

## GROMACS - Bug #157

### Index selection by group name

07/17/2007 01:53 PM - Anonymous

<b>Status:</b> Closed	
<b>Priority:</b> Normal	
<b>Assignee:</b> Erik Lindahl	
<b>Category:</b> analysis tools	
<b>Target version:</b> 3.3.1	
<b>Affected version - extra info:</b>	<b>Difficulty:</b> uncategorized
<b>Affected version:</b>	
<b>Description</b>	
<p>When selecting an index group by using its name the routine that parses the input does not catch an error where it should. See:</p> <pre>Just take any pdb you like, get an index file echo q   make_ndx -f x.pdb  editconf -n -f x.pdb</pre> <p>and select, e.g. "MyBackbone", or "FIRST_sol", or "Protein_Part1", then editconf will not report an Error, but select backbone, SOL or Protein.</p> <p>Within a larger script this can cause nasty bugs as you don't get an error (Which happened to me).</p> <p>Cheers from Goettingen, Jochen</p>	

### History

#### #1 - 07/18/2007 11:17 AM - David van der Spoel

Easy bugs are nice :)  
On line 644 in index.c replace  
strstr(string,s)  
by  
strstr(s,string)

fixed it in CVS as well.