

## GROMACS - Feature #3045

### Option to remove v-sites in trjconv?

07/19/2019 04:42 PM - Cathrine Bergh

<b>Status:</b>	New	
<b>Priority:</b>	Low	
<b>Assignee:</b>		
<b>Category:</b>		
<b>Target version:</b>		
<b>Difficulty:</b>	uncategorized	
<b>Description</b>		
Could there be an option to remove v-sites in trjconv? E.g. that v-sites are removed, if they exist, when option 2) Protein-H is selected or to have a separate option for that. I think many people would find this convenient.		