

# Package ‘zernike’

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**Title** Zernike Polynomials

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mvtnorm, lppuw, clue, minpack.lm

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**LinkingTo** Rcpp, RcppArmadillo

**Description** Routines for Manipulation of Zernike polynomials and  
Interferogram fringe analysis

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**License** MIT and GPL

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addfit	<i>Add zernike coefficients to a matrix.</i>
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## Description

Add zernike coefficients to a matrix.

## Usage

```
addfit(..., th = 0, zcm = NULL, theta = numeric(0))
```

**Arguments**

...	One or more fits as from psifit, etc.
th	Rotation angles, in degrees
zcm	The matrix to be added to (defaults to NULL)
theta	The vector of rotation angles to be added to

**Author(s)**

M.L. Peck

aiapsi

*Iterative algorithms for PSI with unknown phase shifts***Description**

Three iterative algorithms for PSI with unknown phase shifts.

**Usage**

```
aiapsi(im.mat, phases, ptol = 0.001, maxiter=20, trace=1)
aiapsiC(im.mat, phases_init, ptol, maxiter, trace)
hkpsi(im.mat, phases, maxiter = 20, ptol = 0.001,
      trace = 1, plotprogress = TRUE)
tiltpsi(im.mat, phases, coords, ptol = 0.01, maxiter = 20, trace = 1)
tiltpsiC(im.mat, phases, coords, ptol, maxiter, trace)
```

**Arguments**

im.mat	a <i>matrix</i> of interferogram values
phases	Starting guess for phase shifts
ptol	Convergence criterion for phase shifts
maxiter	Maximum number of iterations
trace	Boolean: Print some summary data at each iteration.
plotprogress	Plot some summary data for each iteration? Also, for tiltpsi and tiltpsiC
coords	Low order Zernike polynomial matrix

**Details**

The “variable tilt” algorithm now allows an indefinite number of low order Zernike terms to be variable between phase steps. coords can be created with [zpm](#) setting maxorder to a small value, say 4, discarding the first (dc) column and retaining as many as desired. There must be at least two columns for tilts. The third will be defocus, the next two astigmatism, the next two primary coma, ...

aiapsi and tiltpsi are wrappers for the calls to the C++ code in aiapsiC and tiltpsiC with sensible defaults for ptol, maxiter, and trace.

**Value**

A list containing the following elements:

phi	The wrapped phase estimate. This is a vector as long as the number of rows in im.mat.
mod	Modulation estimate.
phases	Phase shift estimates.
iter	Number of iterations.
sse	Sum squared error at each iteration.

Also, for tiltpsi

zcs	Matrix of Zernike coefficients, with one row for each column in coords and number of columns = number of columns of im.mat.
-----	---

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**References**

Zhaoyang Wang and Bongtae Han, "Advanced iterative algorithm for phase extraction of randomly phase-shifted interferograms," *Opt. Lett.* 29, 1671-1673 (2004).

Han, G-S and Kim, S-W,, "Numerical correction of reference phases in phase-shifting interferometry by iterative least squares fitting," *Applied Optics* 33, 7321-7325 (1994),

Lin, B-J et al., "An iterative tilt-immune phase-shifting algorithm," OSA conference Optical Fabrication and Testing 2010.

**See Also**

[psifit](#)

---

astig.bath

---

*Zernike coefficients for astigmatism due to Bath astigmatism.*


---

**Description**

Calculates Bath astigmatism coefficients with optional rotation of phi degrees.

**Usage**

```
astig.bath(D, rc, s, lambda = 1e-06, phi = 0)
```

**Arguments**

D	Diameter
rc	Radius of curvature
s	separation of reference and test beams
lambda	Wavelength – defaults to 1 nm.
phi	angle of image horizontal relative to interferometer axis, in degrees

**Details**

D, rc, s, and lambda must have the same units.

**Value**

The Zernike coefficients for primary astigmatism terms.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

---

brcutpuw

*Branch cut algorithm for phase unwrapping*


---

**Description**

Solves a modification of the assignment problem to minimize the total length of branch cuts.

**Usage**

```
brcutpuw(phase, pen = 0, details = FALSE)
```

**Arguments**

phase	Matrix containing the wrapped phase map
pen	Optional penalty value for connecting a residue to an edge
details	boolean: if TRUE return some extra details for diagnostics

**Value**

A matrix containing the unwrapped wavefront. If details==TRUE a named list starting with the unwrapped wavefront in puw.

**Author(s)**

M.L. Peck

**See Also**[qpuw idiffpuw](#)**Examples**

```

set.seed(1234)
PW <- wrap(matrix((0:100)*pi/10,101,101))
## need a border of NA's
PW <- cbind(rep(NA,101), PW, rep(NA,101))
PW <- rbind(rep(NA,103), PW, rep(NA,103))
PW <- PW + rnorm(103^2)
mtext(rmap(PW, plot=TRUE))
PU <- brcutpuw(PW, details=TRUE)
image(1:103, 1:103, PU$bcuts, col="blue", add=TRUE)
X11()
image(PU$puw, col=grey256, asp=1, useRaster=TRUE)

```

---

circle.pars

*Pupil parameters*


---

**Description**

Automatically determine the center and radius of a circular interferogram image.

**Usage**

```

circle.pars(im, fw=2, qt=0.995, excl=5, refine=2,
  plots=TRUE, ask=TRUE, details=FALSE)

```

**Arguments**

im	A matrix containing an image of a circular disk
fw	Amount to smooth image
qt	Threshold to accept an edge point, expressed as a quantile
excl	number of pixels around border of frame to exclude
refine	radius range in pixels for a second pass estimate
plots	Plot edge candidates and fit?
ask	Wait for input before displaying fit?
obstructed	Logical: is there a central obstruction?

**Details**

This routine partially implements the Canny algorithm for edge detection. After optionally smoothing the input image the gradient is calculated using a Sobel filter, and edge pixels are identified by locating local maxima in the magnitude of the gradient.

The edge pixels with qt percentile largest gradients are passed to lqs in package MASS to determine robustly the best fit circle.

Finally, if refine > 0, all edge points within +/- refine pixels of the previously determined edge are passed to nls for a second estimate of center point and radius.

**Value**

A list with the following components:

xc	X coordinate of the center of the pupil
yc	Y coordinate of the center of the pupil
rx	Horizontal radius of the pupil
ry	Vertical radius of the pupil = rx
obstruct	Obstruction fraction (always = 0)

**Note**

This routine is only effective on modulation estimates, and will almost certainly fail on interferogram images. Since data quality varies widely considerable experimentation may be needed on any given image. Increasing the smoothing parameter fw helps to suppress artifacts. Depending on how strong the actual edge is compared to artifacts qt may need to be either increased or decreased from the default value.

if details==TRUE several more pieces of data are returned. This is mostly for debugging purposes and may be eliminated in the future.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

Many routines require the pupil parameters in the form returned by circle.pars. For example [psifit](#), [fftfit](#), [pupil](#), etc.

---

`col3d`*OpenGL plot*

---

**Description**

Returns a vector of colors similar to `image()` display.

**Usage**

```
col3d(surf, surf.col=topo.colors(256), zlim = NULL, eqa=FALSE)
```

**Arguments**

<code>surf</code>	A matrix of surface values
<code>surf.col</code>	Color palette for surface
<code>zlim</code>	Range of values to display
<code>eqa</code>	Equal area per color

**Value**

A vector of color values the same length as `surf`.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**References**

The **rgl** package is described at <http://rgl.neoscientists.org/about.shtml>, and available from CRAN.

**See Also**

[plot.pupil](#)



---

convolve2d	<i>2D convolution</i>
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---

**Description**

General 2D convolution using FFTs

**Usage**

```
convolve2d(im, kern)
```

**Arguments**

im	A matrix representing an image
kern	the convolution kernel

**Value**

The filtered matrix im.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[gblur](#). Called by [circle.pars](#).

---

crop	<i>Crop an array</i>
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---

**Description**

Crop a matrix or 3D array. Main application is to trim excess pixels from an image array, wavefront, etc.

**Usage**

```
crop(img, cp, npad = 20)
```

**Arguments**

img	Array to be cropped.
cp	A list describing the pupil boundary.
npad	Amount of padding to leave around the edge.

**Details**

cp is the list provided by [circle.pars](#).

**Value**

im	The cropped array
cp	Revised value of cp

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

---

fftfit

---

*Fourier transform interferogram analysis*


---

**Description**

High level routines for FFT analysis of interferograms.

**Usage**

```
fftfit(imagedata, cp = NULL, fringescale = 1,
      sl = c(1, 1), filter = NULL, taper = 2,
      zlist = makezlist(), zc0 = c(1:3, 6:7),
      satarget = c(0, 0), astig.bath = c(0, 0),
      puw.alg = "qual", uselm = FALSE, sgs = 3, plots = TRUE, CROP = FALSE)
```

**Arguments**

imagedata	A matrix containing the interferogram
cp	A list describing the pupil boundary, as returned by <a href="#">pupil.pars</a>
fringescale	Fringe spacing, in waves. Use 1 for single pass, 0.5 for double pass, etc.
sl	Position of sidelobe in the form c(x,y)
filter	Size of background filter around DC
taper	Size of taper applied to edge of half plane cut
zlist	Indexes of Zernike polynomials to fit to wavefront
zc0	Indexes of Zernike coefficients to be removed from net wavefront
satarget	Target 4th and 6th order SA coefficients in non-null tests of aspheres
astig.bath	Astigmatism coefficients for Bath geometry
puw.alg	Algorithm to use for phase unwrapping
uselm	Logical: use lm() for least squares fit
sgs	Sample Grid Spacing for least squares fits to wavefront values
plots	Logical: plot progress?
CROP	Center and crop maps?

## Details

If `is.null(filter)` (the default), `pick.sidelobe` will be called to select a Fourier domain side-lobe and background filter size.

If `is.null(cp)` `circle.pars` is applied to the modulation to estimate the pupil parameters.

See [wf.net](#) for details of the process of creating net and smoothed wavefronts from raw unwrapped wavefront maps.

`puw.alg` Specifies the unwrapping algorithm. If NULL an algorithm that's likely to be successful will be selected. You can specify an algorithm by choosing `puw.alg=c("brcut", "ls", "lp", "modal", "qual")`.

## Value

A list with the following components:

<code>phase</code>	Wrapped phase map
<code>mod</code>	The estimated modulation
<code>cp</code>	A list describing the pupil boundary
<code>cp.orig</code>	The precropped value of <code>cp</code>
<code>wf.net</code>	Net unsmoothed wavefront; a matrix of class " <a href="#">pupil</a> "
<code>wf.smooth</code>	Net smoothed wavefront
<code>wf.residual</code>	Difference between net wavefront and polynomial fit
<code>fit</code>	Return value from <a href="#">fitzernikes</a>
<code>zcoef.net</code>	Net Zernike coefficients from fit

## Note

These functions are based largely on the work of Roddier and Roddier (1987).

## Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

## References

Roddier, C. and Roddier, F. 1987, **Interferogram analysis using Fourier transform techniques**, *Applied Optics*, vol. 26, pp. 1668-1673.

## See Also

[wf.net](#), [pupil.pars](#), [pick.sidelobe](#).

## Description

Miscellaneous utilities for working with 2D images in the Fourier domain.

## Usage

```
wftophase(X, lambda=1)
padmatrix(X, npad, fill = mean(X, na.rm=TRUE))
submatrix(X, size = 255)
fftshift(X)
.up2(nr, nc=nr)
```

## Arguments

<code>X</code>	A matrix
<code>lambda</code>	Value of the wavelength, in the same units as <code>X</code>
<code>npad</code>	Size of padded matrix
<code>fill</code>	Values to be assigned to padded matrix elements
<code>size</code>	Size of returned matrix
<code>nr</code>	A number
<code>nc</code>	A number

## Details

`wftophase` computes the complex phase from wavefront values.

`padmatrix` pads a matrix to size `npad` x `npad`, placing the original matrix in the lower left hand corner of the padded matrix.

`submatrix` extracts a `size` x `size` matrix from the center of a larger matrix.

`fftshift` shuffles the quadrants of a matrix around to put the DC element (1,1) in the center of the transformed matrix, with spatial frequencies increasing to the right and up.

## Value

A matrix transformation of the input matrix `X`.

`.up2` returns the next higher power of 2 than `max(nr, nc)`.

## Note

These low level routines are used by several higher level functions that operate in the Fourier domain.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[startest](#), [fftfit](#).

---

fitzernikes

*Least Squares fit to Zernike polynomials*

---

**Description**

Performs a least squares fit of a specified set of Zernike polynomials to a vector of wavefront measurements.

**Usage**

```
fitzernikes(wf, rho, theta, phi = 0, maxorder = 14, uselm = FALSE)
```

**Arguments**

wf	A vector of wavefront values
rho	A vector of radial coordinates.
theta	A vector of angular coordinates, in radians.
phi	Orientation of the image, in degrees
maxorder	Maximum Zernike polynomial order
uselm	Boolean: use <code>lm()</code> for least squares fit

**Details**

wf, rho, and theta must be the same length.

**Value**

The model fit as returned by [lm](#), or the coefficients of the least squares fit if `uselm` is FALSE.

**Note**

The model fit is of the form  $wf \sim Z_0 + Z_1 + Z_2 + \dots$ . With the standard ordering of Zernikes  $Z_0$  is the piston term,  $Z_1$  and  $Z_2$  are x and y tilts,  $Z_3$  is defocus, etc.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[zpm](#), [psifit](#), [fftfit](#), [wf.net](#).

foucogram

*Simulate a Foucaultgram***Description**

Simulates the appearance of a wavefront under the Foucault test.

**Usage**

```
foucogram(wf, edgex = 0, phradius = 0, slit = FALSE,
          pad = 4, gamma = 1, map = FALSE, lev = 0.5)
```

**Arguments**

wf	An object of class <code>pupil</code> containing wavefront values
edgex	lateral position of knife edge
phradius	radius of light source
slit	Logical: Is source a slit or pinhole?
pad	pad factor for FFT
gamma	Gamma value for graphics display
map	Logical: Overlay contours from wavefront map?
lev	Increment for contour levels, if used

**Details**

The default value of 0 for `phradius` simulates a monochromatic point source. Try values in the range 10-30 to suppress diffraction effects.

**Value**

A matrix of intensity levels in the simulated image.

**Note**

The key approximations here are treating the light source as monochromatic and spatially coherent, which is usually not the case for an extended source. Also, Fraunhofer diffraction theory is used.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**References**

See [http://home.netcom.com/~mpeck1/astro/foucault/ext\\_foucault.pdf](http://home.netcom.com/~mpeck1/astro/foucault/ext_foucault.pdf) for an outline of the mathematical treatment of an extended source.

**See Also**[pupil](#)

---

`gblur`*Gaussian blur*

---

**Description**

Blur an image by fw pixels

**Usage**

```
gblur(X, fw=0, details=FALSE)
```

**Arguments**

X	A matrix representing an image
fw	Width of the Gaussian convolution kernel, in pixels
details	Return convolution kernel?

**Details**

fw is the standard deviation of the Gaussian.

**Value**

The filtered matrix X.

**Note**

the details option is mostly for debugging purposes and may go away.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**[convolve2d](#)

---

gpcapsi	<i>Generalized Principal components algorithm for phase shifting interferometry</i>
---------	---

---

### Description

A generalized principal components algorithm for phase shifting interferometry developed by the author. This is the “low level” implementation.

### Usage

```
gpcapsi(im.mat, ptol = 0.001, maxiter = 20, trace = 1)
gpcapsiC(im.mat, ptol, maxiter, trace)
```

### Arguments

im.mat	Matrix containing the unmasked pixels from a set of interferograms.
ptol	Convergence tolerance for phase shifts
maxiter	Maximum number of iterations
trace	Print progress of nonlinear solver every trace iterations. Use trace=0 for silent operation.

### Details

gpcapsi is a wrapper to the C++ call in gpcapsiC.

### Value

A list with the following items:

phi	Estimated wrapped phase.
mod	Estimated modulation.
phases	Estimated phase shifts.
snr	An estimate of the S/N of the interferograms.
eigen	Eigenvalues of the crossproduct matrix

### Note

This is the low level interface to the algorithm. The matrix im.mat should contain the unmasked pixel values from the input interferogram array. No checks are made for valid data. This should normally be called through the high level function [psifit](#).

### Author(s)

M. L. Peck



**See Also**[pcapsi psifit](#)

---

`gray256`*8 bit Grayscale*

---

**Description**

A vector of gray scale levels

**Usage**

```
gray256
grey256
```

**Value**

Defined as `gray256 <- grey(seq(0,1,length=256))`

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>.

---

`hypot`*Hypotenuse*

---

**Description**

The Euclidean length of a vector

**Usage**

```
hypot(x)
```

**Arguments**

`x`                      a vector

**Value**

the length of the vector

**Author(s)**

M.L. Peck

**Examples**

```
hypot(c(1,2))
```

idiffpuw

*Phase unwrapping by Integrating DIFFerences***Description**

Simple path following algorithm for two dimensional phase unwrapping.

**Usage**

```
idiffpuw(phase, mask = phase, ucall = TRUE, dx = NULL, dy = NULL)
```

**Arguments**

phase	A matrix of wrapped phase values
mask	Matrix the same size as phase indicating masked pixels
ucall	Boolean: User call?
dx	Matrix of x differences
dy	Matrix of y differences

**Details**

mask indicates pixels that shouldn't be unwrapped. In the simplest (default) case these are just pixels where phase is undefined.

**Value**

if(ucall), a matrix of class "[pupil](#)" with unwrapped wavefront values, otherwise a list with items:

puw	Unwrapped phase
uw	Matrix indicating pixels that have been unwrapped.

**Note**

Both [brcutpuw](#) and [modalpuw](#) call [rmap](#) first to check for the presence of residues. If there are none idiffpuw is guaranteed to work and is called to do the phase unwrapping.

If there *are* residues brcutpuw creates a mask then calls idiffpuw to unwrap unmasked portions of the phase map.

This function is user callable as well; use a call of the form idiffpuw(phase).

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>. Thanks to Steve Koehler for programming ideas to considerably speed up the algorithm.

## References

Ghiglia, D.C., and Pritt, M.D., 1998, **Two-Dimensional Phase Unwrapping**, New York: Wiley & Sons, Inc., ISBN 0-471-24935-1.

## See Also

[rmap](#), [brcutpuw](#), [modalpuw](#)

---

load.images

*Read images*

---

## Description

Loads image files in jpeg or tiff format. load.pgm provides legacy support for reading files in pgm format.

## Usage

```
load.images(files, names=files, channels=c(1,0,0), scale=1, FLIP=FALSE)
load.pgm(files, imdiff=NULL)
```

## Arguments

files	A vector of character strings with file names
names	Original files
channels	channel weights
scale	scale factor for image resize
FLIP	flip image left for right?

## Details

set FLIP=TRUE to reverse mirror imaged interferograms.

## Value

An array containing the contents of the image files.

## Note

load.pgm is the original load.images included for legacy support of greyscale portable anymap files.

## Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

---

`lspsi`*Phase Shifting Interferometry*

---

**Description**

Least squares fitting of phase shifted interferograms.

**Usage**

```
lspsi(images, phases, wt = rep(1, length(phases)))  
lspsiC(images, phases, wt)
```

**Arguments**

<code>images</code>	An array containing the interferogram images
<code>phases</code>	A vector of phase shifts
<code>wt</code>	A vector of weights

**Details**

`images` is a 3 dimensional array with dimensions `nrow` x `ncol` x `length(phases)`, where `nrow` and `ncol` are the number of rows and columns in the individual interferogram images.

`lspsi` reshapes the image array into a matrix and calls `lspsiC` which in turn calls the compiled C++ routine.

**Value**

A list containing the following components:

<code>phi</code>	Estimated wrapped wavefront phase.
<code>mod</code>	Estimated modulation

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[psifit](#)

---

mpinv

---

*Moore-Penrose generalized inverse*


---

**Description**

Computes the Moore-Penrose generalized inverse of a matrix using singular value decomposition.

**Usage**

```
mpinv(X)
```

**Arguments**

X                      A matrix

**Value**

Matrix containing the generalized inverse. If X is an n x m matrix the return will have dimension m x n.

**Note**

The threshold for determining if a matrix is rank deficient is `eps <- .Machine$double.eps * max(dim(X)) * S$d[1]`

**Author(s)**

M. L. Peck

**Examples**

```
X <- matrix(rnorm(18), 6, 3) ## this should be full rank almost always
mpinv(X) %*% X

X <- matrix(1:18, 6, 3) ## this is not
mpinv(X) %*% X
```

---

pcapsi

---

*Vargas et al.'s Principal Components method for PSI*


---

**Description**

Compute the phase using the Principal components algorithm.

**Usage**

```
pcapsi(im.mat, BGSUB)
```

**Arguments**

<code>im.mat</code>	A <i>matrix</i> of interferogram values
<code>BGSUB</code>	Logical - subtract the pixelwise mean as background estimate?
<code>diagpos</code>	controls treatment of singular values of the data matrix

**Details**

Images are input into an array by [load.images](#). This must be reshaped into a matrix for this function. Also, a mask should be applied if available prior to the call.

**Value**

A list containing the following elements:

<code>phi</code>	The wrapped phase estimate. This is a vector as long as the number of rows in <code>im.mat</code> .
<code>mod</code>	Modulation estimate.
<code>phases</code>	Phase shift estimates.
<code>snr</code>	An estimate of the signal to noise ratio in the input data.
<code>eigen</code>	Singular values of the crossproduct matrix.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**References**

- J. Vargas, J. Antonio Quiroga, and T. Belenguer, "Phase-shifting interferometry based on principal component analysis," *Opt. Lett.* **36**, 1326-1328 (2011) <http://www.opticsinfobase.org/ol/abstract.cfm?URI=ol-36-8-1326>
- J. Vargas, J. Antonio Quiroga, and T. Belenguer, "Analysis of the principal component algorithm in phase-shifting interferometry," *Opt. Lett.* **36**, 2215-2217 (2011) <http://www.opticsinfobase.org/ol/abstract.cfm?URI=ol-36-12-2215>

**See Also**

[pcafit](#), [wf.net](#)

---

`pick.sidelobe`*Select an interferogram sidelobe in the Fourier domain*

---

**Description**

Interactively locate the center of a first order sidelobe in the FFT of an interferogram, and mark the width of the background filter.

**Usage**

```
pick.sidelobe(imagedata, logm=FALSE, gamma=3)
```

**Arguments**

<code>imagedata</code>	A matrix containing an interferogram image
<code>logm</code>	Logical: pass <code>fn="logMod"</code> to <a href="#">plot.cmat</a> ?
<code>gamma</code>	gamma value for display

**Details**

Uses the basic graphics utility [locator](#).

**Value**

A list with the following components:

<code>sl</code>	The coordinates $c(x, y)$ of the selected sidelobe
<code>filter</code>	Estimated size of background filter

**Note**

The high level FFT interferogram analysis routine [fftfit](#) requires the approximate location of the intended first order interferogram sidelobe to be specified.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[fftfit](#),

---

plot.cmat	<i>Plot a complex matrix</i>
-----------	------------------------------

---

## Description

Plot a real valued function of a complex matrix

## Usage

```
plot.cmat(X, fn = "Mod", col = topo.colors(256),
cp=NULL, zoom=1, gamma=1, ...)
```

## Arguments

X	A complex valued matrix
fn	A function returning a real value
col	Color palette for graph
cp	pupil parameters as returned by <a href="#">pupil.pars</a>
zoom	zoom factor for display
gamma	gamma value for display
...	Other parameters to pass to <a href="#">image.default</a>

## Details

In addition to the functions described in [complex](#) fn can be assigned the values "logMod", which will call an internally defined function returning the value  $\log(1+\text{Mod}(X))$ , "Mod2" to plot the power spectrum, and "logMod2" to plot the logarithm of the power spectrum.

If the parameter cp is passed axes will display spatial frequencies in cycles per pupil radius.

## Value

none

## Note

This is used primarily for displaying FFT's of interferograms. In the case of an interferogram in which the background has not been removed use fn="logMod" to make the first order sidelobes visible.

## Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

## See Also

[localize.sidelobe](#), [fftfit](#).



---

plot.pupil

---

*Pupils and wavefronts*

---

## Description

Plot and summary methods for objects of class "pupil".

## Usage

```
plot.pupil(wf, cp=NULL, col = topo.colors(256), addContours = TRUE, cscale = FALSE,
  eqa=FALSE, zlim=NULL, ...)
summary.pupil(wf)
```

## Arguments

wf	An object of class "pupil"
cp	Pupil parameters; a list as returned by <a href="#">pupil.pars</a>
col	Color palette for plot
addContours	Logical: add contour lines?
cscale	Add a color scale legend?
eqa	Perform an "equal area" plot?
zlim	z limits to pass to image
...	Additional parameters to pass to <a href="#">image.default</a>

## Details

These give simple plot and summary methods for objects of class [pupil](#).

If eqa is TRUE, each color in the palette will be used for an equal number of pixels (as opposed to representing an equal interval). Note: the color scale (when cscale == TRUE) may be inaccurate if a very small number of colors are used.

## Value

none

## Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

## See Also

[pupil](#), [pupilrms](#), [pupilpv](#), [strehlratio](#), [pupil.pars](#).

---

plotn

---

*Wavefront comparison plots*


---

## Description

Plot an arbitrary number of wavefronts and all differences.

## Usage

```
plotn(..., labels = NULL, addContours=FALSE, wftype = "net",
col = rygcb(400), qt = c(0.01, 0.99))
```

## Arguments

<code>...</code>	List of wavefront estimates as returned by <a href="#">wf.net</a> .
<code>labels</code>	Labels to identify the wavefronts.
<code>addContours</code>	Boolean to add contours to top row plots
<code>wftype</code>	If the inputs are from <code>wf.net</code> , one of "net", "smooth", "residual".
<code>col</code>	Color palette for top row of plot
<code>qt</code>	Quantiles of differences to plot in comparisons.

## Details

`...` can be any number of objects containing wavefront estimates as returned for example by [wf.net](#).

Wavefronts are displayed on the top row, and differences of all pairs on subsequent rows. Grayscale is used to render the difference plots, and the color palette given in `col` is used for the wavefronts.

## Value

none

## Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

## See Also

[plot.pupil wf.net](#)

---

plotxs

---

*Plot cross-sections (profiles) through a wavefront map.*


---

## Description

Plots an arbitrary number of cross-sections through a wavefront map, with one highlighted.

## Usage

```
plotxs(wf, cp, theta0 = 0, ylim = NULL, N = 4, n = 101,
col0 = "black", col = "gray", lty = 2)
```

## Arguments

wf	A matrix of wavefront values.
cp	List of pupil parameters as returned by <a href="#">pupil.pars</a> .
theta0	Angle of highlighted profile, in degrees.
ylim	range of heights to plot.
N	Number of cross sections.
n	Number of points for each cross section.
col0	Highlight color.
col	Cross section color.
lty	Line type for plots.

## Details

The cross sections are equally spaced in angle from 0 to  $\pi \cdot (N-1)/N$ . Any angle can be specified for the highlighted profile at `theta0`.

## Value

none

## Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

## See Also

[plot.pupil](#) is the main wavefront plotting routine.

psfit\_options

*Options for package zernike***Description**

Parameters controlling various aspects of PSI algorithms, Zernike polynomial fitting, and data display

**Usage**

```
psfit_options
get_zoptions()
```

**Format**

Default parameter values are in the list psfit\_options:

```
refine TRUE
puw_alg "qual"
fringescale 1
maxiter 20
ptol 1e-04
trace 1
nzcs 2
zc0 c(1, 2, 3, 6, 7)
satarget c(0, 0)
astig,bath c(0, 0)
maxorder 14
uselm FALSE
sgs 1
plots TRUE
crop FALSE
colors topo.colors(256)
```

**Details**

psfit\_options is defined in the package environment. To retrieve a copy use get\_zoptions. The list can then be modified and passed to [psifit](#).

Parameters you might want to set include zc0 which sets the target SA for “numerical nulling.”

ptol sets convergence tolerances for iterative PSI algorithms. These have different definitions and different values may be suitable for different algorithms.

If you don’t like the default color palette there are many other choices. If you like rainbows [rygcb](#) defined in this package produces a relatively perceptually uniform version.

The number of variable Zernike terms in the algorithm [tiltpsi](#) is controlled by nzcs. Set it to 3 to include defocus, 5 to include primary astigmatism.

**Author(s)**

M.L. Peck <mlpeck54@gmail.com>

---

psifit

---

*Phase Shifting Interferometry*


---

**Description**

High level function for Least squares analysis of phase shifted interferograms.

**Usage**

```
psifit(images, phases, cp = NULL, satarget = NULL, psialg = "ls", options = psifit_options)
```

**Arguments**

images	An array containing the interferogram images
phases	A vector of phase shifts
cp	A list describing the pupil boundary, as returned by <a href="#">pupil.pars</a>
satarget	Target 4th and 6th order SA coefficients in non-null tests of aspheres
psialg	String identifying the PSI algorithm to use
options	a list of options

**Details**

images is a 3 dimensional array with dimensions nrow x ncol x length(phases), where nrow and ncol are the number of rows and columns in the individual interferogram images.

The current values recognized for psialg are

**ls** least squares with known phase shifts

**aia** the “advanced iterative algorithm“ [aiapsi](#)

**pc1** pca with group\_diag = "v"

**pc2** pca with group\_diag = "u"

**gpc** my generalized PC algorithm in [gpcapsi](#)

**gpcthentilt** first [gpcapsi](#) the [tiltpsi](#)

**tilt** tiltpsi

**Value**

A list with the following components	
phi	wrapped phase estimate
mod	modulation estimate
phases	phase shifts
cp	the interferogram boundary
wf.net	net, unfiltered wavefront (see <a href="#">wf_net</a> )
wf.smooth	Zernike fit wavefront
wf.residual	the difference
fit	Coefficients of Zernike fit to wavefront
zcoef.net	Net Zernike coefficients
extras	any extra data returned by low level functions

**Author(s)**

M.L. Peck <[mlpeck54@gmail.com](mailto:mlpeck54@gmail.com)>

**See Also**

[lspsi](#), [aiapsi](#), [tiltpsi](#), [gpcapsi](#), [pcapsi](#)

---

pupil	<i>Pupils and wavefronts</i>
-------	------------------------------

---

**Description**

Create a pupil object and optionally fill it with a wavefront. For our purposes a “pupil” is defined to be a matrix representation of a circular or annular aperture. Simple plot and summary methods are also provided.

**Usage**

```
pupil(zcoef=NULL, zlist=makezlist(), phi=0, piston=0,
      nrow=256, ncol=nrow, cp=list(xc=128,yc=128,rx=127,ry=127,obstruct=0),
      obstruct=NULL)
pupil.arb(zcoef=NULL, zlist=makezlist(), phi=0, piston=0,
          nrow=256, ncol=nrow, cp=list(xc=128,yc=128,rx=127,ry=127,obstruct=0),
          obstruct=NULL)
```

**Arguments**

zcoef	A vector of Zernike coefficients
zlist	List of indexes the same length as zcoef
phi	Amount to rotate image, in degrees
piston	Constant to add to wavefront values
nrow	Number of rows in output matrix
ncol	Number of columns in output matrix
cp	A list with items xc - x coordinate of central pixel, yc - y coordinate of central pixel, rx - x radius in pixels, ry - y radius in pixels, obstruct - central obstruction fraction.
obstruct	Obstruction fraction

**Details**

[plot.pupil](#) and [summary.pupil](#) provide simple plot and summary methods for objects of class "pupil".

`pupil.arb` will accept an arbitrary list of Zernikes.

`pupil` requires a complete set of Zernikes as returned by [makezlist](#).

**Value**

A matrix of size `nrow` x `ncol`. The matrix is assigned to the class "pupil". NA's are used to fill the matrix outside the defined area of the pupil.

**Note**

The parameter `cp` is used to define the dimensions of the pupil. See [pupil.pars](#) for details.

`obstruct` is included twice for backward compatability.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[Zernike](#), [makezlist](#), [pupilrms](#), [pupilpv](#), [strehlratio](#), [pupil.pars](#), [circle.pars](#).

**Examples**

```
wf <- pupil(zcoef=rnorm(length(makezlist())$n), 0, 0.01))
plot(wf, addContours=FALSE)
summary(wf)
```

---

pupil.pars

*Pupil parameters*


---

### Description

Interactively determine the center, radius, and obstruction fraction of a circular or annular interferogram image.

### Usage

```
pupil.pars(im = NULL, obstructed = FALSE)
```

### Arguments

im	A matrix containing an interferogram image
obstructed	Logical: is there a central obstruction?

### Details

In `pupil.pars`, if the image has already been plotted `im` can be `NULL`, which is the default.

### Value

A list with the following components:

xc	X coordinate of the center of the pupil
yc	Y coordinate of the center of the pupil
rx	Horizontal radius of the pupil
ry	Vertical radius of the pupil
obstruct	Obstruction fraction

### Note

`pupil.pars` uses the basic graphics library routine [locator](#) to interactively mark the edge of the pupil, and optionally the edge of the obstruction. After right clicking to terminate `locator()` a least squares fit is performed to the marked points to determine the center and radius of the pupil.

Note that all routines that make use of Zernikes implicitly assume a circular pupil, or an annular one with small obstruction. We allow `rx != ry` for imaging sensors with non-square aspect ratios.

### Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

### See Also

Many routines require the pupil parameters in the form returned by `pupil.pars`. For example [psifit](#), [fftfit](#), [pupil](#), etc.



---

pupil.rhotheta	<i>Polar coordinates</i>
----------------	--------------------------

---

**Description**

Calculate matrixes of polar coordinates for [pupil](#)'s.

**Usage**

```
pupil.rhotheta(nrow, ncol, cp)
```

**Arguments**

nrow	Number of rows in interferogram images
ncol	Number of columns in interferogram images
cp	A list describing the pupil boundary, as returned by <a href="#">pupil.pars</a>

**Value**

A list with the following components:

rho	A matrix of radial coordinates
theta	A matrix of angular coordinates

**Note**

My Zernike polynomial routines work in polar coordinates, which this function provides. Also, NA's are used to fill the matrix outside the pupil boundary, making the returned values convenient for selecting pixels inside interferograms.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[Zernike](#), [pupil](#).

pupilrms

*Wavefront statistics***Description**

Compute basic statistics of wavefronts stored in "pupil" objects.

**Usage**

```
pupilrms(pupil)
pupilpv(pupil)
strehlratio(rms)
```

**Arguments**

pupil	A matrix of class "pupil"
rms	An rms wavefront error

**Value**

Estimates of the RMS and P-V wavefront errors. strehlratio calculates Mahajan's approximation to the Strehl ratio.

**Note**

pupilrms simply calculates the standard deviation of finite values in the matrix pupil. This is a crude, but usually accurate enough estimate of the true RMS wavefront error.

[summary.pupil](#) calls these functions.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**References**

Schroeder, D.J. 2000, *Astronomical Optics, 2nd Edition*, Academic Press, chapter 10.

**See Also**

[summary.pupil](#).

**Examples**

```
zcoef <- rnorm(length(makezlist())$n), 0, 0.01)
wf <- pupil(zcoef=zcoef)
plot(wf)
summary(wf)
sqrt(crossprod(zcoef)) # A more accurate estimate of RMS
```

---

qpuw*Quality guided algorithm for phase unwrapping*

---

**Description**

Quality guided algorithm for two dimensional phase unwrapping.

**Usage**

```
qpuw(phase, qual)
```

**Arguments**

phase	A matrix of wrapped phase values
qual	A matrix of quality values the same size as phase.

**Value**

puw	A matrix of class " <a href="#">pupil</a> " with the unwrapped wavefront.
-----	---

**Note**

This is a straightforward implementation of the quality guided algorithm of G&P.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**References**

Ghiglia, D.C., and Pritt, M.D., 1998, **Two-Dimensional Phase Unwrapping**, New York: Wiley & Sons, Inc., ISBN 0-471-24935-1.

**See Also**

[idiffpuw](#), [brcutpuw](#), [modalpuw](#), [lspuw](#)

---

readjpeg	<i>Read a jpeg or tiff file</i>
----------	---------------------------------

---

**Description**

Reads a jpeg or tiff file and combines the channels to produce a monochrome image in a matrix.

**Usage**

```
readjpeg(filename, channels)
readtiff(filename, channels)
```

**Arguments**

filename	File name
channels	A vector of length 3 with the channel weights

**Details**

Values in channels should be non-negative, but need not add to one.

**Value**

A double precision matrix with the image data.

**Note**

The matrix must have rows reversed and transposed to display properly with `image()`.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

---

rescale	<i>Rescale an image.</i>
---------	--------------------------

---

**Description**

Rescale a matrix containing a bitmapped image using bilinear interpolation.

**Usage**

```
rescale(im, scale)
```

**Arguments**

im	A matrix with image data.
scale	Scale factor.

**Details**

A value <1 will shrink the image.

**Value**

A matrix containing the rescaled image data.

**Note**

NA's are OK.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

Called by [load.images](#) if necessary.

---

rmap

*Utilities for phase unwrapping*

---

**Description**

Utility functions for use in 2D phase unwrapping.

**Usage**

```
rmap(phase, dx = NULL, dy = NULL, plot = FALSE, ...)  
wrap(phase)
```

**Arguments**

phase	Matrix of wrapped phase values
dx	Matrix of x differences
dy	Matrix of y differences
plot	Boolean: plot residue positions?
...	additional arguments for image

**Details**

dx and dy must have the same dimension as phase.

**Value**

In rmap if plot == TRUE

nr                    the number of residues identified in the map

otherwise

phase                wrapped phase returned by wrap

residues             Matrix the same size as phase with residues marked as + or - 1.

**Note**

These are primarily intended for internal use but can be used interactively. Calling rmap(phase, plot=TRUE) will plot the positions of residues and return nothing. If (plot==FALSE) in the call to rmap a matrix the same size as phase is returned with residues identified with values of +1 or -1.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>. Steve Koehler is responsible for the efficient implementation of the wrap function.

**References**

Ghiglia, D.C., and Pritt, M.D., 1998, **Two-Dimensional Phase Unwrapping**, New York: Wiley & Sons, Inc., ISBN 0-471-24935-1.

**See Also**

Called by [modalpuw](#), [idiffpuw](#), [brcutpuw](#).

---

rygcb

*A better rainbow.*


---

**Description**

Produces a rainbow color palette with colors ranging from "red" to "blue" or "magenta". Perceptual uniformity should be superior to R's rainbow.

**Usage**

```
rygcb(n)
rygcbm(n)
```

**Arguments**

n                    Number of color levels

Details

The palette is created using colorRampPalette.

Value

A vector of colors.

Note

The call to colorRampPalette sets space="Lab" and interpolate="spline" with the intent of creating a more perceptually uniform rainbow.

Author(s)

M.L. Peck

See Also

[grey256](#)

Examples

```
plotsp <- function(spectrum) {
  sl <- length(spectrum)
  rgbv <- col2rgb(spectrum)
  plot((0:(sl-1))+0.5, rgbv[1,], type="l", col="red", xlim=c(0,sl),ylim=c(0,300),xlab="Index",ylab="Channel value")
  points((0:(sl-1))+0.5, rgbv[2,], type="l", col="green")
  points((0:(sl-1))+0.5, rgbv[3,], type="l", col="blue")
  grid()
  rect(0:(sl-1), 260, 1:sl, 300, col=spectrum, density=NA)
}
plotsp(rygcb(400))
X11()
plotsp(rygcbm(500))
```

---

sconic	<i>Sconic</i>
--------	---------------

---

Description

twice the radial height difference between a sphere and conic surface

Usage

```
sconic(D, rc, b = -1, lambda = 632.8, nmax = 6)
```

**Arguments**

D	Diameter (mm)
rc	Radius of curvature (mm)
b	conic constant
lambda	source or test wavelength (nm)
nmax	maximum polynomial order

**Value**

Zernike polynomial coefficients

**Note**

This estimates twice the radial distance between a sphere and conic surface with same paraxial radius of curvature, and returns Zernike coefficients of polynomial expansion. Intended for “numerical nulling” when testing an asphere at center of curvature, and should be more accurate than the vertical height difference calculated by [zconic](#) for that purpose.

**Author(s)**

M.L. Peck

**See Also**

[zconic](#)

**Examples**

```
2.*zconic(1000,5000)
sconic(1000,5000)
```

---

separate.wf

*Separate wavefronts*

---

**Description**

Separate “polished in” from “instrumental” aberrations if possible

**Usage**

```
separate.wf(zcm, theta, maxorder = 14)
```

**Arguments**

zcm	Matrix of observed Zernike coefficients
theta	Vector of rotation angles (in radians)
maxorder	Maximum Zernike order to extract



**Value**

zcb                      Table of extracted coefficients and standard errors

**Author(s)**

M.L. Peck

---

startest

*Star test simulator*


---

**Description**

Simulates a star test.

**Usage**

```
startest(wf=NULL, zcoef=NULL, zlist=makezlist(), phi=0,
lambda = 1, defocus=5,
nrow = 255, ncol = nrow,
cp = list(xc=128,yc=128,rx=127,ry=127,obstruct=0),
obstruct=NULL, npad = 4,
gamma=2, psfmag=2, displaymtf=TRUE, displaywf=FALSE)
```

**Arguments**

wf	A matrix of class pupil containing wavefront values
zcoef	Vector of Zernike coefficients
zlist	Indexes of Zernike coefficients
phi	Angle to rotate wavefront
lambda	Wavelength, in same units as coefficients
defocus	Amount of defocus in waves
nrow	# rows in pupil matrix
ncol	# columns in pupil matrix
cp	pupil parameters
obstruct	Obstruction fraction
npad	Pad factor for FFT
gamma	Gamma value for graphics display
psfmag	Magnification factor for in focus PSF display
displaymtf	Logical: Display MTF?
displaywf	Logical: Display calculated wavefront?

**Details**

If wf is NULL the wavefront is calculated from the the Zernike coefficients (which should be non-NULL).

**Value**

A list with the following components:

psf	The in focus point spread function.
otf	The complex optical transfer function, a complex matrix of size pupilsize.
mtf	The modulation transfer function, a real matrix of size pupilsize.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**References**

Born, M. and Wolf, E. 1999, *Principles of Optics, 7th Edition*, Cambridge University Press.  
 Suiter, H. R., 1994, *Star Testing Astronomical Telescopes*, Willman-Bell, Inc.

**See Also**

[Zernike](#), [pupil](#).

**Examples**

```
# a random, but probably almost diffraction limited, wavefront

temp <- startest(zcoef=rnorm(length(makezlist())$n), mean=0, sd=0.01), zlist=makezlist(), displaywf=TRUE)
```

---

synth.interferogram     *Synthetic interferogram*

---

**Description**

Compute and display a synthetic interferogram.

**Usage**

```
synth.interferogram(wf = NULL, zcoef = NULL, zlist = NULL,
  nr = nrow(wf), nc = ncol(wf), cp = NULL,
  phi = 0, addzc = rep(0, 4), fringescale = 1, plots = TRUE)
```

**Arguments**

wf	A matrix of wavefront values
zcoef	A vector of Zernike coefficients
zlist	A list of Zernike indexes
nr	Number of rows in the output matrix
nc	Number of columns in the output matrix
cp	A list describing the pupil boundaries, as created by <a href="#">pupil.pars</a>
phi	Amount to rotate the wavefront, in degrees
addzc	A 4-vector with piston, tilt, and defocus terms to be added
fringescale	Fringe scale. Should be 1 for single pass, 0.5 for double, etc.
plots	Logical: Plot the interferogram?

**Details**

Either wf or zcoef should be non-null, but not both. If zcoef is specified zlist must be as well. Additional piston, tilt, and defocus terms can be added to the calculated wavefront using addzc.

**Value**

A matrix of intensity levels in the calculated interferogram, assigned class "[pupil](#)".

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[pupil.](#)

**Examples**

```
# create a list of zernikes
zcoef <- rnorm(length(zlist.fr$n), mean=0, sd=0.01)

iwf <- synth.interferogram(zcoef=zcoef, zlist=zlist.fr)

X11()

# show again with some tilt

iwf <- synth.interferogram(zcoef=zcoef, zlist=zlist.fr, addzc=c(0,5,5,0))
```

---

turbwf	<i>Kolmogorov Turbulence</i>
--------	------------------------------

---

### Description

Simulates the optical effects of atmospheric turbulence using Noll's (1976) calculation of the covariance matrix of Zernike polynomials under Kolmogorov turbulence.

### Usage

```
turbwf(friedratio = 1, zlist = makezlist(2, 40), reps = 1)
```

### Arguments

friedratio	Ratio of pupil diameter to Fried parameter
zlist	A list of Zernikes, as returned for example by <a href="#">makezlist</a>
reps	Number of draws to simulate

### Details

The default value of zlist has 440 elements, which may be more than necessary for a reasonable representation of an “atmospheric” wavefront.

### Value

A list with the following components:

zcoef.turb	A reps x length(zlist\$n) matrix of simulated draws of Zernike coefficients.
V	Covariance matrix of the indexed Zernikes.

### Note

Typos in the original source material have been corrected in the code. Note that scintillation is not modelled.

### Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

### References

Noll, R.J. 1976, **Zernike polynomials and atmospheric turbulence**, *J. Opt. Soc. Am.*, Vol. 66, No. 3, p. 207.

### See Also

[Zernike](#), [pupil](#).

## Examples

```
# Simulate a single draw from a turbulent atmosphere
zcoef.turb <- turbwf(friedratio=5, zlist=makezlist(2,30), reps=1)$zcoef.turb
# Warning: this can take a while
wf <- pupil(zcoef=zcoef.turb, zlist=makezlist(2,30))
plot(wf)
summary(wf)
```

---

wf3d.pupil

OpenGL wavefront plot

---

## Description

Interactive plot of a wavefront using the OpenGL package **rgl**. This is a 3D plotting method for objects of class "[pupil](#)".

## Usage

```
wf3d.pupil(wf, cp=NULL, zoom.wf = 1, surf.col = topo.colors(256), bg.col = "black",
  eqa=FALSE)
```

## Arguments

wf	A matrix of wavefront values
cp	A list describing the pupil boundary
zoom.wf	Zoom factor for heights
surf.col	Color palette for surface
bg.col	Background color
eqa	Equal area per color?

## Details

The default color palette will match the colors in the default version of [plot.pupil](#).

## Value

none

## Author(s)

M.L. Peck <mpeck1@ix.netcom.com>

## References

The **rgl** package is described at <http://rgl.neoscientists.org/about.shtml>, and available from CRAN.

See Also

[plot.pupil](#)

Examples

```
# create a random wavefront

wf <- pupil(zcoef=rnorm(length(makezlist())$n), mean=0, sd=0.01))
# the default method

plot(wf)

#this is more fun

wf3d(wf)
```

---

wf_net	<i>Wavefront smoothing</i>
--------	----------------------------

---

Description

Calculate net and smoothed wavefronts from a raw wavefront containing low order nuisance aberrations.

Usage

```
wf_net(wf.raw, cp, options)
```

Arguments

- wf.raw            Raw wavefront to be processed
- cp                a list describing the pupil boundary
- options           a list of options. See [psfit\\_options](#)

Details

Called by [psifit](#)

Value

A list with the following components:

- wf.net            Net unsmoothed wavefront; a matrix of class "[pupil](#)"
- wf.smooth        Net smoothed wavefront
- wf.residual      Difference between net wavefront and polynomial fit
- fit               Return value from [fitzernikes](#)
- zcoef.net        Net Zernike coefficients from fit

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

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zconic

*Zernike coefficients for a conic surface*

---

**Description**

Calculates the radially symmetric Zernike coefficient values up to order nmax for a conic surface relative to a sphere of the same paraxial radius of curvature.

**Usage**

```
zconic(D, rc, b = -1, lambda = 1e-06, nmax = 6)
```

**Arguments**

D	Diameter
rc	Radius of curvature
b	Conic constant
lambda	Wavelength – defaults to 1 nm.
nmax	Maximum radial polynomial order

**Details**

D, rc, and lambda must have the same units.

**Value**

A vector of length nmax/2-1 of coefficient values, in increasing radial order, n=c(4,6, ...).

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**See Also**

[Zernike](#)

**Examples**

```
zconic(200,2000)
zconic(10, 20, b=-1.05, lambda=632.8E-9, nmax=12)
```

Zernike

*Zernike Polynomials***Description**

Routines for creating and manipulating Zernike polynomials.

**Usage**

```
Zernike(rho, theta, n, m, t)
DZernike(rho, theta, n, m, t)
DTZernike(rho, theta, n, m, t)
rzernike(rho, n, m)
drzernike(rho, n, m)
```

**Arguments**

rho	normalized radius, $0 \leq \rho \leq 1$
theta	angular coordinate
n	radial polynomial order
m	azimuthal order
t	character for trig function: one of c("n", "c", "s")

**Note**

These functions return Zernikes scaled such that they form an orthonormal basis set for the space of functions defined on the unit circle. Note that this is not the most commonly used definition (as given e.g. in *Born and Wolf*). The definition I use is often associated with *Noll (1976)*.

The function `zmult` can be used to convert between normalized and conventionally defined vectors of Zernike coefficients.

The basic low level functions `rzernike` and `drzernike` use numerically stable recurrence relationships for the radial Zernikes.

**Author(s)**

M.L. Peck <mpeck1@ix.netcom.com>

**References**

Born, M. and Wolf, E. 1999, *Principles of Optics, 7th Edition*, Cambridge University Press, chapter 9 and appendix VII.

Noll, R.J. 1976, **Zernike polynomials and atmospheric turbulence**, *J. Opt. Soc. Am.*, Vol. 66, No. 3, p. 207.

<http://wyant.opt-sci.arizona.edu/zernikes/zernikes.htm>

<http://mathworld.wolfram.com/ZernikePolynomial.html>



**See Also**

[makezlist](#), [zlist.fr](#), [zmult](#), [fillzm](#), [pupil](#), [pupilrms](#), [pupilpv](#), [strehlratio](#).

**Examples**

```
Zernike(1, 0, 4, 0, "n") # == sqrt(5)

# A slightly more complex example

rho <- seq(0, 1, length = 101)
theta <- rep(0, 101)

plot(rho, Zernike(rho, theta, 6, 0, "n"), type="l",
      ylim=c(-3.5,3.5), main="Some 6th order Zernike Polynomials")
lines(rho, Zernike(rho, theta, 5, 1, "c"), lty=2)
lines(rho, Zernike(rho, theta, 4, 2, "c"), lty=3)
lines(rho, Zernike(rho, theta, 3, 3, "c"), lty=4)
```

---

zlist

---

*Lists of Zernike polynomial indexes*


---

**Description**

Ordered lists of Zernike polynomial indexes.

**Usage**

```
makezlist(minorder = 2, maxorder = 14)
zlist.fr
zmult(zlist = makezlist())
```

**Arguments**

minorder	minimum value of n+m
maxorder	maximum value of n+m
zlist	a list of the form returned by makezlist

**Details**

Zernike polynomials are indexed by a radial index n, an azimuthal index m, and include cosine, sine, and radial terms. These routines return lists of indexes using a popular ordering scheme for Zernike polynomials.

**Value**

makezlist and zlist.fr return lists with the following components:

n	radial order
m	azimuthal order
t	one of c("c", "s", "n")

zmult returns a vector the same length as the components of zlist.

**Note**

zlist.fr is an augmented “Fringe” set of Zernike polynomials equivalent to makezlist(2,12).

makezlist returns a complete list of indexes for all orders from minorder through maxorder, where “order” is the value of n+m.

**Author(s)**

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**See Also**

Virtually all high level functions that work with Zernike polynomials use these lists. See for example [pupil](#), [psifit](#), [fftfit](#).

**Examples**

```
zlist <- makezlist(2,12)
zcoef <- rnorm(length(zlist))
zcoef # a vector of normalized Zernike coefficients
zcoef*zmult(zlist) # Coefficients in conventional representation
sqrt(crossprod(zcoef)) # This is the RMS error of the wavefront
# constructed from these Zernikes
```

---

zmoments

*Zernike moments*


---

**Description**

Calculate Zernike moments from a vector of coefficients

**Usage**

```
zmoments(zcoef, maxorder = 14)
```

**Arguments**

zcoef	Zernike coefficients
maxorder	Maximum order to return

**Value**

A table of the moments along with radial and azimuthal orders

**References**

M.L. Peck

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zpm	<i>Matrixes of Zernike polynomials</i>
-----	--

---

**Description**

Create a matrix of Zernike polynomial values, or their derivatives or gradient.

**Usage**

```
zpm(rho, theta, phi= 0 , maxorder = 14)
zpmC(rho, theta, maxorder)
zpm.arb(rho, theta, phi = 0, zlist = makezlist())
filldzm(rho, theta, phi = 0, zlist = makezlist())
fillgradientzm(rho, theta, phi = 0, zlist = makezlist())
```

**Arguments**

rho	A vector of radial coordinates.
theta	A vector of angular coordinates, in radians.
phi	Orientation of the image, in degrees
zlist	A list of indexes, as returned by <a href="#">makezlist</a>
maxorder	The maximum Zernike polynomial order

**Details**

rho and theta must be the same length.

**Value**

zpm.arb and filldzm return a matrix of size  $\text{length}(\text{rho}) \times \text{length}(\text{zlist}\$n)$  with values of Zernike polynomials or their radial derivatives evaluated at the polar coordinates (rho,  $\text{theta} - \pi \times \text{phi} / 180$ ).

zpm and zpmC return a matrix of size  $\text{length}(\text{rho}) \times (\text{maxorder}/2+1)^2$  of Zernike polynomial values including a piston term.

zpmC is the C++ routine that does the computations for zpm. No column names are returned.

fillgradientzm returns the gradient, in polar coordinates, of Zernikes in a  $2 \times \text{length}(\text{rho}) \times \text{length}(\text{zlist}\$n)$  matrix. Rows 1:length(rho) contain the radial derivative, followed by 1/rho times the tangential derivative.

**Note**

These are used by various routines to make least squares fits of sets of Zernike polynomials to measured wavefront values.

**Author(s)**

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**See Also**

[Zernike](#), [makezlist](#), [zlist.fr](#), [fitzernikes](#)

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