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Subject: Re: chunk indexing like

Posted by [Wout De Nolf](#) on Sun, 10 Jan 2010 13:52:14 GMT

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On Sat, 9 Jan 2010 01:37:18 -0800 (PST), chris <[rogass@googlemail.com](mailto:rogass@googlemail.com)> wrote:

```
> h=histogram(total(n>0,/CUMULATIVE,/int)-1,/  
> BINSIZE,MIN=0,REVERSE_INDICES=ri )  
> nh=n_elements(h)  
> chinkind=ri[0:nh-1]-ri[0]  
> ind2=where([1,chinkind[1:]-chinkind[0:nh-2]] ne 0)  
> l1=lindgen(nh)-ind2[chinkind]
```

The code below solves the histogram approach error but even when excluding the histogram from timing (I need chunk indexing anyway), the loop approach is the fastest.

```
h=histogram(total(n>0,/CUMULATIVE,/int)-1,/BINSIZE,MIN=0,REVERSE_INDICES=ri)  
nh=n_elements(h)  
chinkind=ri[0:nh-1]-ri[0]
```

```
t0=systime(1)  
tmp=[1,chinkind[1:]-chinkind[0:nh-2]]  
ind2=where(tmp ne 0)  
tmp=total((tmp-1)>0,/cum,/int)  
l2=lindgen(nh)-ind2[chinkind-tmp]  
print,'Histogram-approach: ',systime(1)-t0,' seconds for  
,n_elements(n), ' elements'
```

Timing:

CR-Rebin/mod-approach: 1.4690001 seconds for , 10000 elements  
Histogram-approach: 2.3290000 seconds for , 10000 elements  
Loop-approach: 0.82800007 seconds for , 10000 elements

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