

● mutation
■ Protein Domain

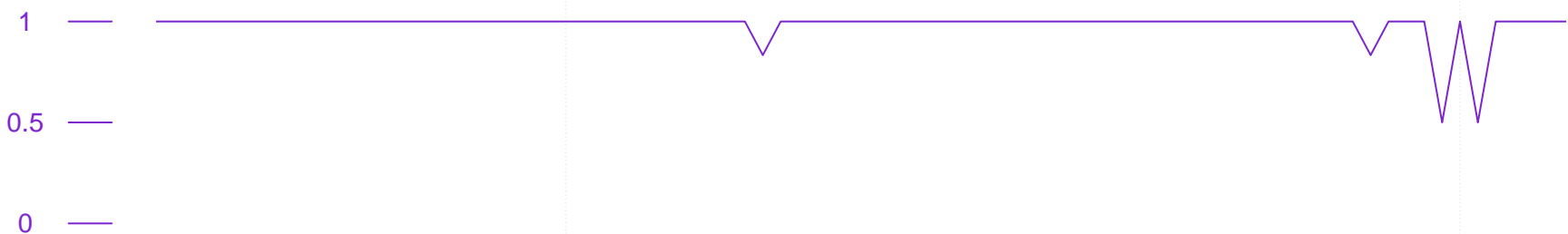
Reference

(KTRSMLLSSEQLKLCVLEYQGKY I LKVCGCDEYFLEKYPLSQYKY I RSC I MLGRMPNLMLMAKESLYSQLPMDCTMP

PIK3CA



Conservation



Score



250

300

Amino Acid Position