

## GROMACS - Feature #1029

### -tu option in g\_rmsf

11/05/2012 04:36 PM - Francesco Oteri

<b>Status:</b> Closed	
<b>Priority:</b> Low	
<b>Assignee:</b>	
<b>Category:</b>	
<b>Target version:</b>	
<b>Difficulty:</b> uncategorized	
<b>Description</b> Good morning, I think it is useful adding the option -tu in g_rmsf. Nowadays people are simulating for several nanoseconds so using yet the picosecond notation is, in my opinion, very annoying. Thank you Francesco	
<b>Related issues:</b> Is duplicate of GROMACS - Feature #1030: adding -tu option to some analysis t... <b>Fix uploaded</b>	

#### History

**#1 - 01/07/2014 02:16 PM - Rossen Apostolov**

- Project changed from 9 to GROMACS

**#2 - 05/26/2014 11:01 AM - Rossen Apostolov**

- Is duplicate of Feature #1030: adding -tu option to some analysis tools added

**#3 - 06/01/2014 10:10 AM - David van der Spoel**

- Status changed from New to Closed