## Subject: Re: MORPH\_XYZ and values keyword Posted by btt on Wed, 25 Jun 2003 14:17:04 GMT

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## Karsten Rodenacker wrote:

- > Hallo, maybe some information concerning VALUE keywords. Let ste the
- > structuring element and val the values array.
- > 1. Only val\*(ste ne 0) is of importance!
- > 2. As far as I know are the val values subtracted or added from the
- > input data befor the max or min according erode or dilate is performed.
- > Hence, flat value arrays do not make sense.
- > 3. I attach two routines (for 1-d morphological filtering) for
- > illustration. gen\_sphere to generate the value array and mm\_filter to
- > apply a sequence of increasing filter steps. Unluckily the routines are
- > not documented. Maybe they are comprehensible without further words.

>

## Thanks Karsten,

This helps very much. I'm still not clear on why a flat-topped VALUES do not make sense. Here's what I get from the online docs for ERODE and DILATE. They don't look symmetric to me - so it seems like a flat-topped kernal would have a meaningful effect. But, I'm new at this and feeling like I'm on pretty thin ice.

Erode... result = min(Image) - VALUE

"Each pixel of the result is the minimum of Image less the corresponding elements of VALUE."

Dilate... result = max(Image + VALUE)

"Each pixel of the result is the maximum of the sum of the corresponding elements of VALUE and the Image pixel value. "

The Tophat filter is supposed to be...

result = (Erosion followed by Dilation) - OriginalImage

I have mocked up an 1-d example (below) using your gen\_sphere routine. It looks to me like I might be better off \*not\* using the VALUES anyway.

Just type morphtest from the command line. The top plot is the tophat without values specified. The second is with the fat-topped (the VALUES structure is shown hovering over the plot on the right.) And the last is using a non-flat-topped structure (also shown at hovering at right.)

Thanks.

Ben

```
******begin code
function gen_sphere, rad, TYPE=typ
  if not keyword_set(typ) then typ=0
  case typ of
   1:v = sqrt(rad^2-(rad-findgen(rad^2+1))^2)
   2:v = rad^2-(rad-findgen(rad^2+1))^2
   3:v = 0.05/(1.+sqrt(rad^2-(rad-findgen(rad^2+1))^2))
   4:v = 0.05/(1.+rad^2-(rad-findgen(rad^2+1))^2)
   5:v = rad-sqrt(rad^2-(rad-findgen(rad^2+1))^2)
   6:v = rad^2-(rad-findgen(rad^2+1))^2
   -1:v = replicate(1.0, rad*2+1)
   else: v = rad-abs(-findgen(rad*2+1)+rad)
  ENDCASE
  mv = max(v)
  IF mv GT (rad+1) THEN v = v/mv*(rad+1)
  return. v
end
Pro MorphTest
x = [19,19,21,23,24,23,22,21,20,19,20,20,19,17,15,
13,11,11,12,14,17,21,24,25,24,23,21,21,23,26,29,30,27,$
22,17,12,10,10,12,14,16,19,20,21,20,17,12,8,$
4,3,3,6,10,14,19,23,26,27,28,29,29,29,29,27,$
23,19,16,14,14,16,18,19,19,18,15,13,11,7,$
4,1,0,0,3,6,10,12,12,11,8,5,2,1,1,4,7,11,14,17,20,23,27]
n = 10
old p = !P
!P.CharSize = 2
!P.Multi = [0,1,3]
Dim = get Screen Size()
Window, /Free, ys = dim[1]*0.9
plot, x, title = 'orig with simple tophat'
oplot, thick = 2, $
   morph_tophat(x, [0, replicate(1,n-1), 0])
plot, x, title = 'orig with tophat values set with flat top'
values = [0, replicate(9,n-1),0]
```