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Subject: Re: For-loop vs. Dimensional Juggling relative performance

Posted by [Gray](#) on Thu, 11 Feb 2010 17:22:01 GMT

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On Feb 10, 10:51 pm, Jeremy Bailin <astroco...@gmail.com> wrote:

> On Feb 9, 9:54 pm, Gianguido Cianci <gianguido.cia...@gmail.com>  
> wrote:

>  
>  
>

>> On Feb 8, 10:26 pm, Gray <grayliketheco...@gmail.com> wrote:

>

>>> Hi folks,

>

>>> I recently wrote my own version of SRCOR from the NASA Astrolib. Just  
>>> as a reminder, the program takes two lists of 2D coordinates and finds  
>>> matches where the distance is less than some cutoff. SRCOR uses a for-  
>>> loop to step through the first list, comparing the distance of each  
>>> coordinate-pair from every point in the second list. My version uses  
>>> matrix multiplication and dimensional juggling to avoid the for-loop.

>

>>> For  $n_1 = 2143$  and  $n_2 = 2115$ , SRCOR is faster (0.16 seconds to my 0.53  
>>> on my macbook); however, for  $n_1 = 25$  and  $n_2 = 26$ , mine is faster  
>>> ( $1.8e-4$  seconds to  $4.2e-4$ ). Is there any way to predict what kind of  
>>> list sizes will be faster with each method, without making some random  
>>> data and using brute force?

>

>>> The relevant code is:

>

>>> SRCOR (dcr2 is the cutoff, option eq 2 ignores the cutoff) -->

>

```
>>> FOR i=0L,n1-1 DO BEGIN
>>>   xx = x1[i] & yy = y1[i]
>>>   d2=(xx-x2)^2+(yy-y2)^2
>>>   dmch=min(d2,m)
>>>   IF (option eq 2) or (dmch le dcr2) THEN BEGIN
>>>     ind1[nmch] = i
>>>     ind2[nmch] = m
>>>     nmch = nmch+1
>>>   ENDIF
>>> ENDFOR
```

>

>>> My code -->

>

```
>>> lkupx = rebin(indgen(n1),n1,n2)           ;make index lookup
>>> tables, so as not to
>>> lkupy = rebin(transpose(indgen(n2)),n1,n2) ;worry about confusing
>>> 1D vs 2D
```

```

>>> ;use matrix multiplication and dim. juggling to fast compute
>>> sqrt((x2-x1)^2+(y2-y1)^2)
>>> dists =
>>> sqrt(rebin(x1^2.+y1^2,n1,n2)+rebin(transpose(x2^2.+y2^2),n1, n2)-2*(x1#x2+y1 #y2))
>>> min_x = min(dists,xmatch,dimension=2) ;find the minima in both
>>> directions...
>>> min_y = min(dists,ymatch,dimension=1) ;this is given in 1D indices
>>> xm = lkupy[xmatch] ;convert to 2D indices
>>> ym = lkupx[ymatch]
>>> ;remove elements w/ distance greater than max_dist, and where the
>>> two lists don't match
>>> nomatch_x = where(ym[xm] ne indgen(n1) or min_x gt max_dist, nmx)
>>> if (nmx gt 0) then xm[nomatch_x] = -1
>>> nomatch_y = where(xm[ym] ne indgen(n2) or min_y gt max_dist, nmy)
>>> if (nmy gt 0) then ym[nomatch_y] = -1
>
>>> Thanks!!
>>> --Gray (first time poster)
>
>> Gray, have you tried the inbuilt DISTANCE_MEASURE ? I'd be curious to
>> know if it's any faster.
>
>> --Gianguido
>
> I'd wager that JD's match_2d will knock the socks off both of those...
>
> -Jeremy.

```

Oy... wish I'd known about match\_2d before I spent so much time on mine. Yes, it kicks both routines' collective butt.