Subject: Re: Specification for a new array slicing function Posted by steinhh on Thu, 20 May 1999 07:00:00 GMT

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In article <374317CC.E1AC89EA@ssec.wisc.edu> Liam Gumley <Liam.Gumley@ssec.wisc.edu> writes:

```
> Please find below a suggested specification for a new array slicing
> function, formatted as a standard IDL prolog. The intention here is to
> provide a means to extract n-dimensional array 'slices' from an existing
> array in memory. The caller can choose to skip elements along any or all
> dimensions if desired.
> Comments are invited. There's no code yet, so now is the time.
>
[..]
 : INPUT KEYWORD PARAMETERS:
                  Set this keyword to a vector containing the start
> :
     START
[..]
     STRIDE
                  Set this keyword to a vector containing the
[..]
> ;
    COUNT
                  Set this keyword to a vector containing the
[..]
> : EXAMPLE:
> ; ;Extract every other element along each dimension
> ; array = findgen(1, 10, 5, 6, 7)
> ; ndims = size( array, /n dimensions )
> ; stride = replicate( 2L, ndims )
> ; result = array_slice( array, stride=stride )
> ; help, result
>
> ;;RESULT
                   FLOAT
                             = Array[1, 5, 2, 3, 3]
```

IMO, the use of keyword parameters for START, STRIDE and COUNT is a bit "wordy" for my liking. And these items are really essential to the routine as such. So why not use positional parameters?

For something that really ought to be a part of the IDL syntax, I would also like a shorter name (despite the possibility for name conflicts), like "arex", short for array_extract.

My suggestion would be something a bit more like the native Fortran 9X syntax (not that I actually *know* exactly how that syntax works!), e.g.:

a(0:5:2,:,5:9) would be translated into

```
arex(a,[0,5,2],-1,[5,9])
```

I.e., each positional parameter signifies the extraction operation for one array dimension. There are some issues that I would like to clear up, though: What exactly does the 0:5:2 sequence mean? Does it mean elements 0:5, sampled with a stride of 2? Or does it mean 5 elements sampled with a stride of 2, starting from 0? Or is it START:STRIDE:COUNT, meaning 2 elements, sampled with a stride of 5?

Just curious.... And I would strongly recommend following Fortran conventions, whatever they are....

Anyway, the three elements in each parameter appear in "optionality" order: start [, stride [, count]] (if that's what the syntax is supposed to be).

Looking at the example above, you may wonder what the "-1" is doing there... Well, the idea is that one could use a nonnegative *scalar* parameter to signify extraction of a slice at a given position, whilst -1 really means "*", in IDL notation.

Since we now have one extra "degree of freedom" in that a start position (and nothing else) may be specified in two similar ways, as e.g. 0 or [0]... I have great use for this (since we're at now at liberty to rewrite the rules..:-) I've always disliked the way this works:

```
a = fltarr(5,5,5)
surface,a(*,3,*)
```

% SURFACE: Array must have 2 dimensions: <FLOAT Array[5, 1, 5]>.

I mean - if I'm extracting an "image" out of a "cube", why would I want the last dimension to stick around...???

So, I would like to be able to say

```
surface, arex(a, -1, 3, -1)
```

with no error messages! On the other hand, if I do want the dangling dimension, I could specify it:

```
surface, arex(a, -1, [3], -1)
```

(and get the error message :-)

Or the other way around, if people feel strongly about leaving this dimension in.....

I would also like to see a corresponding index function, returning the one-dimensional indices to the extracted elements instead of the elements themselves. This could be used for assignments. I.e.:

$$a(arexi(a,-1,[3],[0,2])) = data_block$$

Just some thoughts...

Regards,

Stein Vidar