

# russmisc-2.15.1

December 24, 2012

---

aspect

*Solve right-triangles systems involving (monitor) aspect ratios*

---

## Description

I designed this to calculate values for monitor aspect ratios, but I imagine it is more general. Given any two or three values, solve for the missing values. Note, the values are not checked so the principle of GIGO applies.

## Usage

```
aspect(w = NULL, h = NULL, d = NULL, aspect = NULL)
```

## Arguments

w	Width
h	Height
d	Diagonal
aspect	Aspect ratio (w/h)

## Value

numeric The length of the unspecified side

## References

[http://en.wikipedia.org/wiki/Pythagorean\\_theorem](http://en.wikipedia.org/wiki/Pythagorean_theorem)

## Examples

```
pythag(a=3, c=pythag(3, 4))
```

---

aspell_add_word	<i>Add a word to the end of an aspell dictionary</i>
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---

**Description**

Actually this adds the value in the 'word' parameter to the end of any file specified by the 'file' argument.

**Usage**

```
aspell_add_word(word, dict)
```

**Arguments**

word	The word to be added to the dictionary
dict	The location of the dictionary

**Examples**

```
#mydictloc <- "c:\\rsp\\tools\\aspell\\bin\\aspell.exe"  
#aspell_add_word("pythagorean",mydictloc)
```

---

bmp.save	<i>Save current plot as a bmp</i>
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**Description**

Save current plot as a bmp

**Usage**

```
bmp.save(filename, ...)
```

**Arguments**

filename	The filename for the saved plot
...	options passed to dev.copy

---

center	<i>Center a variable</i>
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**Description**

By centering I mean take the numeric variable and subtract the mean from each value. Then return each value.

**Usage**

```
center(x)
```

**Arguments**

x	Variable to be centered
---	-------------------------

**Value**

Numeric  $x - \text{mean}(x)$

**Examples**

```
center(c(1,2,3))
```

---

checkFarm	<i>Check a farm</i>
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**Description**

See `farm()` for details on farms. This function checks for a file based on the `farmName` parameter called `farmName.res.Rda`. If that file exists it loads it and returns the object stored by the farm in the object `farmName`. If that file does not exist, then the farm is not done processing, and a warning and `NULL` are returned. Note that a rapid loop through `checkFarm()` without `Sys.sleep` produced an error during development.

**Usage**

```
checkFarm(farmName)
```

**Arguments**

farmName	This is the name of the farm, used for creating and destroying filenames. This should be saved from when the <code>farm()</code> is created
----------	---

**See Also**

[farm](#) [waitForFarm](#)

## Examples

```
.tmp <- "This is a test of farm()"
exampleFarm <- farm("
print(.tmp)
helloFarm <- 10+2
farmName <- helloFarm
save(farmName,file=farmResult)
")
example.result <- checkFarm(exampleFarm)
while (is.null(example.result)) {
  example.result <- checkFarm(exampleFarm)
  Sys.sleep(1)
}
print(example.result)
```

---

Conv2Dto3D

*Calculate a real world position of a projected object*

---

## Description

Given the position of an object on the projection plane, the real depth of the object, and the depth of the projection plane, calculate the position of the real-world object

## Usage

```
Conv2Dto3D(ScreenPosition, TargetDepth, ScreenDepth)
```

## Arguments

ScreenPosition	The position of the object on the screen
TargetDepth	The targets' position in depth coordinates
ScreenDepth	The position of the projection plane in depth coordinates

---

Conv3Dto2D

*Calculate the projection position of a real world object*

---

## Description

Given the position of an object in 3D space and the depth of the projection plane, calculate the position of the projection of that object on projection plane

## Usage

```
Conv3Dto2D(TargetPosition, TargetDepth, ScreenDepth)
```

## Arguments

TargetPosition	The targets' position in width and/or height coordinates
TargetDepth	The targets' position in depth coordinates
ScreenDepth	The position of the projection plane in depth coordinates

**Value**

numeric Projection position in the coordinate type specified in TargetPosition

---

convertID	<i>Convert Excel Column ID</i>
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**Description**

This function converts R numeric column IDs to Excel letter IDs and vice versa.

**Usage**

```
convertID(ID)
```

**Arguments**

ID                      Either a character or numeric value specifying the ID to convert.

**Value**

character or numeric; the reverse of what was specified by ID

**Note**

This function is not properly vectorized, it can only handle a single ID

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debackslash	<i>Debackslash</i>
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**Description**

Take the contents of the clipboard and replace backslashes with forward slashes

**Usage**

```
debackslash()
```

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degrees	<i>Convert radians to degrees</i>
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**Description**

Convert radians to degrees

**Usage**

```
degrees(radians)
```

**Arguments**

radians	A value in radians
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**Value**

The value provided in radians converted to degrees

**See Also**

[radians](#)

**Examples**

```
degrees(pi)
```

---

dist3D.l	<i>Helper function for dist3D to handle lists</i>
----------	---

---

**Description**

Helper function for dist3D to handle lists

**Usage**

```
dist3D.l(point1, point2)
```

**Arguments**

point1	I think it wants a list. Frankly, I don't remember how this works.
point2	I think it wants a list. Frankly, I don't remember how this works.

**Value**

numeric ... I think

**Note**

TODO: Figure out how this works.

---

dist3D	<i>Calculate the distance between two 3D points</i>
--------	---

---

**Description**

Calculate the distance between two 3D points

**Usage**

```
dist3D(point1, point2)
```

**Arguments**

point1	The location of the first point as a numeric vector of length 3 c(X,Y,Z)
point2	The location of the second point as a numeric vector of length 3 c(X,Y,Z)

**Value**

numeric distance between two points

**References**

<http://www.calculatorsoup.com/calculators/geometry-solids/distance-two-points.php>

**Examples**

```
dist3D(c(5,6,2),c(-7,11,-13))
dist3D(list(c(0,0),c(0:0),c(0,0)),list(1:2,3:4,5:6))
dist3D(list(5,6,2),list(-7,11,-13))
```

---

droplead0	<i>droplead0</i>
-----------	------------------

---

**Description**

Drop a leading zero from a number and convert to a character string, as in reporting p-values.

**Usage**

```
droplead0(val)
```

**Arguments**

val	The numeric value from which to drop the lead 0
-----	---

---

factory

*Catch errors and warnings and store them for subsequent evaluation*


---

### Description

Factory is not my code. Factory generates a function which is appropriately wrapped by error handlers. The result produced is a list. This is a nice way to recover from a loop problems that may have occurred during loop evaluation. Check the references for additional related functions. I have not included the other factory functions because they did not play well with the return item as an S4 object.

### Usage

```
factory(fun)
```

### Arguments

fun                      The function to be turned into a factory

### Value

list[3]: item 1 is the result from the function and \$warn and \$err contain warning and error messages respectively.

### Author(s)

Martin Morgan; Packaged by Russell S. Pierce

### References

<http://stackoverflow.com/questions/4948361/how-do-i-save-warnings-and-errors-as-output-from-a-f>

### Examples

```
f.log <- factory(log)
f.log("a")
f.as.numeric <- factory(as.numeric)
f.as.numeric(c("a", "b", 1))
```

---

farm

*Create a farm*


---

### Description

A farm is an external self-terminating instance of R to solve a time consuming problem in R. Think of it as a (very) poor-person's multi-core. For a usage example, see checkFarm. Known issues: May have a problem if the library gdata has been loaded.// If a farm produces warnings or errors you won't see them If a farm produces an error... it never will produce a result.



**Usage**

```
farm(commands,
      farmName = paste("farm-", as.integer(Sys.time()) + runif(1), sep = ""),
      Rloc = NULL)
```

**Arguments**

commands	A text string of commands including line breaks to run. This must include the result being saved in the object farmName in the file farmResult (both are variables provided by farm() to the farm).
farmName	This is the name of the farm, used for creating and destroying filenames. One is randomly assigned that is plausibly unique.
Rloc	The location of R.exe. The default loads the version of R that is stored in the windows registry as being \"current\".

**Value**

The farm name is returned to be stored in an object and then used in checkFarm()

**See Also**

[checkFarm](#) [waitForFarm](#)

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fft.binFreqs

---

*Report the frequencies of the the bins in an FFT*


---

**Description**

Report the frequencies of the the bins in an FFT

**Usage**

```
fft.binFreqs(sfreq, Nsample)
```

**Arguments**

sfreq	numeric The number of samples per unit of time, e.g. Hz
Nsample	numeric The number of samples in the time series

**Note**

This code might not work

---

fft.binN	<i>Find the number of FFT bins</i>
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---

**Description**

Find the number of FFT bins

**Usage**

```
fft.binN(Nsample)
```

**Arguments**

Nsample	numeric	The number of samples in the time series
---------	---------	--

---

fft.binSize	<i>Find the size of an FFT bin</i>
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---

**Description**

Find the size of an FFT bin

**Usage**

```
fft.binSize(sfreq, Nsample)
```

**Arguments**

sfreq	numeric	The number of samples per unit of time, e.g. Hz
Nsample	numeric	The number of samples in the time series

---

glibrary	<i>Try to load a library, if that fails, install it, then load it.</i>
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---

**Description**

glibrary short for (get)library. The primary aim of this function is to make loading packages more transparent. Given that we know we want to load a given package, actually fetching it is a formality. glibrary skims past this formality to install the requested package.

**Usage**

```
glibrary(..., lib.loc = NULL, quietly = FALSE,
  warn.conflicts = TRUE, pickmirror = FALSE,
  countrycode = "us")
```

**Arguments**

<code>...</code>	comma seperated package names
<code>lib.loc</code>	See <a href="#">require</a>
<code>quietly</code>	See <a href="#">require</a>
<code>warn.conflicts</code>	See <a href="#">require</a>
<code>pickmirror</code>	If TRUE, glibrary allows the user to select the mirror, otherwise it auto-selects on the basis of the country code
<code>countrycode</code>	glibrary compares this value to results from <code>getCRANmirrors()</code> to select a mirror in the specified country

**Value**

logical; TRUE if glibrary was a success, an error if a package failed to load

**Note**

`keep.source` was an arguement to `require` that was deprecated in R 2.15

This warning `Warning in install.packages: InternetOpenUrl failed: 'The operation timed out'` indicates that the randomly selected repository is not available. Check your internet connection. If your internet connection is fine, set `pickmirror=TRUE` and manually select an operational mirror.

**Examples**

```
#glibrary(lattice,MASS) #not run to prevent needless dependency
```

---

in2m	<i>Convert inches to meters</i>
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---

**Description**

Convert inches to meters

**Usage**

```
in2m(inches)
```

**Arguments**

<code>inches</code>	A value in inches
---------------------	-------------------

**Value**

numeric a value in meters

---

km2mi	<i>Convert kilometers to miles</i>
-------	------------------------------------

---

**Description**

Convert kilometers to miles

**Usage**

km2mi(km)

**Arguments**

km                      numeric value in kilometers (km)

**Value**

numeric a value in miles

---

latex.p	<i>Format a p-value for LaTeX</i>
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---

**Description**

All ps greater than .01 will be reported exactly to two decimal places. All ps greater than .001 but less than .01 will be reported exactly to 3 decimal places. All other ps will be reported at  $p < .001$ .

**Usage**

latex.p(p.val)

**Arguments**

p.val                      A numeric p value

---

latexFigure	<i>Insert a figure in LaTeX</i>
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---

**Description**

Insert a figure in LaTeX

**Usage**

```
latexFigure(file, caption = NULL,
  placementSpecifier = "tdp", clearpage = FALSE)
```

**Arguments**

<code>file</code>	The name of the picture to insert
<code>placementSpecifier</code>	Options for LaTeX that go in brackets when starting a figure. Tells LaTeX where to put the picture. Note that the placement specifier here is the latex default, tdp. When using many images other packages, e.g. floats, morefloats may be useful. Alternatively, a <code>\clearpage</code> can clear out the floats.
<code>clearpage</code>	Whether to trigger a <code>\clearpage</code> after the figure.
<code>caption</code>	(optional) The caption for the figure in LaTeX

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<code>latinsquare.digram</code>	<i>Create a digram-balanced latin square</i>
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---

**Description**

This function creates a digram-balanced latin square as described in Keppel & Wickens, 2004 pg 384-386. The condition orders for subjects are listed by rows, such that row one is the order of conditions for subject 1, and so on. This method of counter-balancing is preferred to so-called cyclic counter-balancing because it balances the order of presentation such that conditions occur in each position equally and they also balance the order of presentation such that each condition occurs before and after each other presentation an equal number of times. See Keppel & Wickens, 2004 for details.

**Usage**

```
latinsquare.digram(conds)
```

**Arguments**

<code>conds</code>	A vector of condition IDs or condition names
--------------------	--

**Details**

This is the deprecated name for [latinSquareDigram](#)

**Value**

Digram balanced latin squares come in pairs when there are an odd number of conditions, but only one square when there are an even number of conditions. This function returns a list with a square (in a matrix) for each item.

---

latinSquareDigram	<i>Create a digram-balanced latin square</i>
-------------------	--

---

### Description

This function creates a digram-balanced latin square as described in Keppel & Wickens, 2004 pg 384-386. The condition orders for subjects are listed by rows, such that row one is the order of conditions for subject 1, and so on. This method of counter-balancing is preferred to so-called cyclic counter-balancing because it balances the order of presentation such that conditions occur in each position equally and they also balance the order of presentation such that each condition occurs before and after each other presentation an equal number of times. See Keppel & Wickens, 2004 for details.

### Usage

```
latinSquareDigram(conds)
```

### Arguments

conds	A vector of condition IDs or condition names
-------	--

### Value

Digram balanced latin squares come in pairs when there are an odd number of conditions, but only one square when there are an even number of conditions. This function returns a list with a square (in a matrix) for each item.

### Author(s)

Russell S. Pierce <Russell.S.Pierce@gmail.com>

### References

Keppel & Wickens, 2004 pp 384-386

### Examples

```
latinsquare.digram(1:3)
latinsquare.digram(LETTERS[1:4])
```

---

lawOfCos	<i>Use the law of sines Use the law of cosines to calculate an angle</i>
----------	--

---

### Description

Given that A, B, and C are the legs of a triangle solve the the opposite angles

### Usage

```
lawOfCos(A, B, C)
```

**Arguments**

A	numeric Length of side A
B	numeric Length of side B
C	numeric Length of side C

**Value**

list(a=numeric angle opposite A,b=numeric angle opposite B,c=numeric angle opposite C)

**References**

<http://oakroadsystems.com/twt/solving.htm#eq30>

**Examples**

```
lawOfCos(3,4,5)
lawOfCos(3,3,3)
stimsize.val <- 10 #the stimulus size
dist <- 15 #the distance
visangle(10,15)-lawOfCos(stimsize.val,pythag(stimsize.val/2,dist),pythag(stimsize.val/2,dist))$a
offset <- 5
visangle(10,15,5)-lawOfCos(stimsize.val,pythag(stimsize.val/2+offset,dist),pythag(stimsize.val/2+offset,dist))$a
```

---

longframe

---

*Transform a wide dataset into a long dataset.*


---

**Description**

There is almost certainly a better way to do this. The aim here is to turn a wide dataset into a long dataset.

**Usage**

```
longframe(wide.data, btwnsubnames, wide.var.names,
  value.name = NA, dropcol = c())
```

**Arguments**

wide.data	The wide dataset
btwnsubnames	The between subject identifiers
wide.var.names	The variables that are in the wide format
value.name	unknown
dropcol	Columns to drop from the long format

**See Also**

[widenames](#)

**Examples**

```
widenames(c("Age", "Workload"), c(2,2))
```

---

m2in	<i>Convert meters to inches</i>
------	---------------------------------

---

**Description**

Convert meters to inches

**Usage**

m2in(m)

**Arguments**

m	A value in meters
---	-------------------

**Value**

numeric a value in inches

---

MAD	<i>Calculate the median absolute deviation (MAD)</i>
-----	--

---

**Description**

Calculate the median absolute deviation (MAD)

**Usage**

MAD(x)

**Arguments**

x	A set of numeric scores
---	-------------------------

---

makewave	<i>Make wave</i>
----------	------------------

---

**Description**

Generate a vector of numeric data points representing a sine wave

**Usage**

```
makewave(freq, phase, amp, Nsamples = time * samplerate,  
          samplerate, time = Nsamples/samplerate,  
          as.time.series = TRUE)
```



**Arguments**

freq	The frequency of cycles per unit time (e.g. Hz) of the sine wave
phase	The frequency in degrees of the sine wave
amp	The amplitude of the sine wave
Nsamples	The number of samples to generate, this defaults to the time * sample rate.
samplerate	The number of samples to create per unit of time
time	The number of units of time represented by the generated sine wave
as.time.series	Should the result be returned as a time series ts()

**Value**

A numeric vector or a time series representing the specified sine wave

**Note**

coh is calculated as  $\text{coh}[i + (j - 1) * (j - 2)/2] <- \text{Mod}(\text{pgram}[i, j])^2 / (\text{spec}[i] * \text{spec}[j])$   
 $\text{Mod}(\text{pgram}[i, j])^2$  #cross-amplitude values

**Examples**

```
plot(makewave(2,0,3.889*1.6,180,60))
sp <- spectrum(ts.union(makewave(15,30,1,180,60),makewave(15.01,60,1,180,60)),span=c(3),taper=0)
with(sp,data.frame(freq=freq,sta.freq=freq/60,phase=round(phase,4),coh=round(coh,4),phase.deg=round(phase/
plot(sp$freq,sp$phase/(2*pi)*180,type="l")
plot(sp,plot.type="phase")
plot(sp,plot.type="coh")
plot(ts.union(makewave(15,30,1,180,60),makewave(15.01,60,1,180,60)))
Mod(mvfft(ts.union(makewave(15,30,1,180,60),makewave(15.01,60,1,180,60))))
```

---

MEANS.STD.CORR

*Write out the variables from X in a MPLUS MEANS STD CORR format*

---

**Description**

Take data.frame x which only has the variables of interest and convert to MEAN STD CORR format.  
Note this has not been tested at all. Also note that this will overwrite any pre-existing file.

**Usage**

```
MEANS.STD.CORR(x, filename = "output.dat")
```

**Arguments**

x	The data.frame with the variables to export
filename	The filename to export to.

**Examples**

```
x <- data.frame(X1=rnorm(20),X2=rnorm(20),X3=rnorm(20)) #example data
MEANS.STD.CORR(x)
```

---

mi2km	<i>Convert miles to kilometers</i>
-------	------------------------------------

---

**Description**

Convert miles to kilometers

**Usage**

mi2km(mi)

**Arguments**

mi                      numeric value in miles (mi)

**Value**

numeric a value in km

---

MSE	<i>Calculate the mean squared error</i>
-----	---

---

**Description**

Calculate the mean squared error

**Usage**

MSE(x, y, ...)

**Arguments**

x                      Numeric data series  
y                      Numeric data series  
...                    further arguments for mean

**Value**

numeric

---

popscale	<i>Population scale/Z scores</i>
----------	----------------------------------

---

**Description**

The built-in R function `scale` uses the sample standard deviation when its `scale` option is set to `TRUE`. This function uses the population standard deviation instead.

**Usage**

```
popscale(x, center = TRUE, scale = TRUE)
```

**Arguments**

<code>x</code>	A numeric vector, matrix, or data.frame
<code>center</code>	A logical indicating whether the scores in the columns in <code>x</code> should have their column means subtracted
<code>scale</code>	A logical indicating whether the scores in the columns in <code>x</code> should be divided by their column standard deviations

**Author(s)**

Borrowed from Ryne Sherman on 2012/09/06, his function is named `scale2`

---

popsd	<i>Calculate the population standard deviation of x Usage popsd(x)</i>
-------	--

---

**Description**

Calculate the population standard deviation of `x` Usage `popsd(x)`

**Usage**

```
popsd(x, nomiss = 0.8)
```

**Arguments**

<code>x</code>	A numeric vector
<code>nomiss</code>	A numeric vector specifying the proportion of valid cases in <code>x</code> (i.e. data that must not be NA) for the sd to be returned

**Note**

R's built-in `sd` function divides the sum of the squared deviations from the mean by the number of observations minus 1 ( $N-1$ ). However, there are times where one would prefer to use the formula with  $N$  in the denominator (e.g. if one is working with the entire population of scores). This function does just that.

**Author(s)**

Borrowed from Ryne Sherman on 2012/09/06

---

product	<i>Take the product of all values in a vector</i>
---------	---

---

**Description**

Take the product of all values in a vector

**Usage**

```
product(x)
```

**Arguments**

x	A vector of type numeric
---	--------------------------

**Value**

numeric The result of systematically multiplying the items in the vector x.

**Examples**

```
product(1:3)
product(c(2,2,4))
```

---

pythag	<i>Calculate the pythagorean theorem to solve for A, B, or C</i>
--------	--

---

**Description**

Given that A and B are the legs of a right triangle and C is its hypotenuse: solve for A, B, or C given any other two.

**Usage**

```
pythag(a = NULL, b = NULL, c = NULL)
```

**Arguments**

a	Leg 1
b	Leg 2
c	Hypotenuse

**Value**

numeric The length of the unspecified side

**References**

[http://en.wikipedia.org/wiki/Pythagorean\\_theorem](http://en.wikipedia.org/wiki/Pythagorean_theorem)

**Examples**

```
pythag(a=3,c=pythag(3,4))
```

---

r2t	<i>Convert r to a t</i>
-----	-------------------------

---

**Description**

Convert r to a t

**Usage**

```
r2t(r, df)
```

**Arguments**

r	The correlation coefficient value
df	the df for the correlation coefficient

---

radians	<i>Convert degrees to radians</i>
---------	-----------------------------------

---

**Description**

Convert degrees to radians

**Usage**

```
radians(degrees)
```

**Arguments**

degrees	A value in degrees
---------	--------------------

**Value**

The value provided in degrees converted to radians

**Examples**

```
radians(90)
```

---

russmisc-package	<i>russmisc-package placeholder</i>
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---

### Description

A repository for a variety of useful functions.

### Details

A repository for a variety of useful functions which may be of interest or otherwise reused. Unless otherwise stated in the function description, all functions are authored by Russell S. Pierce.

### Author(s)

Russell S. Pierce <Russell.S.Pierce@gmail.com>

---

russmisc.refresh	<i>Refresh the russmisc package</i>
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---

### Description

This is just a convenience to unload and reload the russmisc package, useful for rapid development

### Usage

```
russmisc.refresh()
```

---

simpleFarm	<i>Create a one-line simple farm</i>
------------	--------------------------------------

---

### Description

This is a convenience wrapper function that uses farm to create a single farm appropriate for processing single line commands.

### Usage

```
simpleFarm(command,
  farmName = paste("farm-", as.integer(Sys.time()) + runif(1), sep = ""),
  Rloc = NULL)
```

### Arguments

command	A single command
farmName	This is the name of the farm, used for creating and destroying filenames. One is randomly assigned that is plausibly unique.
Rloc	The location of R.exe. The default loads the version of R that is stored in the windows registry as being \"current\".

**Value**

The farm name is returned to be stored in an object and then used in `checkFarm()`

**See Also**

[farm](#), [checkFarm](#), and [waitForFarm](#)

**Examples**

```
a <- 5
b <- 10
farmID <- simpleFarm("a + b")
waitForFarm(farmID)
```

---

SSE	<i>Calculate the sum of squared errors</i>
-----	--

---

**Description**

Calculate the sum of squared errors

**Usage**

```
SSE(x, y, ...)
```

**Arguments**

x	Numeric data series
y	Numeric data series
...	further arguments for mean

**Value**

numeric

---

stimsizes	<i>Calculate a stimulus size given a visual angle and viewing distance</i>
-----------	--

---

**Description**

Calculate a stimulus size given a visual angle and viewing distance

**Usage**

```
stimsizes(visangle.val, viewdist.val)
```

**Arguments**

visangle.val	A value in degrees of visual angle
viewdist.val	A viewing distance

Value

The stimulus size that when centrally presented would have a visual angle (in degrees) of `visangle.val`. Stimulus size is reported in the same units as `viewdist.val`.

See Also

`viewdist` `stimsizesize`

Examples

```
stimsizesize(visangle(10,100),100)
```

---

t2r	Convert t to a r
-----	------------------

---

Description

Convert t to a r

Usage

```
t2r(t, df)
```

Arguments

t	The t statistic
df	The df for the t statistic

---

table.png	Print LaTeX Table Direction to an Image
-----------	---

---

Description

Print LaTeX Table Direction to an Image

Usage

```
table.png(obj, name)
```

Arguments

obj	The matrix to turn into a table
name	The name of the png to produce (without extention)

Author(s)

Found here: <http://stackoverflow.com/questions/9298765/print-latex-table-directly-to-an-image-png-or-other>



---

trimval	<i>Trim values Usage trimval(x, tr=.2, na.rm=TRUE)</i>
---------	--

---

**Description**

Trim values Usage trimval(x, tr=.2, na.rm=TRUE)

**Usage**

```
trimval(x, tr = 0.2, na.rm = TRUE)
```

**Arguments**

x	A numeric vector to be trimmed
tr	The proportion of values in x to be trimmed
na.rm	A logical indicating whether missing values should be removed prior to trimming

**Author(s)**

borrowed from Ryne Sherman on 2012/09/06. Modified to return NA for removed values by Russell S. Pierce

---

unfactor	<i>Unfactor a vector</i>
----------	--------------------------

---

**Description**

Take a factor vector and return a value a vector of either type character or numeric by replacing the factors with their associated labels. If all label names are numeric, a numeric vector will be returned, otherwise a character vector will be returned.

**Usage**

```
unfactor(factors)
```

**Arguments**

factors	The vector of factors to be unfactored
---------	--

**Value**

character or numeric

**Examples**

```
unfactor(factor(c(3,2,1)))
```

---

uniquePerms

*Get a set of minimially replicated permutations*


---

### Description

This function provides a random set of sequences from the full possible set of permutations. If you request more sequences than there are unique ones it will give you a full set of unique permutations/sequences (or more than one) and then randomly select the remaning sequences. The arguments are conds (like in latinsquare.digram) and N (number of sequences requested)

### Usage

```
uniquePerms(conds, N)
```

### Arguments

conds (vector) A of condition IDs or condition names  
N (numeric) The number of sequences desired

### Value

N x conds (matrix) of sequences (by rows)

### Author(s)

Russell S. Pierce <Russell.S.Pierce@gmail.com>

### Examples

```
uniquePerms(LETTERS[1:4],12)
uniquePerms(LETTERS[1:4],36)
```

---

updateTex

*Update a .tex file from .rnw file if needed*


---

### Description

Check the file dates. If .Rnw is newer than .Tex, then re-weave.

### Usage

```
updateTex(filename)
```

### Arguments

filename character the root of the filename, e.g. "writing" when refering to "Writing.Rnw"

---

viewdist	<i>Calculate a viewing distance given a visual angle and stimulus size</i>
----------	--

---

**Description**

Calculate a viewing distance given a visual angle and stimulus size

**Usage**

```
viewdist(visangle.val, stimsized.val)
```

**Arguments**

visangle.val	A value in degrees of visual angle
stimsized.val	A stimulus size

**Value**

The viewing distance of an object of stimsized.val that, when centrally presented, would have a visual angle (in degrees) of visangle.val. Viewing distance is reported in the same units as stimsized.val.

**Author(s)**

Russell S. Pierce <Russell.S.Pierce@gmail.com>

**See Also**

[viewdist visangle](#)

**Examples**

```
viewdist(visangle(10,100),10)
```

---

visangle	<i>Calculate visual angle given a stimulus size and viewing distance</i>
----------	--

---

**Description**

Calculate visual angle given a stimulus size and viewing distance

**Usage**

```
visangle(stimsized.val, viewdist.val, offset = 0)
```

**Arguments**

stimsized.val	The stimulus size
viewdist.val	The viewing distance
offset	The offset of the center of the object from the center of the viewing angle

**Value**

Degrees in visual angle for the given stimulus size

**Note**

The units of measure should be the same for both parameters

**See Also**

`stimsize viewdist link{visangle3D}`

**Examples**

```
visangle(10,100)
visangle(visangle(0.75438,45),viewdist(visangle(0.75438,45),0.75438))
```

---

visangle3D.l	<i>Helper function for visangle3D to handle lists</i>
--------------	---

---

**Description**

Helper function for visangle3D to handle lists

**Usage**

```
visangle3D.l(viewerloc, stimloc, stimdim)
```

**Arguments**

- |           |  |
|-----------|--|
| viewerloc | The location of the center of the viewer as a list   |
| stimloc   | The location of the center of the stimulus as a list |
| stimdim   | The stimulus dimensions as a numeric list            |

---

visangle3D	<i>Calculate visual angle given a 3D viewer position, stimulus size, and stimulus position</i>
------------	--

---

**Description**

This function calculates the visual angle of a stimulus given a viewer position, stimulus size, and stimulus position. Here I make fewer assumptions and calculate the distance between the viewer and the edges of the object in order to use the law of cosines to calculate the associated angle.

**Usage**

```
visangle3D(viewerloc, stimloc, stimdim)
```

**Arguments**

viewerloc	The location of the center of the viewer as a numeric vector of length 3 c(X,Y,Z)
stimloc	The location of the center of the stimulus as a numeric vector of length 3 c(X,Y,Z)
stimdim	The stimulus dimensions as a numeric vector of length 2 c(width,height)

**Value**

Degrees in visual angle for the given stimulus dimentions and positions of viewer and stimuli

**Note**

The units of measure should be the same for all parameters. Z is assumed to be the depth plane.

**See Also**

[visangle](#)

**Examples**

```
#corresponds to an offset of 10 and 10 and a viewer distance of 20
viewerloc <- c(0,0,0)
stimloc<-c(10,10,-20)
stimsz.val <- 10
stimdim <- c(stimsz.val,1)
visangle(stimsz.val,20,pythag(stimloc[1],stimloc[2]))
visangle3D(viewerloc,stimloc,stimdim)
#corresponds to an offset of 10 and 0 and a viewer distance of 20
viewerloc <- c(0,0,0)
stimloc<-c(10,0,-20)
stimsz.val <- 10
stimdim <- c(stimsz.val,1)
visangle(stimsz.val,20,pythag(stimloc[1],stimloc[2]))
visangle3D(viewerloc,stimloc,stimdim)
visangle3D(list(c(1,1),c(1,1),c(1,1)),list(c(2,2),c(2,2),c(2,2)),list(c(3,3),c(3,3)))
visangle3D(list(1,2,3),list(3,4,5),list(2,2))
```

---

waitForFarm

*Wait for a farm result*


---

**Description**

This function repeatedly checks for a farm, when the farm is found it returns the harvest (the farm result object). If the farm terminated with an error or there is some other sort of coding error, waitForFarm will be an infinite loop. As checkFarm produces errors on checks when the harvest is not ready, waitForFarm hides these errors in the factory error-catching wrapper.

**Usage**

```
waitForFarm(farmName, noCheck = FALSE)
```

Arguments

farmName	This is the name of the farm, used for creating and destroying filenames. This should be saved from when the farm() is created
noCheck	If this value is TRUE the check for the farm's .r is skipped. If it is FALSE, the existance of the appropriate .r is checked for before entering a potentially unending while loop.

---

widenames	<i>Generate Names for a Wide Stuctured Dataset</i>
-----------	--

---

Description

Generate Names for a Wide Stuctured Dataset

Usage

widenames(names, levels)

Arguments

names	A vector of type character with the names of the factors
levels	The number of levels for each factor listed in names

See Also

[longframe](#)

Examples

widenames(c("Age", "Workload"),c(2,2))

---

winvar	<i>Calculate the winsorized variance of x Usage winvar(x, tr=.2, na.rm=TRUE) This function is borrowed directly from Rand Wilcox's function and is kept with this set of functions for capability with a large number of functions.</i>
--------	---

---

Description

Calculate the winsorized variance of x Usage winvar(x, tr=.2, na.rm=TRUE) This function is borrowed directly from Rand Wilcox's function and is kept with this set of functions for capability with a large number of functions.

Usage

winvar(x, tr = 0.2, na.rm = TRUE)

**Arguments**

x	A numeric vector of which to get the winsorized variance
tr	The proportion of scores to winsorize
na.rm	A logical indicating whether missing values should be removed prior to calculation.

**Author(s)**

Rand Wilcox, borrowed from Ryne Sherman on 2012/09/06

---

Zmodified

*Calculate the Iglewicz and Hoaglin (1993) Z modified*

---

**Description**

Calculate the Iglewicz and Hoaglin (1993) Z modified

**Usage**

Zmodified(x)

**Arguments**

x	A set of numeric scores
---	-------------------------

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