

# ESCRT I complex

E-MAP score

6  
4  
2  
0  
-2

R108G H141R R108A R108L R108Y T34Q R108Q Y157A K143W R108I T34L T34A D79S F58A T34E K101R WT A180T K132H G80A T34G Q147E R78K

Gsp1 mutant

complex member ● SRN2 ● STP22

