share/template/gromacs/selmethod.h										
src/gmxlib/selection/compiler c src/gmxlib/selection/evaluate c src/gmxlib/selection/params c src/gmxlib/selection/paraetree c	src/gmxlib/selection/scanner_internal.c. src/gmxlib/selection/selection.c.	src/gmxlib/selection/selelem.c. src/gmxlib/selection/selbeln.c.	elp.c src/gmxlib/selection/selmethod.c src/gmxlib/selection	selection/sm_compare.c src/gmxlib/selection/sm_distance.c	src/gmxlib/selection/sm_insolidangle.csrc/gmxlib/selection/sm_keywords	rds.c src/gmxlib/selection/sm merge.c s	src/amxlib/selection/sm_permute.c	.c/gmxlib/selection/sm_position.c src/gmxlib/sc	election/sm same.c src/gmxlib/selection/sm simple.c	src/gmxlib/traiana/traiana.c
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