
Subject: performing multiple histograms without loops
Posted by [Jeremy Bailin](#) on Tue, 02 Feb 2010 05:25:32 GMT
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I thought it would be worth expanding on the technique that I used in response to Ed's question, because it's a very useful one. The basic idea is this:
suppose I want to use HISTOGRAM not on one a single set of N data points, but independently on multiple (say M) sets each of N data points. The simple solution is to use a for loop, but if M is large the IDL loop penalty soon becomes a problem. Is it possible to avoid a loop?

The answer is yes, and the trick is to modify the data in each of the M data sets so that they don't overlap, and then use a single HISTOGRAM on all of them at once.

For a concrete example, let's say that we have 5 data sets, each with 10 data points that are small integers (for convenience - if your data doesn't look like this but contains individual values, then you can use a combination of UNIQ and VALUE_LOCATE to turn it into this form, or just VALUE_LOCATE if you need to bin different values together).

```
set1 = [4,1,2,3,1,3,2,2,1,1]
set2 = [2,3,1,3,4,2,2,0,0,4]
set3 = [2,2,0,3,4,1,2,3,1,1]
set4 = [1,4,2,4,1,4,2,4,3,3]
set5 = [0,4,1,2,1,4,2,2,3,4]
set6 = [3,4,1,0,0,1,1,0,2,1]
datasets = [[set1],[set2],[set3],[set4],[set5],[set6]]
```

datasets is now a 10x6 array with all of the data:

```
IDL> print, datasets
```

```
   4   1   2   3   1   3   2   2
1   1
   2   3   1   3   4   2   2   0
0   4
   2   2   0   3   4   1   2   3
1   1
```

	1	4	2	4	1	4	2	4
3	3							
	0	4	1	2	1	4	2	2
3	4							
	3	4	1	0	0	1	1	0
2	1							

If we want to print out the histogram of each one, the traditional way would be
to put a HISTOGRAM command inside a for loop:

```

datasetsize=size(datasets,/dimen)
ndatasets=datasetsize[1]
minval=min(datasets)
maxval=max(datasets)
for i=0,ndatasets-1 do print, histogram(datasets[:,i], min=minval,
max=maxval)

```

0	4	3	2	1
2	1	3	2	2
1	3	3	2	1
0	2	2	2	4
1	2	3	1	3
3	4	1	1	1

But we can do it within a single histogram by adding 5 to all values in set2
so that it runs from 5 to 9, 10 to all the values in set3 so that it runs
from 10 to 14, etc.

```

datasets_new = datasets - minval
dataspan = maxval-minval+1
datasets_new += rebin(transpose(lindgen(ndatasets)*dataspan), size
(datasets,/dimen))

```

```

IDL> print, datasets_new

```

	4	1	2	3	1
3					
	2	2	1	1	
	7	8	6	8	9
7					
	7	5	5	9	
	12	12	10	13	14
11					
	12	13	11	11	
	16	19	17	19	16
19					

	17	19	18	18	
	20	24	21	22	21
24					
	22	22	23	24	
	28	29	26	25	25
26					
	26	25	27	26	

Now, if we perform a histogram of `datasets_new`, the 1s from set1 don't interfere with the 1s from set2 (which are now 6s), or the 1s from set3 (which are now 11s), etc. A single histogram will effectively perform a histogram of each set independently:

```
h_new = histogram(datasets_new, min=0, max=ndatasets*daspan-1,
bin=1)
```

But the 6 histograms are all jammed up against each other inside `h`!
How do we get them out?

```
h_new = reform(h_new, daspan, ndatasets)
```

```
IDL> print, h
```

0	4	3	2	1
2	1	3	2	2
1	3	3	2	1
0	2	2	2	4
1	2	3	1	3
3	4	1	1	1

We can compare to the loop version and see that it does indeed give the right answer.

"Alright," you say, "but the main reason I use HISTOGRAM is because of REVERSE_INDICES. How do I get those out?"

If you want the reverse indices for data set `j`, they can be easily extracted from the reverse indices of the full histogram. If we create the histogram as:

```
h_new = reform(histogram(datasets_new, min=0,
max=ndatasets*daspan-1, bin=1, $
reverse_indices=ri), daspan, ndatasets)
```

then the `i`-vector (in JD's terminology) for data set `j` runs from

ri[j*daspan] to ri[(j+1)*daspan-1]. These can be used directly to index the o-vector. To get the original element in data set j from the value in the o-vector, subtract j*daspan[0]. For example, where are the 3's in datasets[* ,1]?

```
IDL> print, ri[ri[1*daspan+3]:ri[1*daspan+3+1]-1] - 1*daspan[0]
1      3
```

A more interesting question is "where are the 3's in all of the datasets"?

This can in fact be done without loops! First, let's look at the loop version:

```
for i=0,ndatasets-1 do begin
  h = histogram(datasets[* ,i], min=minval, max=maxval,
reverse_indices=ri1)
  if h[3] gt 0 then print, ri1[ri1[3]:ri1[3+1]-1]
endfor
3      5
1      3
3      7
8      9
8
0
```

Using a combination of chunk indexing and "chunk index generation" (i.e. the solution to Wox's problem of a few weeks ago):

```
n = h_new[lindgen(ndatasets)*daspan+3]
h2=histogram(total(n>0,/CUMULATIVE,/int)-1,/
BINSIZE,MIN=0,REVERSE_INDICES=ri2)
nh=n_elements(h2)
chunkind=ri2[0:nh-1]-ri2[0]
l1=((l=lindgen(((nm=ndatasets>(max(n))))),nm) mod nm))[where((l lt $
(rebin(transpose(n),nm,nm,/sample))))]
where3 = [[chunkind],[ri[ri[chunkind*daspan+3]+l1] mod daspan[0]]]
[0]]
```

where3 is now a N3 x 2 array containing the data set and index within that data set of every value of 3 (of which there are N3=10 in this example):

```
IDL> print, transpose(where3)
0      3
0      5
```

1	1
1	3
2	3
2	7
3	8
3	9
4	8
5	0

One caveat with this method is that it can be quite wasteful of memory.

The full histogram contains $\text{dataspan} \times \text{ndatasets}$ entries, but really only ndatasets^2 of them can be non-zero. If dataspan is much larger than ndatasets , as might be the case if the values from each data set don't appear in the other data sets, then you might run into memory problems pretty quickly.

You can try to work around this a bit by using UNIQ to compress the values that get fed into the histogram, but it makes it much more complicated to extract the original information back out.

-Jeremy.
