# NAME

pgmkernel - generate a convolution kernel

# SYNOPSIS

**pgmkernel** [ **–weight** w ] width [ height ]

# DESCRIPTION

Generates a portable graymap array of size *width* x *height* (or *width* x *width* if *height* is not specified) to be used as a convolution file by **pnmconvol**. The data in the convolution array K are computed according to the formula:

 $K(i,j) = 1 / (1 + w * sqrt((i-width/2)^2 + (j-height/2)^2))$ 

where w is a coefficient specified via the -weight flag, and width and height are the X and Y filter sizes.

The output PGM file is always written out in ASCII format.

### **OPTIONS**

The optional -weight flag should be a real number greater than -1. The default value is 6.0.

#### BUGS

The computation time is proportional to *width* \* *height*. This increases rapidly with the increase of the kernel size. A better approach could be using a FFT in these cases.

### SEE ALSO

pnmconvol(1), pnmsmooth(1)

### AUTHOR

Alberto Accomazzi (alberto@cfa.harvard.edu).