

# Package ‘test2norm’

May 8, 2026

**Type** Package

**Title** Normative Standards for Cognitive Tests

**Version** 0.3.0.1

**Description** Package test2norm contains functions to generate formulas for normative standards applied to cognitive tests. It takes raw test scores (e.g., number of correct responses) and converts them to scaled scores and demographically adjusted scores, using methods described in Heaton et al. (2003) <doi:10.1016/B978-012703570-3/50010-9> & Heaton et al. (2009, ISBN:9780199702800). The scaled scores are calculated as quantiles of the raw test scores, scaled to have the mean of 10 and standard deviation of 3, such that higher values always correspond to better performance on the test. The demographically adjusted scores are calculated from the residuals of a model that regresses scaled scores on demographic predictors (e.g., age). The norming procedure makes use of the mfp2() function from the 'mfp2' package to explore nonlinear associations between cognition and demographic variables.

**License** GPL (>= 2)

**Encoding** UTF-8

**LazyData** true

**Imports** mfp2

**Depends** R (>= 3.5.0)

**RoxygenNote** 7.2.3

**NeedsCompilation** no

**Repository** CRAN

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**Author** Anya Umlauf [aut, cre]

**Maintainer** Anya Umlauf <aumlau@health.ucsd.edu>

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PsychTestData	<i>Neuropsychological test data</i>
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### Description

A simulated data containing raw test scores and demographic characteristics for 250 persons, 200 in the control group and 50 in the test group. The raw test scores are to be converted to demographically corrected normed scores, adjusting for effects of age and sex. The control group is used to generate the norming formulas, which are then applied to all scores.

### Usage

```
PsychTestData
```

### Format

A data frame with 250 rows and 4 variables:

**rawscore** raw test score on a neuropsychological test, ranging 0-36, with higher values indicating better test performance

**age** age of the participant, in years

**male** sex of the participant, male (1) or female (0)

**group** norming group the participant belongs to (control or test)

### Examples

```
data(PsychTestData)
test2norm(data = PsychTestData, test = "rawscore",
           test.min = 0, test.max = 36, test.better = "High",
           group.id = "group", control.id = "control",
           demographics = c("age", "male"))
```

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raw2scaled	<i>Convert raw neuropsychological test scores to scaled scores.</i>
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## Description

Convert raw neuropsychological test scores to scaled scores.

## Usage

```
raw2scaled(
  data = NULL,
  test = NULL,
  test.min = NULL,
  test.max = NULL,
  test.better = c("High", "Low"),
  group.id = NULL,
  control.id = NULL,
  all.controls = FALSE,
  rnd.s = TRUE
)
```

## Arguments

<code>data</code>	a data frame containing the test score
<code>test</code>	a character string specifying the name of the variable containing the test score
<code>test.min</code>	a real number indicating the smallest possible test score
<code>test.max</code>	a real number indicating the largest possible test score
<code>test.better</code>	a character string indicating direction of the scores. Use "High" if high test scores imply better performance, use "Low" otherwise.
<code>group.id</code>	a character string specifying the name of the variable containing group identification (i.e. control vs exposed/test/risk). Ignored, if <code>all.controls = TRUE</code> .
<code>control.id</code>	a character string specifying the label of the control group within <code>group.id</code> variable. Ignored, if <code>all.controls = TRUE</code> .
<code>all.controls</code>	a logical indicating whether all observations should be treated as controls. Overwrites <code>group.id</code> and <code>control.id</code> .
<code>rnd.s</code>	a logical indicating whether the scaled scores should be rounded. Default is <code>TRUE</code> .

## Details

The `raw2scaled()` function can be used by neuropsychologists, who wish to convert raw test scores to scaled scores (mean=10, SD=3), using methods described in Heaton et al. (2003 & 2009). The raw test scores that have many decimal digits should be rounded to fewer digits prior to the application of the `raw2scaled()` function. This will significantly reduce software running time. The

recommended number of decimal digits is 4 or fewer. Values below `test.min` or above `test.max` will result in NA. Detailed description of the procedure are found in Umlauf et al. (2024).

Note that the function does not guarantee that the calculated scaled scores range from 0 to 20, because it uses normal distribution quantiles. Therefore, scaled scores outside 0-20 range are possible.

### Value

A list consisting of 3 objects. The first two are vectors containing the original raw test scores and the calculated scaled scores. The third object in the list, called `SS.maps`, contains conversions from raw scores to scaled scores in a form of a table with two columns, one representing scaled scores (one per row) and one representing raw scores (a single value or range of raw values corresponding to each scaled score). Note that this table shows rounded scaled scores regardless of the value for `rnd.s`.

### Author(s)

Anya Umlauf

### References

Umlauf A et al. (2024) Automated procedure for demographic adjustments on cognitive test scores. <doi:10.1080/23279095.2023.2288231>

Heaton RK, Taylor MJ, & Manly J (2003) Demographic effects and use of demographically corrected norms with the WAIS-III and WMS-III. In: Tulskey D et al. (Eds.) *Clinical Interpretation of the WAIS-III and WMS-III*. San Diego, CA: Academic Press, 183-210.

Heaton RK, Ryan L, & Grant I (2009) Demographic influences and use of demographically corrected norms in neuropsychological assessment. In Grant I & Adams KM (Eds.) *Neuropsychological Assessment of Neuropsychiatric and Neuromedical Disorders*. New York, NY: Oxford University Press, 127-155.

### Examples

```
data(PsychTestData)
raw2scaled(data = PsychTestData, test = "rawscore",
            test.min = 0, test.max = 36, test.better = "High",
            group.id = "group", control.id = "control")
```

---

score2adjust

*Convert neuropsychological test scores to demographically adjusted norms.*

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### Description

Convert neuropsychological test scores to demographically adjusted norms.

**Usage**

```
score2adjust(
  data = NULL,
  test.score = NULL,
  group.id = NULL,
  control.id = NULL,
  all.controls = FALSE,
  demographics = NULL,
  mfp.alpha = 1,
  rnd.a = TRUE,
  mean.a = 50,
  sd.a = 10
)
```

**Arguments**

<code>data</code>	a data frame containing the variables needed for the norming process. The current version of the function does not accommodate missing data. For best results, exclude cases with missing test scores or missing demographics before applying this function.
<code>test.score</code>	a character string specifying the name of the test to be normed, usually the output of the <code>raw2scaled()</code> function.
<code>group.id</code>	a character string specifying the name of the variable containing group identification (i.e. control vs exposed/test/risk). Ignored, if <code>all.controls = TRUE</code> .
<code>control.id</code>	a character string specifying the label of the control group within <code>group.id</code> variable. Ignored, if <code>all.controls = TRUE</code> .
<code>all.controls</code>	a logical indicating whether all observations should be treated as controls. Overwrites <code>group.id</code> and <code>control.id</code> .
<code>demographics</code>	a single or multiple character strings (concatenated by <code>c()</code> function) specifying the names of demographic predictors to be included into normative formulas. Demographic variables should be numeric or binary (0/1).
<code>mfp.alpha</code>	a numeric value between 0 and 1 that sets significance level for inclusion of demographic predictors into normative formula. Passed to the <code>mfp2()</code> function (its <code>select</code> argument). Default value is 1 for inclusion of all predictors regardless of their significance.
<code>rnd.a</code>	a logical indicating whether the adjusted scores (T-scores) should be rounded. Default is <code>TRUE</code> .
<code>mean.a</code>	numeric value for the mean of adjusted score (T-score) distribution. Default is 50.
<code>sd.a</code>	numeric value for the standard deviation of adjusted score (T-score) distribution. Default is 10.

**Details**

The `score2adjust()` function can be used by neuropsychologists, who wish to construct normative formulas for cognitive tests that adjust for expected effects of demographic characteristics (e.g.,

age), using methods described in Heaton et al. (2003 & 2009). The adjusted scores are sometimes referred to as T-scores in the literature. The norming procedure makes use of the `mfp2()` function from the `mfp2` package to explore nonlinear associations between cognition and demographic variables. Detailed description of the procedure are found in Umlauf et al. (2024). (Previous versions of the function depended on `mfp` package.)

### Value

A list consisting of 3 objects. The first two are vectors containing the non-adjusted test scores and the calculated demographically adjusted scores. The last item in the output list is also a list called `MFP.formulas`. It contains the information for calculation of adjusted scores, including variable transformations (if any), multiple fractional polynomial (MFP) model coefficients, the standard deviation of residuals resulting from the MFP modeling, and a matrix with number of rows equal to the number of predictors and 2 columns containing powers (in numeric form) selected for variable transformations.

### Author(s)

Anya Umlauf

### References

Umlauf A et al. (2024) Automated procedure for demographic adjustments on cognitive test scores. <doi:10.1080/23279095.2023.2288231>

Heaton RK, Taylor MJ, & Manly J (2003) Demographic effects and use of demographically corrected norms with the WAIS-III and WMS-III. In: Tulskey D et al. (Eds.) *Clinical Interpretation of the WAIS-III and WMS-III*. San Diego, CA: Academic Press, 183-210.

Heaton RK, Ryan L, & Grant I (2009) Demographic influences and use of demographically corrected norms in neuropsychological assessment. In Grant I & Adams KM (Eds.) *Neuropsychological Assessment of Neuropsychiatric and Neuromedical Disorders*. New York, NY: Oxford University Press, 127-155.

Benner A (2005) `mfp`: Multivariable fractional polynomials. *R News* 5(2): 20–23.

### Examples

```
data(PsychTestData)
PsychTestData$scaledscore <- raw2scaled(data=PsychTestData, test="rawscore",
                                       test.min=0, test.max=36,
                                       test.better="High", group.id="group",
                                       control.id="control")[[2]]
score2adjust(data = PsychTestData, test.score = "scaledscore",
             group.id = "group", control.id = "control",
             demographics = c("age", "male"))
```

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test2norm	<i>Convert raw neuropsychological test scores to demographically adjusted norms.</i>
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## Description

Convert raw neuropsychological test scores to demographically adjusted norms.

## Usage

```
test2norm(
  data = NULL,
  test = NULL,
  test.min = NULL,
  test.max = NULL,
  test.better = c("High", "Low"),
  group.id = NULL,
  control.id = NULL,
  all.controls = FALSE,
  demographics = NULL,
  mfp.alpha = 1,
  rnd.s = TRUE,
  rnd.a = TRUE,
  mean.a = 50,
  sd.a = 10
)
```

## Arguments

data	a data frame containing the variables needed for the norming process. The current version of the function does not accommodate missing data. For best results, exclude cases with missing test scores or missing demographics before applying this function.
test	a character string specifying the name of the test to be normed
test.min	a real number indicating the smallest possible test score
test.max	a real number indicating the largest possible test score
test.better	a character string indicating direction of the scores. Use "High" if high test scores imply better performance, use "Low" otherwise.
group.id	a character string specifying the name of the variable containing group identification (i.e. control vs exposed/test/risk). Ignored, if all.controls = TRUE.
control.id	a character string specifying the label of the control group within group.id variable. Ignored, if all.controls = TRUE.
all.controls	a logical indicating whether all observations should be treated as controls. Overrides group.id and control.id.

demographics	a single or multiple character strings (concatenated by <code>c()</code> function) specifying the names of demographic predictors to be included into normative formulas. Demographic variables should be numeric or binary (0/1).
mfp.alpha	a numeric value between 0 and 1 that sets significance level for inclusion of demographic predictors into normative formula. Passed to the <code>mfp2()</code> function. Default value is 1 for inclusion of all predictors regardless of their significance.
rnd.s	a logical indicating whether the scaled scores should be rounded. Default is TRUE.
rnd.a	a logical indicating whether the adjusted scores (T-scores) should be rounded. Default is TRUE.
mean.a	numeric value for the mean of adjusted score (T-score) distribution. Default is 50.
sd.a	numeric value for the standard deviation of adjusted score (T-score) distribution. Default is 10.

### Details

The `test2norm()` function can be used by neuropsychologists, who wish to construct normative formulas for cognitive tests that adjust for expected effects of demographic characteristics (e.g., age), using methods described in Heaton et al. (2003 & 2009). The norming procedure makes use of the `mfp2()` function from the `mfp2` package to explore nonlinear associations between cognition and demographic variables. The raw test scores that have many decimal digits should be rounded to fewer digits prior to the application of the `test2norm()` function. This will significantly reduce software running time. The recommended number of decimal digits is 4 or fewer. Detailed description of the procedure are found in Umlauf et al. (2024). (Previous versions of the function depended on `mfp` package.)

### Value

A list consisting of 6 objects. The first four are vectors containing the original raw test scores and the calculated scaled scores, demographically adjusted scores, and deficit scores. The fifth object in the list, called `SS.maps`, contains conversions from raw scores to scaled scores in a form of a table with two columns, one representing scaled scores (one per row) and one representing raw scores (range of raw values corresponding to each scaled score). The last item in the output list is also a list called `MFP.formulas` and contains the information for calculation of adjusted scores, including variable transformations (if any), multiple fractional polynomial (MFP) model coefficients, and the standard deviation of residuals resulting from the MFP modeling.

### Author(s)

Anya Umlauf

### References

Umlauf A et al. (2024) Automated procedure for demographic adjustments on cognitive test scores. <doi:10.1080/23279095.2023.2288231>

Heaton RK, Taylor MJ, & Manly J (2003) Demographic effects and use of demographically corrected norms with the WAIS-III and WMS-III. In: Tulskey D et al. (Eds.) *Clinical Interpretation of the WAIS-III and WMS-III*. San Diego, CA: Academic Press, 183-210.

Heaton RK, Ryan L, & Grant I (2009) Demographic influences and use of demographically corrected norms in neuropsychological assessment. In Grant I & Adams KM (Eds.) *Neuropsychological Assessment of Neuropsychiatric and Neuromedical Disorders*. New York, NY: Oxford University Press, 127-155.

Benner A (2005) mfp: Multivariable fractional polynomials. *R News* 5(2): 20–23.

### Examples

```
data(PsychTestData)
test2norm(data = PsychTestData, test = "rawscore",
          test.min = 0, test.max = 36, test.better = "High",
          group.id = "group", control.id = "control",
          demographics = c("age", "male"))
```

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