Subject: Re: Array sorting by row Posted by JD Smith on Tue, 22 Aug 2006 20:48:59 GMT

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On Fri, 18 Aug 2006 11:12:36 -0700, Ed Hyer wrote:

- > Try this one out for size (or rather, for speed):
- > a = [[4,2,0,5],[9,0,1,5],[0,4,2,1],[1,2,3,4]]; or any 2-d array
- > sz=size(a,/dimensions); get dimensions
- > cols=rebin(indgen(sz[0]),sz[0],sz[1]); label points by column
- > h2=hist_nd(transpose([[a[*]],[cols[*]]]),[1,1],reverse_indic es=ri2)
- > ; syntax is a wee messy, but this is JD Smith's beautiful HIST_ND()
- > function:
- > ; http://www.dfanning.com/documents/programs.html#HIST_ND
- > starti=n_elements(h2)+1; beginning of indices in \$RI2
- > sorted=transpose(reform(a[ri2[starti:*]],sz[0],sz[1]))

>

> This one should be blazing fast right up to memory limits.

Interesting method. However, a (usual sort of) problem occurs if your input array is sparse, e.g.:

```
a=[[4,2,0,5],[9,0,1,5],[0,4,2,1],[1,21212121212L,3,4]]
```

In this case, you will quickly fill up memory with countless zeroes in a nearly empty 2D histogram.

A related method which avoids this sparseness issue, and allows arbitrary floating point numbers, is as follows:

```
sz=size(a,/DIMENSIONS)
s=sort(a)
h=histogram(s mod sz[0],REVERSE_INDICES=ri)
sorted_inds=transpose(reform(s[ri[sz[0]+1L:*]],sz))
sorted=a[sorted_inds]
```

Basically, you sort the entire array, and then bin the sorted list by column number using HISTOGRAM. The histogram is guaranteed to contain only as many bins as you have columns in your input array, which is a small number even for very large arrays. This means the algorithm offers a constant runtime depending only on array size.

There's one other variant possible. If you know in advance your values are not sparsely distributed (e.g. at least 1 in 10 number are represented, on average), you can see further speedup (depending on your memory resources) by changing:

s=sort(a)

in the above prescription to

h=histogram(a,REVERSE_INDICES=ri) s=ri[n_elements(h)+1:*]

i.e. just sort directly using HISTOGRAM. This really has no business being this much faster than SORT, but it is. Here are some timings comparing the three methods:

; 100x100 array, on average every number represented (1 in 1 sparse)

HIST ND method: 0.17189288

SORT+HISTOGRAM method: 0.0094909668 HISTOGRAM+HISTOGRAM method: 0.0018260479

; 500x500 array, 1 in 1 sparse

HIST_ND method: 10.859160

SORT+HISTOGRAM method: 0.37013221
HISTOGRAM+HISTOGRAM method: 0.17653203

; 500x500 array 10 in 1 sparse (numbers repeated 10 times on average)

HIST ND method: 1.3364871

SORT+HISTOGRAM method: 0.37795591 HISTOGRAM+HISTOGRAM method: 0.10262513

; 500x500 array, 1 in 10 sparse (histogram only 10% filled)

HIST_ND method: <array too large to allocate>

SORT+HISTOGRAM method: 0.35415411 HISTOGRAM+HISTOGRAM method: 0.35962415

And, just to show that in some small part of parameter space the first two are similar.

; 2000x2000 array, 100 in 1 sparse (numbers repeated 100 times on average)

HIST_ND method: 11.669180

SORT+HISTOGRAM method: 10.190833 HISTOGRAM+HISTOGRAM method: 2.4765849

And to really puts a hurt on the pure HISTOGRAM based methods, by cranking up the sparseness (HIST_ND is already out of its league):

; 1000x1000 array, 1 in 5 sparse

SORT+HISTOGRAM method: 2.0481181 HISTOGRAM+HISTOGRAM method: 1.2417412

; 1000x1000 array, 1 in 10 sparse

SORT+HISTOGRAM method: 2.0289030 HISTOGRAM+HISTOGRAM method: 1.6590791 ; 1000x1000 array, 1 in 50 sparse

SORT+HISTOGRAM method: 2.0890758 HISTOGRAM+HISTOGRAM method: 4.4074631

; 2000x2000 array, 1 in 50 sparse

SORT+HISTOGRAM method: 10.333820

HISTOGRAM+HISTOGRAM method: <array too large to allocate>

So there you have it. These results are highly reminiscent of an old post giving the various trade-offs using SORT and HISTOGRAM for list matching and set operations, see http://www.dfanning.com/tips/set_operations.html. The advice offered is so similar, I'll quote myself:

"So, if you want a generic solution which works in the same n log(n) time using a fixed amount of memory for any type of integer input data, sparse or not, use SORT. If you know your data is non-sparse (better than 1 in 10 say), you can see a speedup of a few with HISTOGRAM."

JD