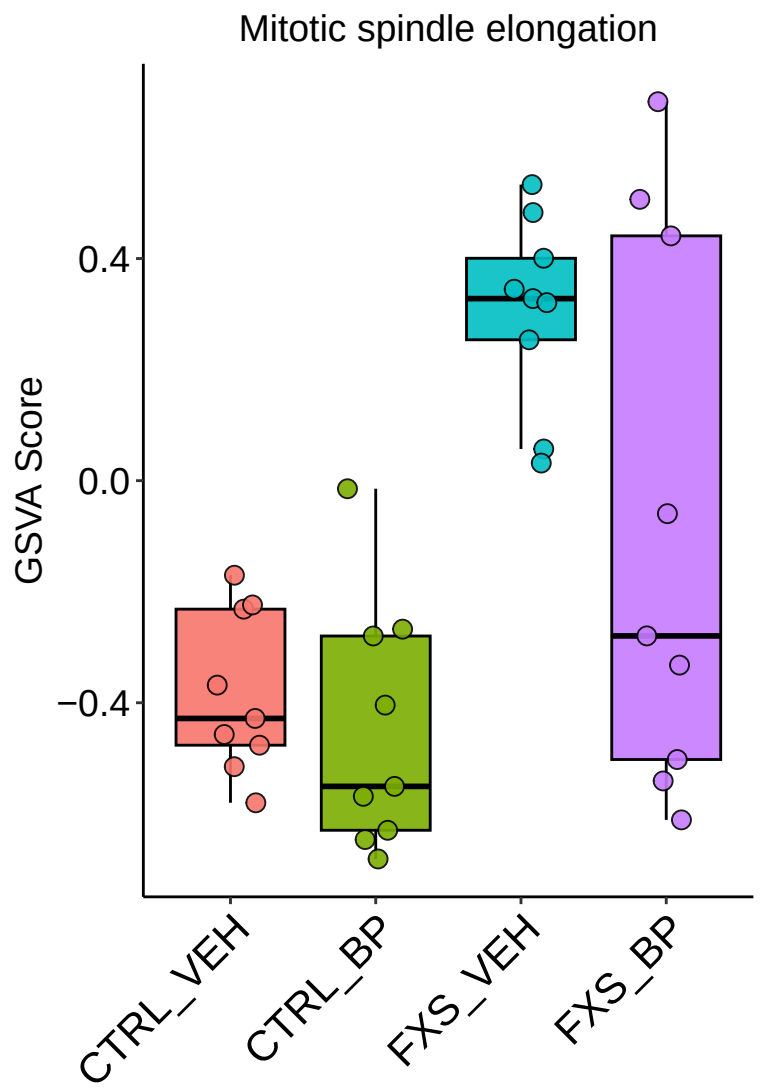
Mitotic spindle elongation
GSVA Score
0.4
0.0
−0.4
CTRL_VEH
CTRL_BP
FXS_VEH
FXS_BP