

Issues

#	Project	Tracker	Status	Priority	Subject	Assignee	Updated	Target version
2984	GROMACS	Feature	New	Normal	More Extensive Selections Examples	Dallas Warren	06/18/2019 11:50 PM	
2260	GROMACS	Bug	Closed	Normal	grompp running out of memory when excluded bonded neighbours is large		01/03/2018 02:02 PM	
2289	GROMACS	Feature	New	Normal	gmx spatial add ability to perform over multiple reference structures		11/08/2017 12:31 AM	
1602	GROMACS	Feature	New	Low	gmx order S per slice along all 3 axes		09/22/2014 07:25 AM	
1619	GROMACS	Feature	Closed	Low	gangle add x and y axis to second vector group options	Teemu Murtola	03/05/2015 10:26 AM	5.1
3107	GROMACS	Bug	Closed	Normal	editconf not updating atom names correctly	Mark Abraham	09/27/2019 02:33 PM	2019.4
2951	GROMACS	Bug	Closed	Normal	Core dump with gmx rdf with some selections		06/11/2019 02:57 PM	2019.3