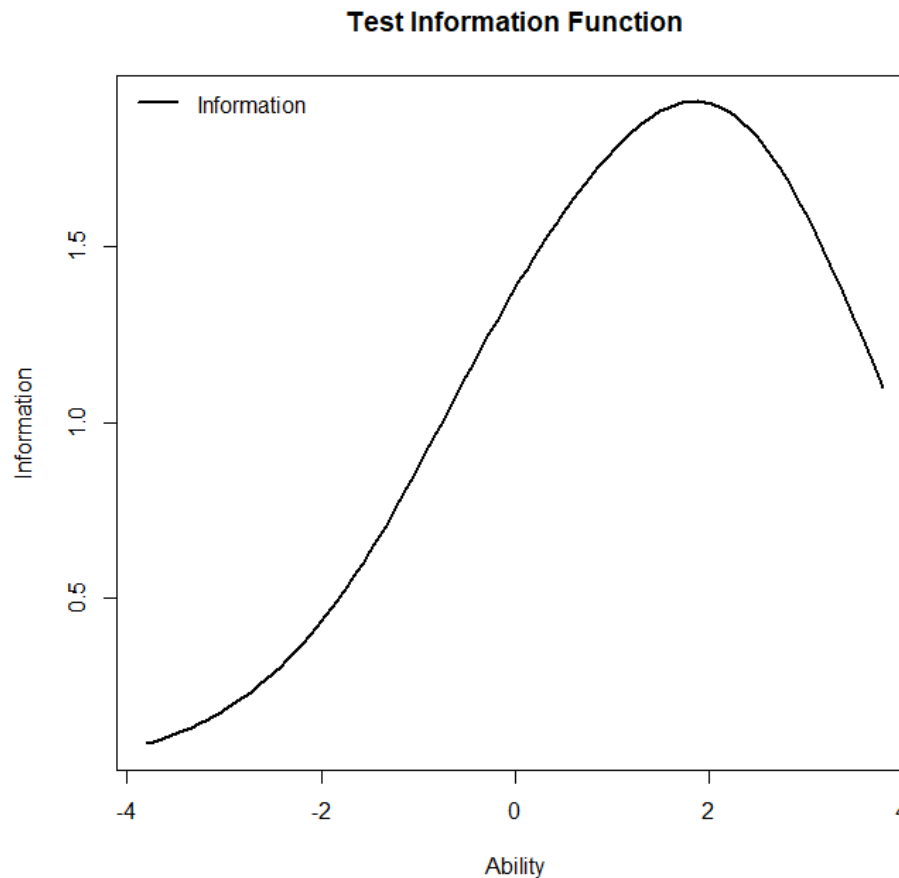
Rasch Model

`plot(modRasch, type = "IIC", lwd = 2, legend = TRUE)`
E) #Item Information Curve



Rasch Model

```
plot(modRasch,type = "IIC", lwd = 2,legend =TRUE,  
item=0) #Test Information Curve
```



Rasch Model

```
summary(modRasch)
```

```
## Model Summary:
```

```
##      log.Lik      AIC      BIC
## -1850.108 3720.216 3761.106
```

```
##
```

```
## Coefficients:
```

```
##           value std.err  z.vals
## Dffclt.V1    1.4478  0.1348 10.7407
## Dffclt.V2-    0.0404  0.1188  0.3402
## Dffclt.V3     3.0063  0.1951 15.4113
## Dffclt.V4     2.9379  0.1910 15.3784
## Dffclt.V5-    1.7446  0.1422 12.2693
## Dffclt.V6-    0.1942  0.1192  1.6287
## Dffclt.V7     2.5267  0.1700 14.8651
## Dffclt.V8-   -0.3053  0.1191 -2.5629
## Dffclt.V9-    2.4010  0.1645 14.5949
## Dffclt.V10    1.7999  0.1436 12.5328
## Dscrmn       1.0000      NA      NA
```

Rasch Model

```
thetaRasch <- ability(data, est(data, model = "1PL"  
, rasch = "T", engine = "ltm"), method = "BME")
```

```
round(head(thetaRasch), 2)
```

```
##           est    sem    n  
## [1, ]    0.30  0.63  10  
## [2, ]   -0.56  0.69  10  
## [3, ]   -0.56  0.69  10  
## [4, ]    2.45  0.59  10  
## [5, ]   -1.07  0.74  10  
## [6, ]    1.06  0.60  10
```

Rasch Model

```
item.fit(modRasch)
```

```
##
```

```
## Item-Fit Statistics and P-values
```

```
##
```

```
## Alternative: Items do not fit the model
```

```
## Ability Categories: 10
```

```
##
```

```
##           X^2 Pr(>X^2)
```

```
## V1    31.7051 <0.0001
```

```
## V2-   68.2809 <0.0001
```

```
## V3    40.0526 <0.0001
```

```
## V4    23.1139  0.0016
```

```
## V5-   31.0865  0.0001
```

```
## V6-   59.1941 <0.0001
```

```
## V7    27.8279  0.0002
```

```
## V8-   27.1124  0.0003
```

```
## V9-   24.8152  0.0008
```

```
## V10   57.9610 <0.0001
```

1PL

$$P(\theta) = \frac{1}{1 + \exp[-a(\theta - b_i)]}$$

1PL

```
mod1PL <- rasch(data, IRT.param = TRUE)
```

```
round(coef(mod1PL),2) # Obtain difficulty and discrimination parameter estimates
```

##		Dffclt	Dscrmn
##	V1	0.90	2.14
##	V2-	0.05	2.14
##	V3	1.81	2.14
##	V4	1.77	2.14
##	V5-	1.08	2.14
##	V6-	0.14	2.14
##	V7	1.53	2.14
##	V8-	-0.17	2.14
##	V9-	1.46	2.14
##	V10	1.11	2.14

1PL

```
summary(mod1PL)
```

```
## Model Summary:
```

```
##      log.Lik      AIC      BIC  
## -1762.771 3547.542 3592.522
```

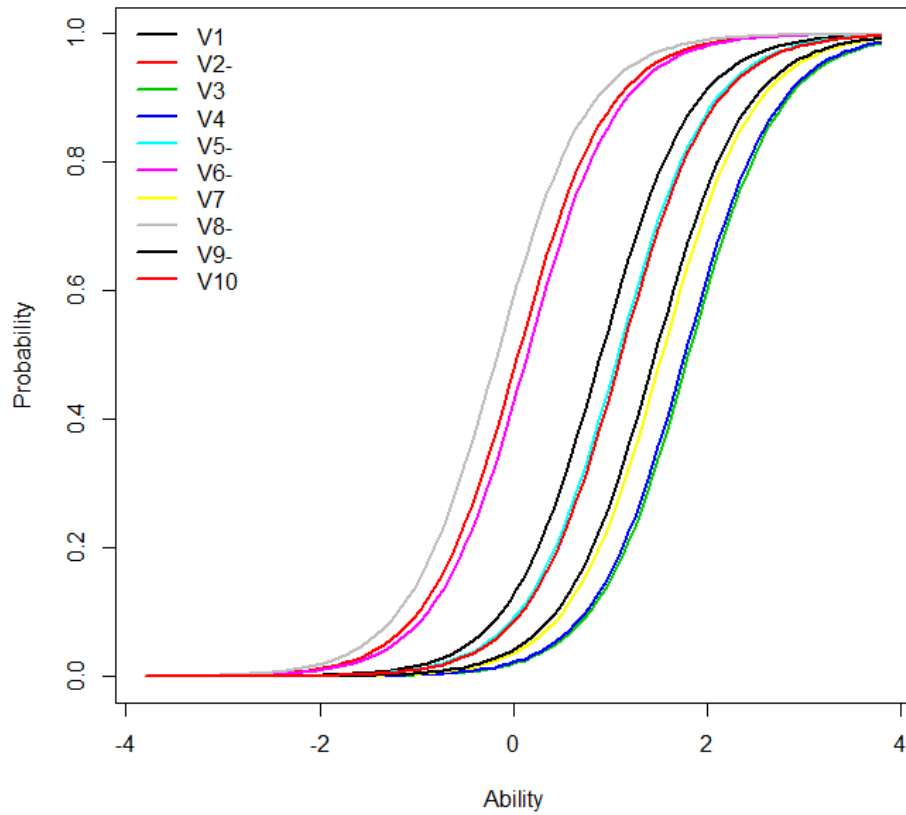
```
##
```

```
## Coefficients:
```

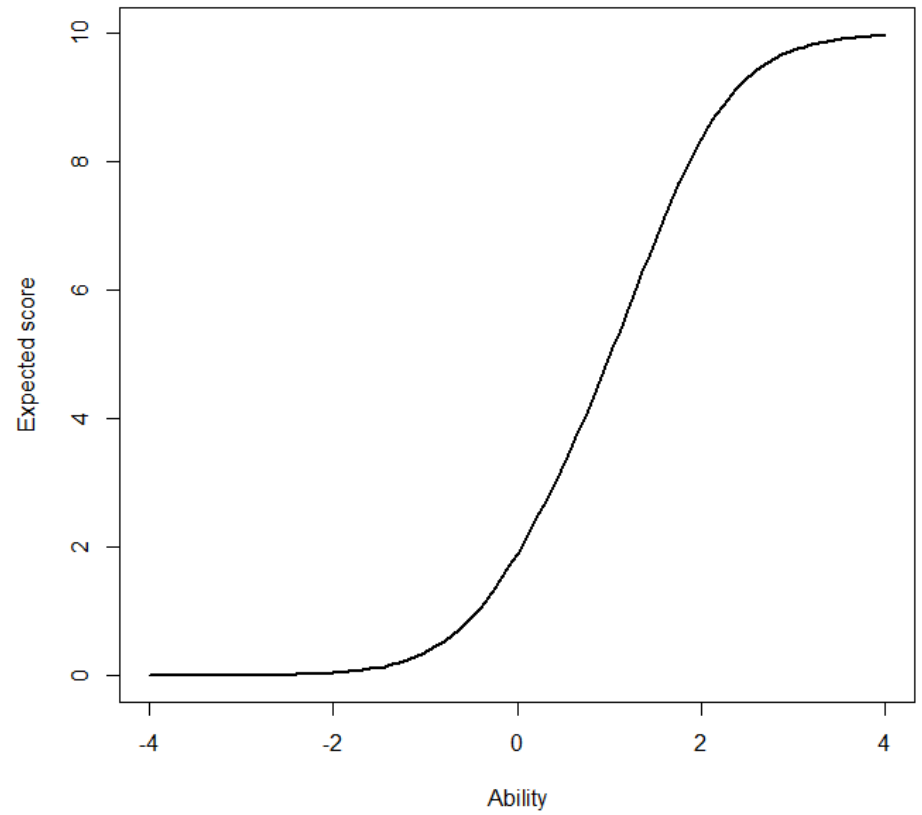
```
##      value std.err  z.vals  
## Dffclt.V1  0.9008  0.0899 10.0225  
## Dffclt.V2-  0.0454  0.0772  0.5881  
## Dffclt.V3   1.8090  0.1323 13.6714  
## Dffclt.V4   1.7700  0.1298 13.6321  
## Dffclt.V5-  1.0771  0.0958 11.2414  
## Dffclt.V6-  0.1406  0.0773  1.8183  
## Dffclt.V7   1.5336  0.1162 13.1996  
## Dffclt.V8- -0.1692  0.0779 -2.1726  
## Dffclt.V9-  1.4608  0.1124 12.9933  
## Dffclt.V10  1.1093  0.0970 11.4374  
## Dscrmn     2.1385  0.1212 17.6424
```

1PL

Item Characteristic Curves

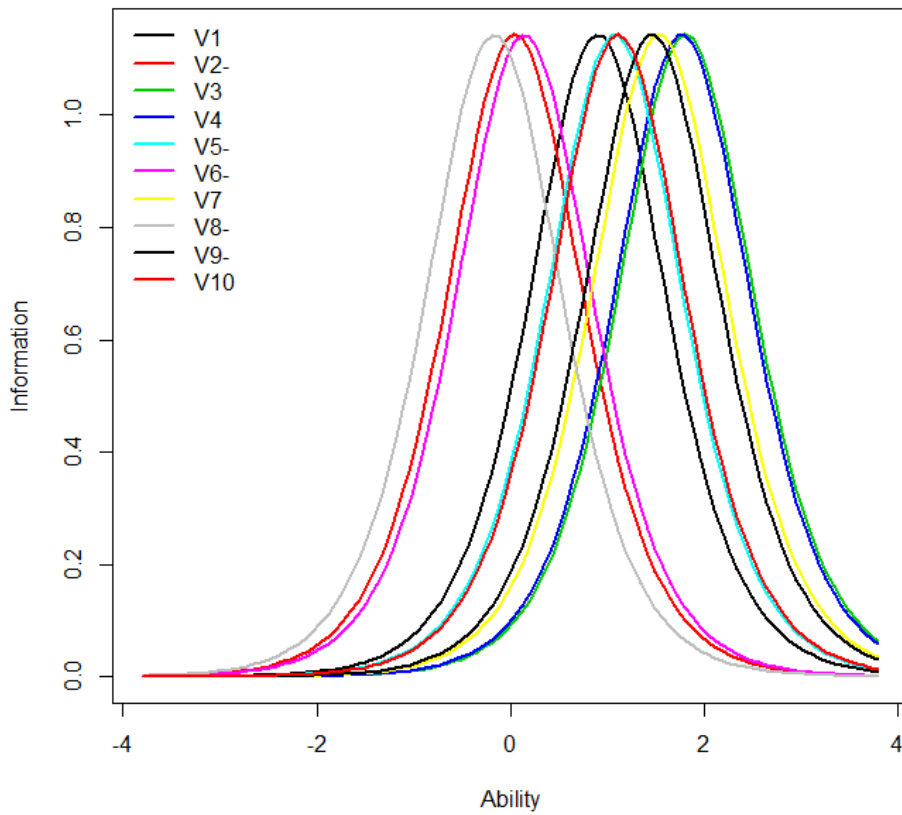


Test response function

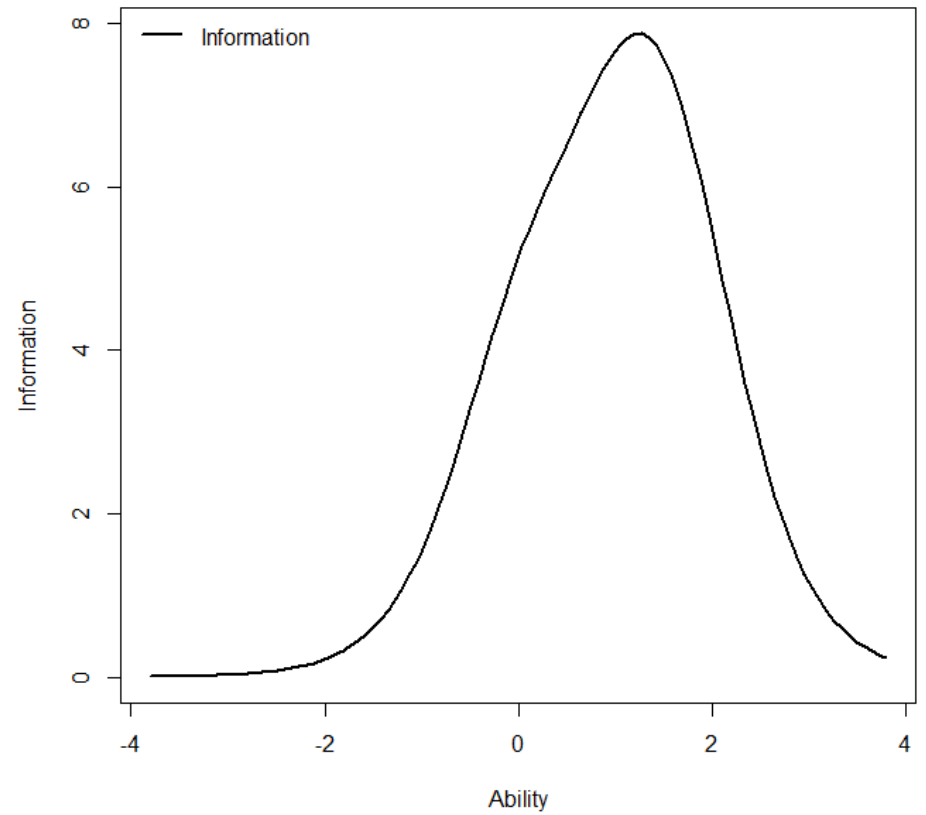


1PL

Item Information Curves



Test Information Function



1PL

```
item.fit(mod1PL)
```

```
##
```

```
## Item-Fit Statistics and P-values
```

```
## Alternative: Items do not fit the model
```

```
## Ability Categories: 10
```

```
##
```

```
##           X^2 Pr(>X^2)
```

```
## V1      7.5789  0.3712
```

```
## V2-    28.5334  0.0002
```

```
## V3     18.5744  0.0096
```

```
## V4      8.8948  0.2603
```

```
## V5-    15.4554  0.0306
```

```
## V6-    20.0431  0.0055
```

```
## V7     11.1756  0.1311
```

```
## V8-    14.8750  0.0376
```

```
## V9-     9.2461  0.2355
```

```
## V10    23.6501  0.0013
```

Comparing Models

Compare Model fit

anova.rasch(modRasch,mod1PL)

##

Likelihood Ratio Table

##		AIC	BIC	log.Lik	LRT	df	p.value
##	modRasch	3720.22	3761.11	-1850.11			
##	mod1PL	3547.54	3592.52	-1762.77	174.67	1	<0.001

2PL

$$P(\theta) = \frac{1}{1 + \exp[-a_i(\theta - b_i)]}$$

2PL

Perform the 2PL analysis with ltm(), and save the results in "mod2PL"

```
mod2PL <- ltm(data ~ z1, IRT.param = TRUE)
```

round(coef(mod2PL),2) # Obtain difficulty and discrimination parameter estimates

##		Dffclt	Dscrmn
##	V1	0.92	2.04
##	V2-	0.05	2.55
##	V3	1.60	3.12
##	V4	1.87	1.87
##	V5-	1.20	1.66
##	V6-	0.14	2.27
##	V7	1.59	1.95
##	V8-	-0.17	1.98
##	V9-	1.62	1.70
##	V10	0.97	3.26

2PL

```
summary(mod2PL)
```

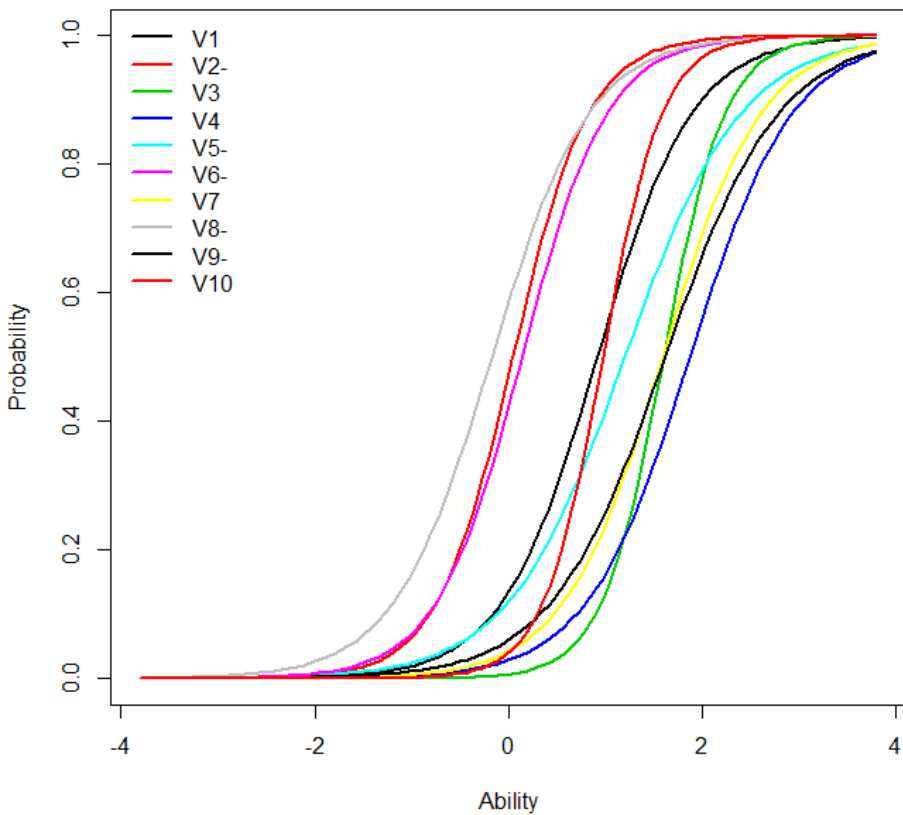
```
##  
## Model Summary:  
##      log.Lik      AIC      BIC  
## -1752.659 3545.319 3627.1
```

```
##  
## Coefficients:  
##           value std.err  z.vals  
## Dffclt.V1    0.9166  0.1021  8.9815  
## Dffclt.V2-   0.0457  0.0727  0.6281  
## Dffclt.V3    1.6011  0.1340 11.9451  
## Dffclt.V4    1.8733  0.1998  9.3748  
## Dffclt.V5-   1.2008  0.1346  8.9246  
## Dffclt.V6-   0.1392  0.0758  1.8353  
## Dffclt.V7    1.5907  0.1593  9.9862  
## Dffclt.V8-  -0.1740  0.0807 -2.1563  
## Dffclt.V9-   1.6169  0.1752  9.2270  
## Dffclt.V10   0.9742  0.0896 10.8790
```

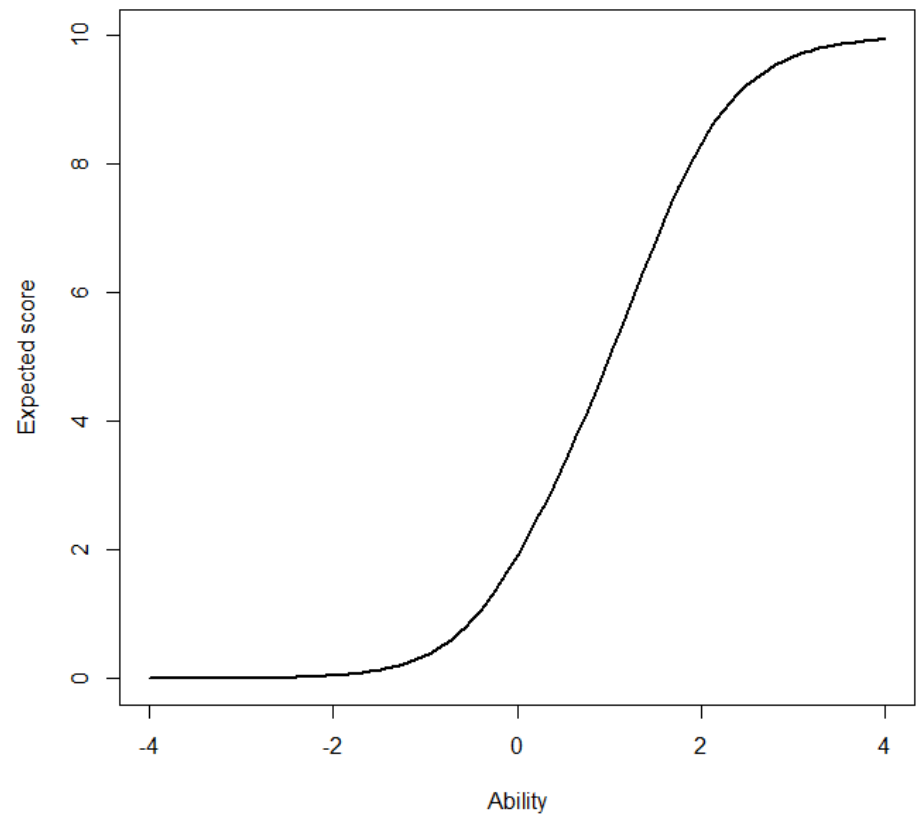
```
## Dscrmn.V1    2.0359  0.2949  6.9049  
## Dscrmn.V2-   2.5474  0.3565  7.1459  
## Dscrmn.V3    3.1185  0.6073  5.1354  
## Dscrmn.V4    1.8743  0.3265  5.7402  
## Dscrmn.V5-   1.6615  0.2396  6.9347  
## Dscrmn.V6-   2.2654  0.3124  7.2518  
## Dscrmn.V7    1.9536  0.3105  6.2920  
## Dscrmn.V8-   1.9781  0.2641  7.4904  
## Dscrmn.V9-   1.6982  0.2697  6.2960  
## Dscrmn.V10   3.2559  0.5954  5.4685
```

2PL

Item Characteristic Curves

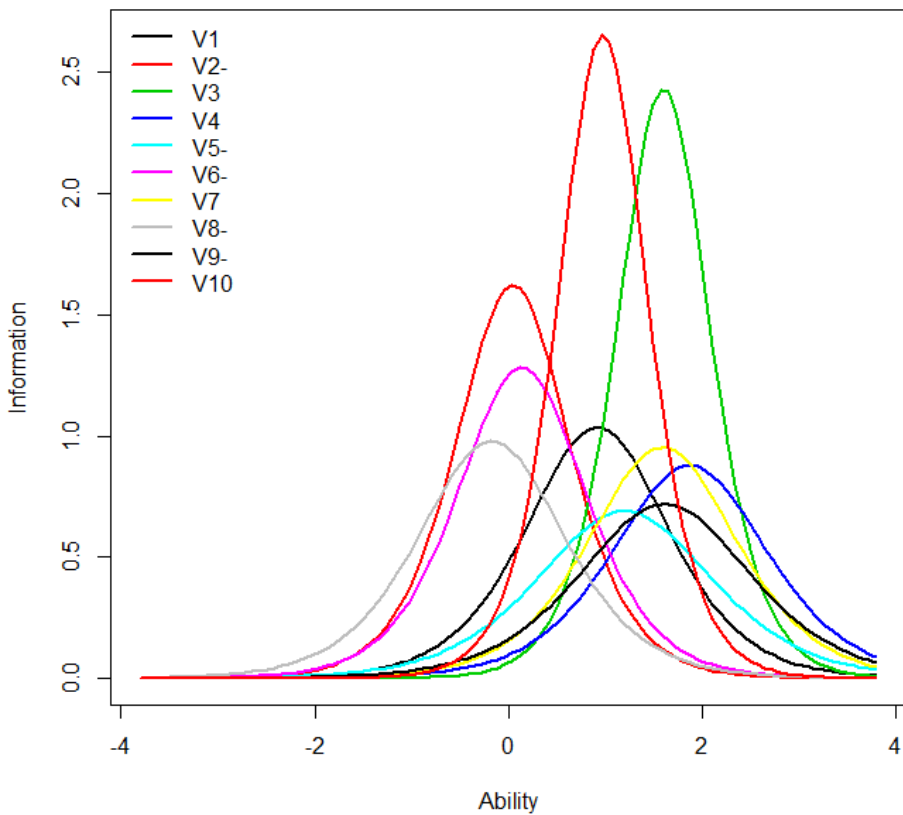


Test response function

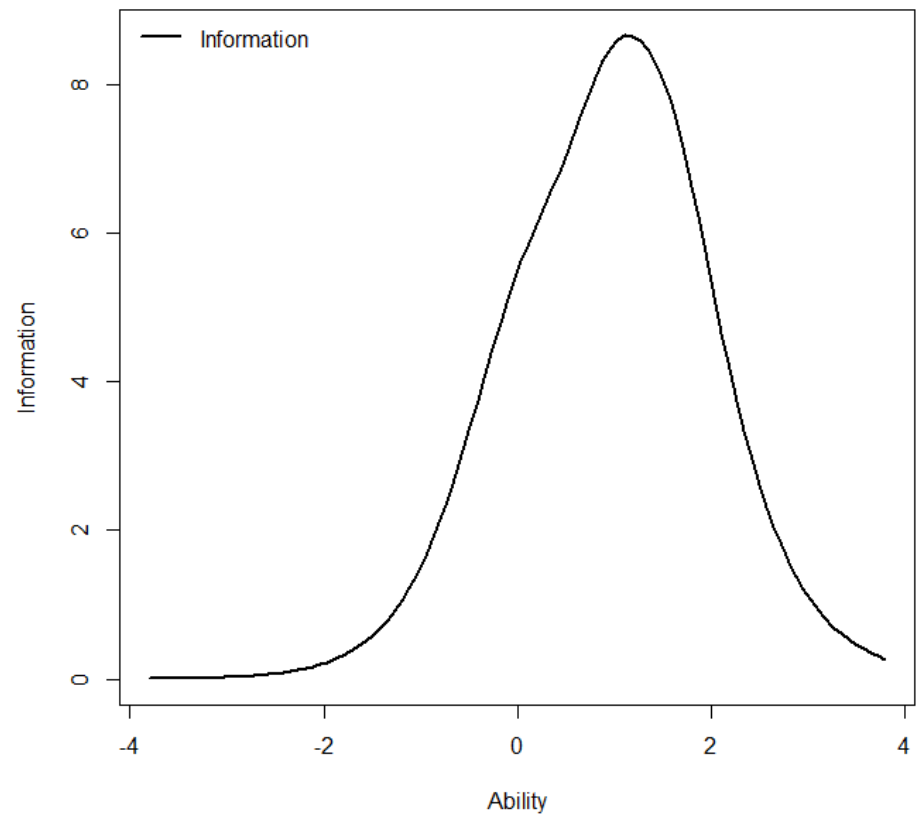


2PL

Item Information Curves



Test Information Function



2PL

```
item.fit(mod2PL)
```

```
##
```

```
## Item-Fit Statistics and P-values
```

```
##
```

```
## Alternative: Items do not fit the model
```

```
## Ability Categories: 10
```

```
##
```

```
##           X^2 Pr(>X^2)
```

```
## V1    14.7882    0.0634
```

```
## V2-   32.8034    0.0001
```

```
## V3    15.1024    0.0572
```

```
## V4    19.3237    0.0132
```

```
## V5-   22.9620    0.0034
```

```
## V6-    8.7313    0.3655
```

```
## V7    13.1027    0.1084
```

```
## V8-   12.4854    0.1308
```

```
## V9-   29.2589    0.0003
```

```
## V10   15.4248    0.0514
```

Comparing Models

Compare Model fit

anova.rasch(mod1PL,mod2PL)

##

Likelihood Ratio Table

##		AIC	BIC	log.Lik	LRT	df	p.value
----	--	-----	-----	---------	-----	----	---------

##	mod1PL	3547.54	3592.52	-1762.77			
----	--------	---------	---------	----------	--	--	--

##	mod2PL	3545.32	3627.10	-1752.66	20.22	9	0.017
----	--------	---------	---------	----------	-------	---	-------

Scoring

BME (Bayesian Modal Estimation)

Person 1: 1, 1, 0, 1, 0, 0, 0, 0, 0, 0

```
person1 <- data[1,]  
## Computing theta  
theta <- seq(-4,4,.001)  
apar <- coef(mod2PL)[,2]  
bpar <- coef(mod2PL)[,1]
```

```
P <- matrix(0,nitems,length(theta))  
L <- matrix(0,nitems,length(theta))  
dis <- dnorm(theta)  
dis <- dis/sum(dis)
```

```
for(m in 1:nitems){  
  P[m,] <- 1/(1+exp(-(apar[m]*(theta-bpar[m]))))}
```

$$L = \prod_{i=1}^{nitems} P_i(u_i|\theta)\phi(\theta)$$

$$P(\theta) = \frac{1}{1 + \exp[-a_i(\theta - b_i)]}$$

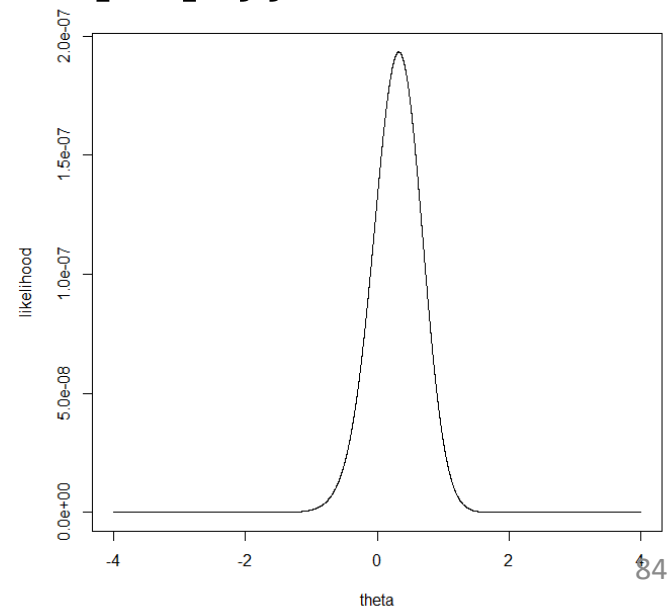
Scoring

BME (Bayesian Modal Estimation)

$$L = \prod_{i=1}^{nitems} P_i(u_i|\theta)\phi(\theta)$$

```
for(k in 1:length(person1)){
  if(person1[k] == 1){ L[k,] <- P[k,] }
  else if(person1[k] == 0) {L[k,] <- 1-P[k,] }}
```

```
likelihood <- dis
for(j in 1:length(person1)){
  likelihood <- likelihood*L[j,]}
plot(theta,likelihood,type="l")
(bme<-theta[which.max(likelihood)])
## [1] 0.325
```



Scoring

BME (Bayesian Modal Estimation)

```
for(k in 1:length(person1)){  
  if(person1[k] == 1){ L[k,] <- P[k,] }  
  else if(person1[k] == 0) {L[k,] <- 1-P[k,] }}
```

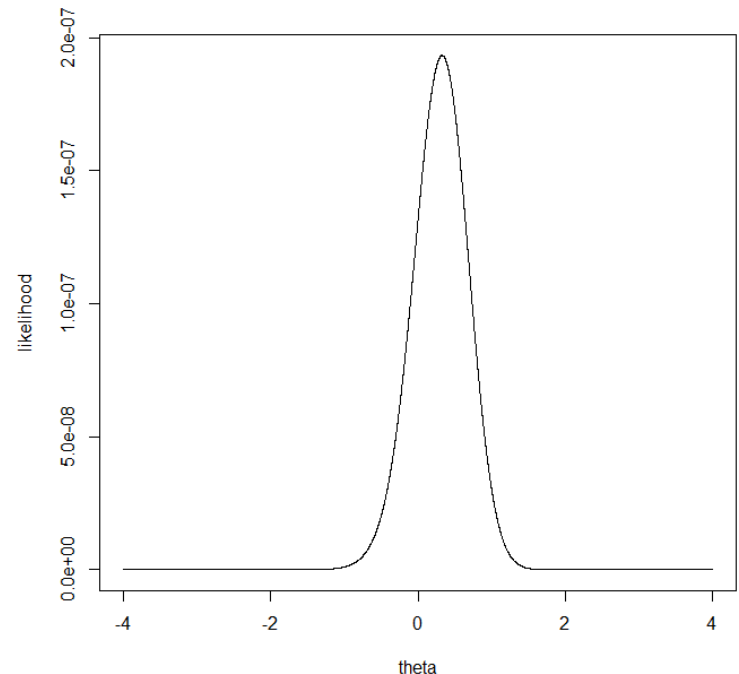
```
likelihood <- dis  
for(j in 1:length(person1)){  
  likelihood <- likelihood*L[j,]}
```

```
plot(theta,likelihood,type="l")  
(bme<-theta[which.max(likelihood)])
```

```
## [1] 0.325
```

```
round(theta2PL[1,],2)
```

```
##      est      sem      n  
## 0.32  0.36 10.00
```



3PL

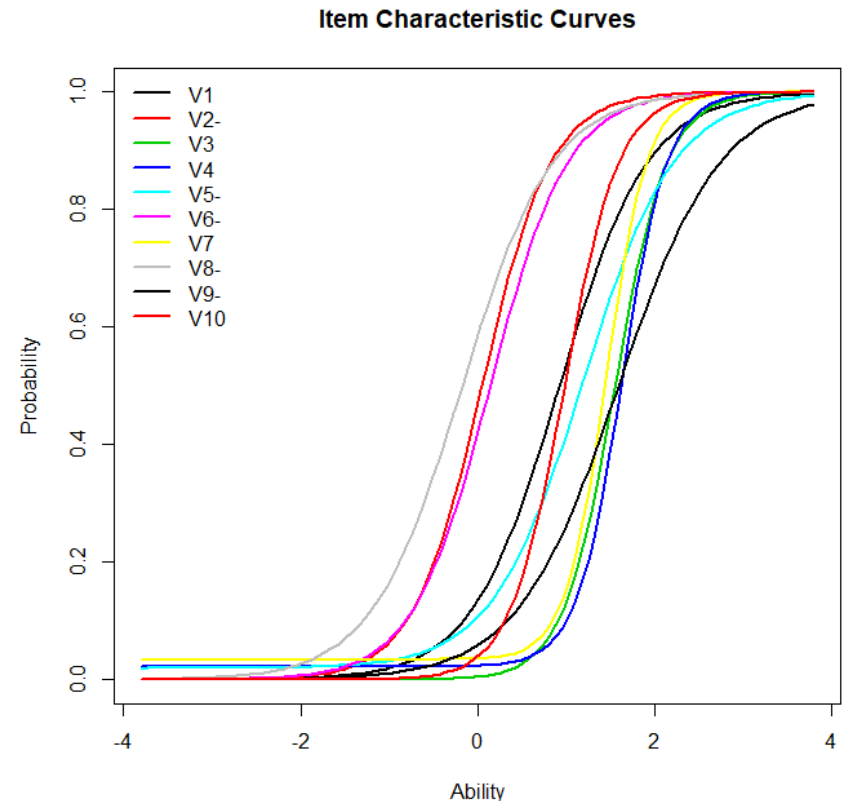
$$P(\theta) = c_i + \frac{1 - c_i}{1 + \exp[-a_i(\theta - b_i)]}$$

3PL

```
mod3PL <- tpm(data, type="latent.trait", IRT.param = TRUE,  
control=c(optimizer="nlminb"))
```

```
round(coef(mod3PL),2) # Obtain difficulty and discrimina  
tion parameter estimates
```

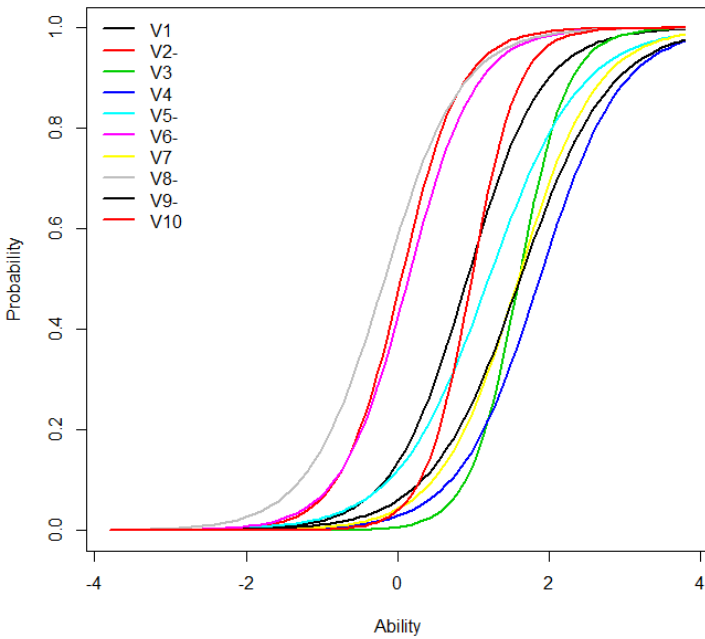
##		Gussng	Dffclt	Dscrmn
##	V1	0.00	0.92	2.01
##	V2-	0.00	0.05	2.56
##	V3	0.00	1.55	3.38
##	V4	0.02	1.63	3.88
##	V5-	0.02	1.20	1.94
##	V6-	0.00	0.14	2.29
##	V7	0.03	1.46	4.30
##	V8-	0.00	-0.17	1.95
##	V9-	0.00	1.59	1.74
##	V10	0.00	0.98	3.25



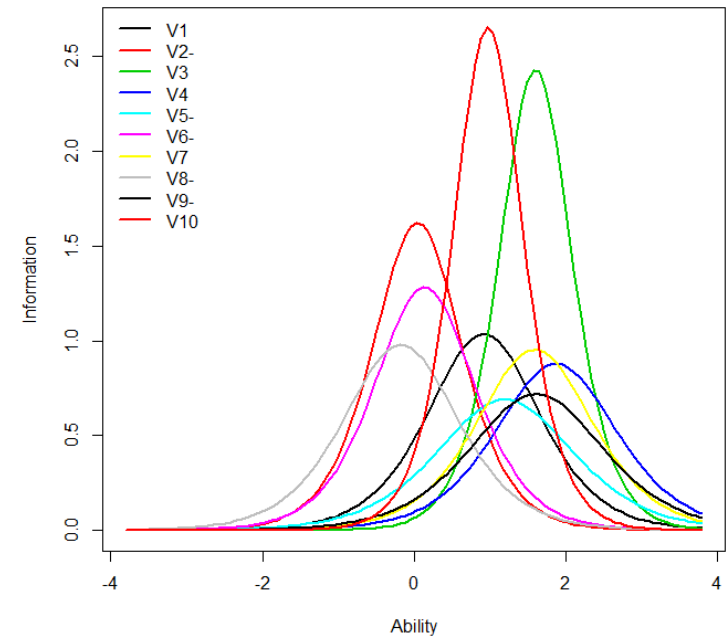
Best Model

2PL

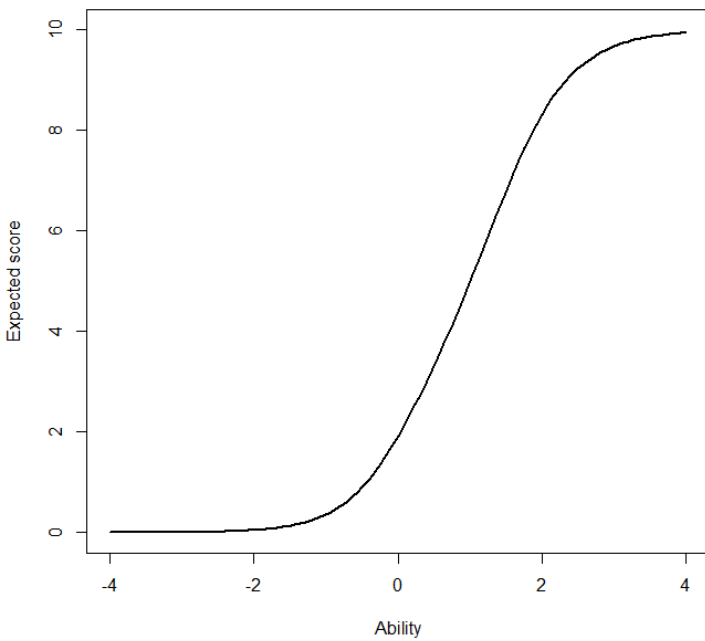
Item Characteristic Curves



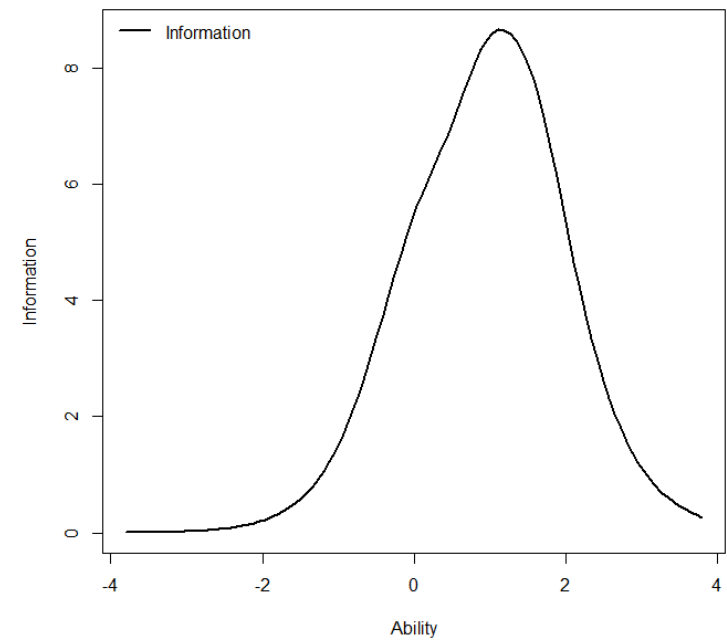
Item Information Curves



Test response function



Test Information Function



2PL

Confirm Unidimensionality: Modified Parallel Analysis

```
unidimTest(mod2PL)
```

```
##
```

```
## Unidimensionality Check using Modified Parallel Analysis
```

```
##
```

```
##
```

```
## Alternative hypothesis: the second eigenvalue of the observed  
## data is substantially larger than the second eigenvalue of  
## data under the assumed IRT model
```

```
##
```

```
## Second eigenvalue in the observed data: 0.8961
```

```
## Average of second eigenvalues in Monte Carlo samples: 0.6274
```

```
## Monte Carlo samples: 100
```

```
## p-value: 0.0594
```

Assessing Local Independence

```
## Assessing Local Dependence using Yen's Q3 statistic, (Yen, 1984)
mod.q3 <- Q3( dat=data, theta=theta2PL[,1] , b=coef(mod2PL)[,1])

## Yen's Q3 Statistic based on an estimated theta score
## *** 10 Items | 45 item pairs
## *** Q3 Descriptives
##      M      SD   Min   10%   25%   50%   75%   90%   Max
## -0.014  0.114 -0.186 -0.131 -0.091 -0.029  0.031  0.138  0.301
```

Yen's Q3 Correlations

```
mod.q3$q3.matrix[upper.tri(mod.q3$q3.matrix,diag=TRUE)] <- 0
round(mod.q3$q3.matrix,2)
```

```
##           V1    V2-    V3    V4    V5-    V6-    V7    V8-    V9- V10
## V1
## V2- -0.06
## V3  0.01 -0.15
## V4  0.01 -0.13  0.22
## V5- -0.03  0.02 -0.09  0.00
## V6- -0.15 0.30 -0.11 -0.17  0.06
## V7  -0.07 -0.12  0.18  0.15 -0.03 -0.13
## V8-  0.01  0.06 -0.13 -0.13 -0.08  0.04 -0.09
## V9- -0.06 -0.03 -0.19 -0.08  0.12  0.01 -0.03  0.03
## V10 0.29 -0.11  0.06  0.00 -0.03 -0.07  0.10 -0.01 -0.03
```

#Criteria relative to average observed residual correlation from Christensen, Maransky & Horton (2017)

```
mod.q3$Q3.stat
```

```
##           M           SD           Min           10%           25%           50%
## -0.01400869  0.11426811 -0.18632406 -0.13112532 -0.09130349 -0.02858531
##           75%           90%           Max
##  0.03106187  0.13792324  0.30136573
```

```
mod.q3$Q3.stat[9]-abs(mod.q3$Q3.stat[1])
## 0.287357
```

Rosenberg Self-Esteem Scale (1965)

1. On the whole, I am satisfied with myself.
2. At times, I think I am not good at all (R)
3. I feel that I have a number of good qualities
4. I am able to do things as well as most other people.
5. I feel I do not have much to be proud of. (R)
6. I certainly feel useless at times. (R)
7. I feel that I'm a person of worth, at least on an equal plane with others.
8. I wish I could have more respect for myself (R)
9. All in all, I am inclined to feel that I am a failure (R)
10. I take a positive attitude toward myself.

Next Steps

- Use θ estimates in future analyses instead of summed scores

```
SelfEsteem <- theta2PL[1,]
```

- Investigate item *and* category functioning using polytomous IRT models

Conclusion

- IRT has clear benefits compared to CTT
- Analysis of scales using IRT models are becoming more accessible and user-friendly
- Improved precision of person estimates for use in future analyses