
Subject: Re: Need Some Advice on Seperating Out Some Data

Posted by [rdellsy](#) on Wed, 09 Aug 2006 20:13:12 GMT

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Thanks for that. I took it, and played around with it a bit to get it to work. [Errors I found were: x and y don't concatenate in the line 'array=transpose([[x],[y]])' and I found I had to comment away the /ISOTROPIC in the plotting.) Unfortunately, it seems that cluster separates on a purely 1 dimensional basis. I tried discarding the histogram related code in favor of a much simpler system in case that was the problem, and it still didn't work. If you look at the data set I provided, the problem should be self evident.

Incidentally, I replaced everything from
h=histogram(c,reverse_indices=ri) down to the second to last line with:

```
--  
plot,x,y,psym=2  
bmax=max(array[0,*],maxsubsc)  
goodc=c[maxsubsc]  
keep=where(c[*] eq goodc)  
--
```

I feel that my code may be a tad more efficient, though I don't know how efficient the WHERE command is.

Anywho, I'm looking CLUSTER_TREE right now, which shows some more promise. If I understand it correctly, it works using distance appart, not coordinates which is a bit more useful, I think, for my problem. I'm just not sure how I can take the output and turn it into a set of clusters.

Thanks for all the help!

- Rob

JD Smith wrote:

> On Tue, 08 Aug 2006 15:20:13 -0700, rdellsy wrote:

>

>> I'm a tad confused about what you're suggesting. I'll try and work it
>> out, but I'm still fairly new to IDL, so if you could give an IDL or
>> pseudo-code example of what you're trying to explain, I would
>> appreciate. If that's too much work, I understand, and I'll just try to
>> puzzle it out on my own.

>

> You might find much of what you need in the HISTOGRAM tutorial:

>

> http://www.dfanning.com/tips/histogram_tutorial.html

>

> But before you go that route, you might first try the CLUSTER function
> in IDL (which I just read up on). Here's an example using a fake

```

> clustered data set with 5 clusters. You'll probably have to experiment
> with the number of clusters.
>
> JD
>
> tvlct,[255,0,0,0,255,255],[0,255,0,255,255,0],[0,0,255,255,0,255],1
> n_clust=5
>
> ;; Make some flake clustered data
> if n_elements(x) ne 0 then begin
>   n=1000
>   clust_fwhm=.2
>   cposx=randomu(sd,n_clust) & cposy=randomu(sd,n_clust)
>   cind=fix(randomu(sd,n)*n_clust)
>
>   x=clust_fwhm
>   fac=2*sqrt(2*log(2))
>   x=randomn(sd,n)*clust_fwhm/fac+cpox[cind]
>   y=randomn(sd,n)*clust_fwhm/fac+cpoy[cind]
> endif
>
> array=transpose([[x],[y]])
> w=clust_wts(array,N_CLUSTERS=n_clust)
> c=cluster(array,w)
> h=histogram(c,REVERSE_INDICES=ri)
> nh=n_elements(h)
>
> plot,x,y,PSYM=4,/ISOTROPIC
>
> cen=make_array(2,nh,VALUE=!VALUES.F_NAN)
> for i=0,nh-1 do begin
>   if ri[i+1] eq ri[i] then continue
>   take=ri[ri[i]:ri[i+1]-1]
>   oplot,x[take],y[take],PSYM=4,COLOR=i+1
>   cen[0,i]=[mean(x[take]),mean(y[take])]
> endfor
>
> ;; Find the lower right cluster
> void=max(cen[0,*]-cen[1,*],lrc,/NAN)
>
> ;; Highlight it
> keep=ri[ri[lrc]:ri[lrc+1]-1]
> oplot,x[keep],y[keep],PSYM=6,SYMSIZE=2
>
> END

```
