Subject: Re: Array subscript for VECTOR must have same size as source expression

Posted by Rohit Deshpande on Sun, 26 Jun 2011 14:22:47 GMT

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On Jun 24, 3:27 pm, Paolo <pgri...@gmail.com> wrote:
> Well,
>
> this is a typical errors that you will encounter many times
 in vour idl career.
>
  The line
  all_barytimes[i,*] = im.BARYTIME
 tries to assign a variable im.BARYTIME into an array all_barytimes,
>
  specifically into a particular column of the array.
 The error specifies that im.BARYTIME does not fit into
> all barytimes[i,*]
> (likely because it has a different number of elements and/or
> dimensions).
 Here's an example showing the problem
> IDL> a=fltarr(4,4)
> IDL> b=fltarr(5)
> IDL> a[1,*]=b
> % Array subscript for A must have same size as source expression.
> % Execution halted at: $MAIN$
> see? can't fit the 5 elements of b into a column of a.
>
> Ciao.
> Paolo
  On Jun 24, 11:27 am, Rohit Deshpande <singlebin...@gmail.com> wrote:
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   Hello Everyone,
>>
>> I am a beginner in IDL and I have been working on a project. I explain
>> it below:
```

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>
>> 1. The Idea: Read a bunch of FITS files in IDL. They have structure so
>> I use MRDFITS. I would like to read each one of them in a separate
>> variable and plot them. Given that each file has X by Y dimension,
>> where X is always = 19 while Y changes but is mostly 1636.
>> 2. The Code:
>> im = fltarr(n)
>> all barytimes = dblarr(19,4000)
>> all_normflux = dblarr(19,4000)
  FOR i = 0, n-1 DO BEGIN
         ; where filenames are the list of fits files I am reading
>>
>> it.
>>
        im = mrdfits(file+'raw_test/'+string(filenames[i]),1,head)
        all barytimes[i,*] = im.BARYTIME
>>
        all_normflux[i,*] = im.AP_CORR_FLUX
>> ENDFOR
>> 3. The Error:
>> IDL> lcs1
>> % READCOL: 21 valid lines read
>> MRDFITS: Binary table. 19 columns by 1639 rows.
>> % Array subscript for ALL BARYTIMES must have same size as source
>> expression.
>> % Execution halted at: LCS1
>> Please let me know how to make it work.
>> Thanks!
Thank you so much. I have now created a double array outside the FOR
loop in the following way. This help and I have no errors:
all_barytimes = dblarr(n,4450)
all normflux = dblarr(n, 4450)
FOR i = 0, n-1 DO BEGIN
im = mrdfits(file+'raw test/'+string(filenames[i]+'.fits'),1,head)
arrayend=n_elements(im.BARYTIME)-1
  all_barytimes[i,0:arrayend] = im.BARYTIME
all_normflux[i,0:arrayend] = im.AP_CORR_FLUX
```