

ProCKSI - Bug #3

Structrues excluded from calculation when not enough atoms

11/26/2007 04:15 PM - Anonymous

Status:	Closed	Start date:	
Priority:	High	Due date:	
Assignee:	Anonymous	% Done:	0%
Category:	ProCKSI	Estimated time:	0.00 hour
Target version:			
Resolution:	fixed		
Description			
<p>When structures do not have enough selected atoms, no .sa file is generated. Nor are .dm, .cm, .ac, .rc, and .cn files generated which are use by USM and [[MaxCMO]].</p> <p>=> These methods do give similarity matrices for the full dataset.</p> <p>=> SSMs cannot be combined.</p> <p>Possible solution (pre-processing):</p> <ul style="list-style-type: none">- Check for minimum number of structures, atoms, selected atoms before request is submitted <p>Possible solution (post-processing):</p> <ul style="list-style-type: none">- Enter all values into database- Generate SSMs on-the-fly- Correct for missing data- Combine SSMs on-the-fly			

History

#1 - 11/23/2007 04:34 PM - Anonymous

- Status changed from New to Closed
- Resolution set to fixed

(In r505) f - New similarity comparison method: Vorolign
e - New design with better navigation
f - Customisable visualisation for contact maps as images and graphs
f - Customisable notification emails for finished experiments/tasks
f - Improved data security with authentication
e - Improved scheduling of all tasks

closes #3, #5, #2, #35, #42, #24, #26, #52, #32, #11, #40, #22

#2 - 11/26/2007 04:15 PM - Anonymous

(In r506) f - New similarity comparison method: Vorolign
e - New design with better navigation
f - Customisable visualisation for contact maps as images and graphs
f - Customisable notification emails for finished experiments/tasks
f - Improved data security with authentication
e - Improved scheduling of all tasks

closes #3, #5, #2, #35, #42, #24, #26, #52, #32, #11, #40, #22

