Subject: Re: array concatenation and optimization Posted by Brian Jackel on Wed, 26 Sep 2001 21:29:19 GMT

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Hi

```
This is what I generally do
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```
dat= {struct, a:fltarr(3), b:intarr(4), c:0b} ;example structure
OPENR, lun, filename, /GET LUN
READU, lun, dat ; read one record
fs= FSTAT(lun); look at current file information
nrec= fs.size/fs.cur_ptr
IF (fs.size GT 100000000L) THEN BEGIN ;avoid out-of-memory errors
 FREE LUN, lun
 MESSAGE, 'Error- file larger than 100Mbytes, returning'
ENDIF
POINT LUN, lun, 0
data= REPLICATE(dat,nrec)
READU, lun, data : read all the data at once
FREE LUN, lun
```

I'm not sure how well this will work with compressed files.

Brian

```
Sean Raffuse wrote:
>
> Hello.
> I am trying to read a bunch of data from a file to a structure array. I'm
> not sure many data entries the file will have until I have read it and so I
> am increasing the size of the structure array after reading each line. I do
> this by concatenating.
>
> adp_struct_single is the structure as a "scalar"
> adp struct
                     is the array
  I concatenate like so:
>
     adp struct = [adp struct, adp struct single]
>
 This is working but it has increased the processing time of my loop by an
> order of magnitude. Is there a better way to do this? Is there a reason
  this is so slow?
> Thanks in advance.
>
```

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