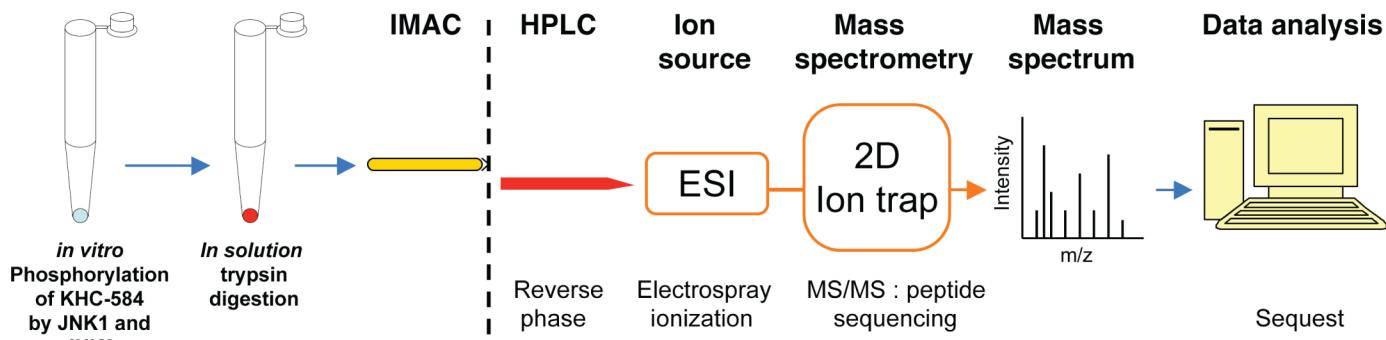


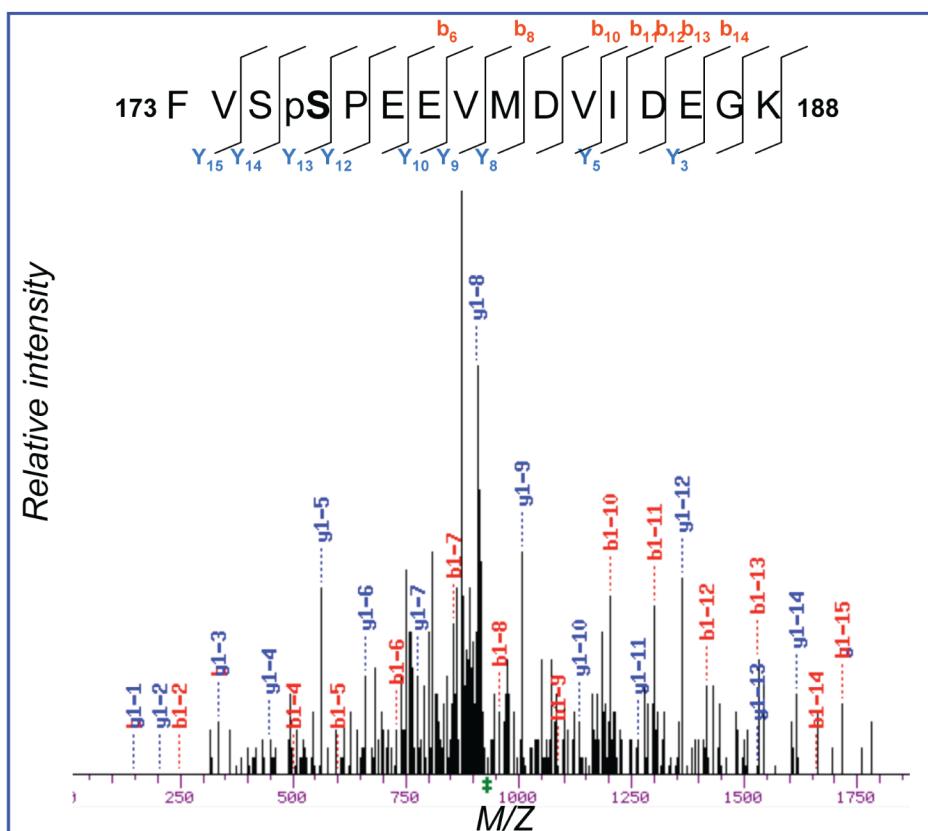
A**B****KHC-584**

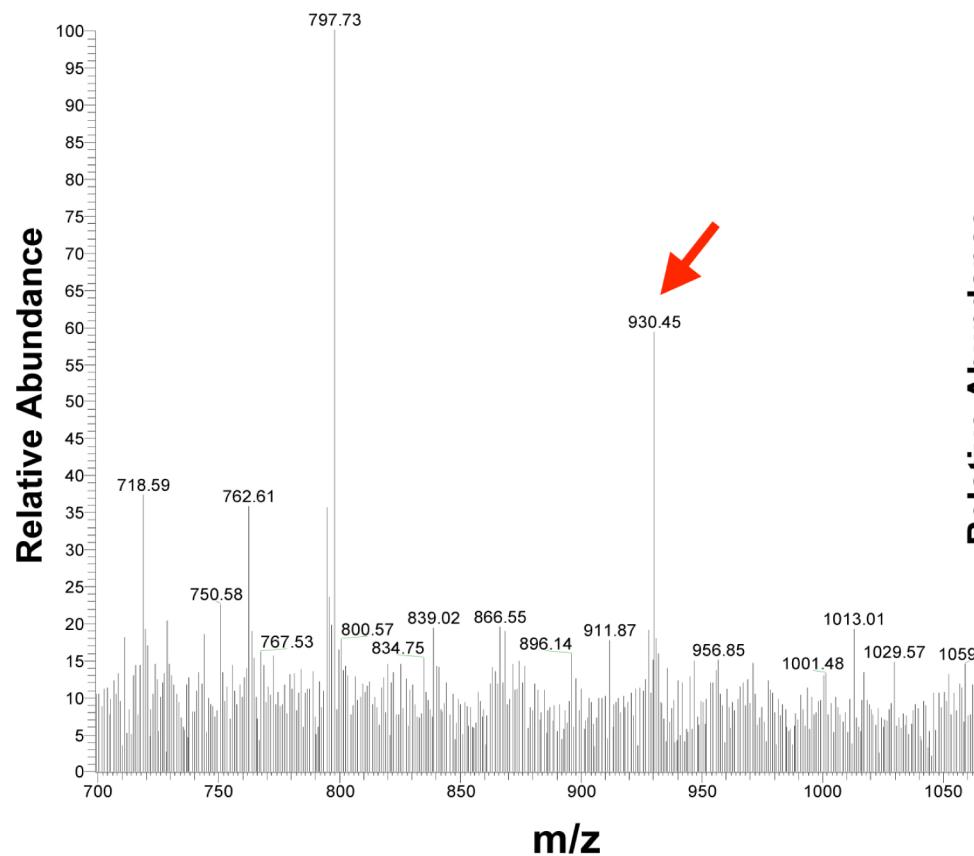
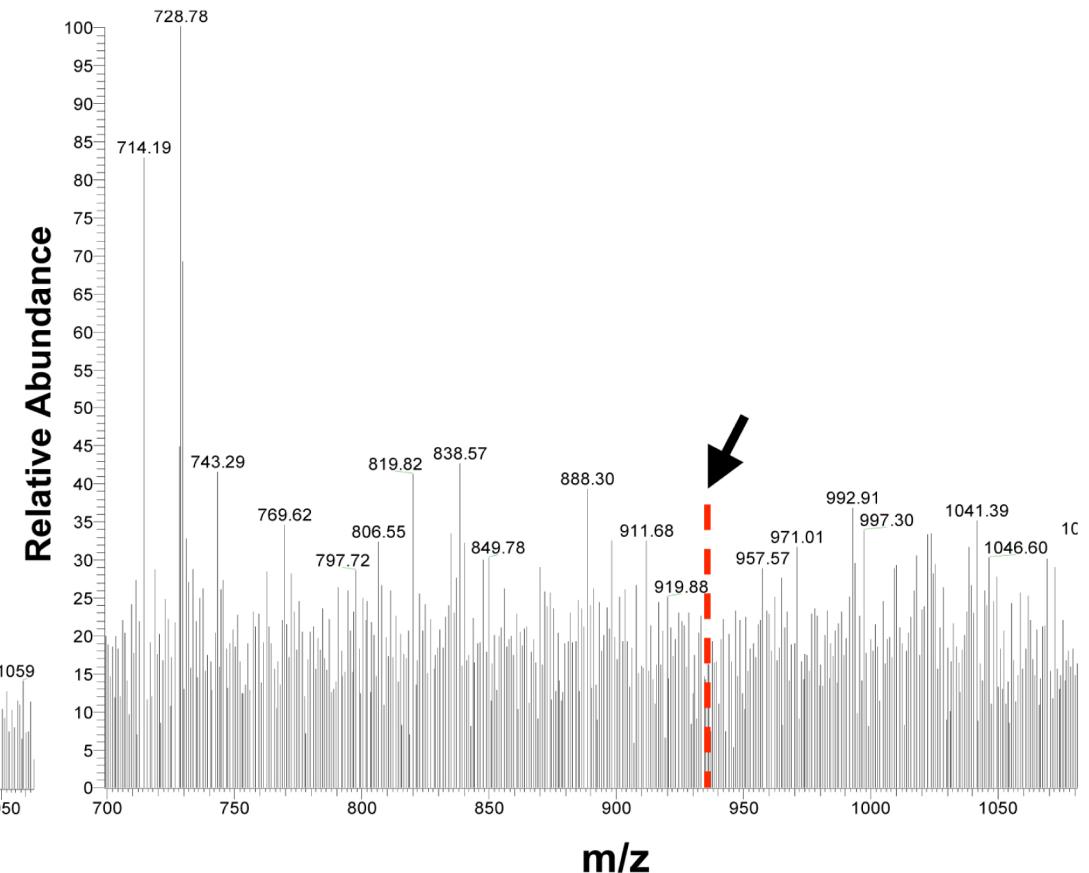
HHHHHHMADPAECSIKVMCRFRPLNEAEILRGDKFIRKFKGEETVVGQGKPYVFDRVLPPNTTQEQQVYNACAKQIVKDLEGYNGTIFAYGQTSSGKTHTMEGKHDPQLMGIIPRIAHDFDHIYSMDENLEFHVKSYFEIYLDKIRDLLDVSKTNLAVHEDKNRVPYVKGCTERFVSS*PEEVMDVIDEGKANRHVAVTNMNEHSSRSHSIFLINIKQNVETEKKLSGKLYLVDLAGSEKVSKTGAEGAVLDEAKNINKSLSALGNVISALAEGTKTHVPYRDSKMTRILQDSLGGNCRTTIVICCSPSVFNEAETKSTLMFGQRAKTIKNTVSVNLELTAAEEWKKKYEKEKEKNKALKSVIQHLEVELNRWRNGEAVPEDIQISAKDHKSLEPCDNTPIIDNITPVVDGISAEEKEKYDEEITSLYRQLDDKDEINQQSQLAEKLKQQMLDQDELLASTRDYEKIQEELTRLQIENEAKDEVKEVLQALEELAVNYDQKSQEVEDKTRANEQLTDELAQKTTTQRELSQLQELSNHQKKRATEILNLLKDLGEIGGIIGTNDVKTLADVNGVIEEEF

Protein coding for the first 584 aa of kinesin1-c from rat. Coverage of the 584 aa fragment is 72%

C

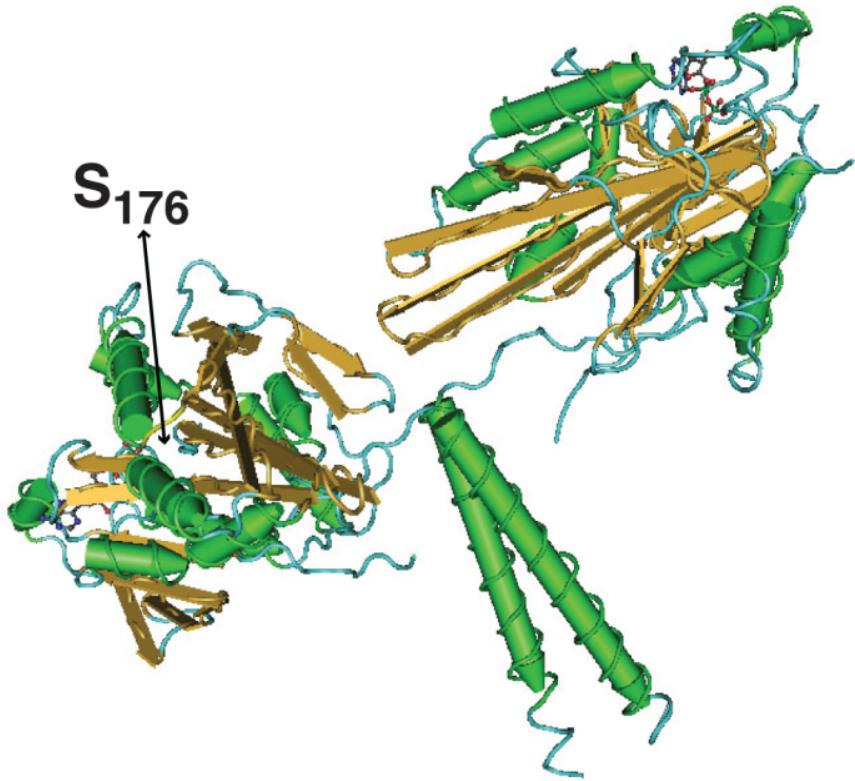
Ion charge (a)	MH+ (ion mass) (b)	Xcorr (c)	dCn (d)	Sequence (e)
2	1862.0 (-1.4)	2.8069	0.395	R.FVSS*PEEVMDVIDEGK.A

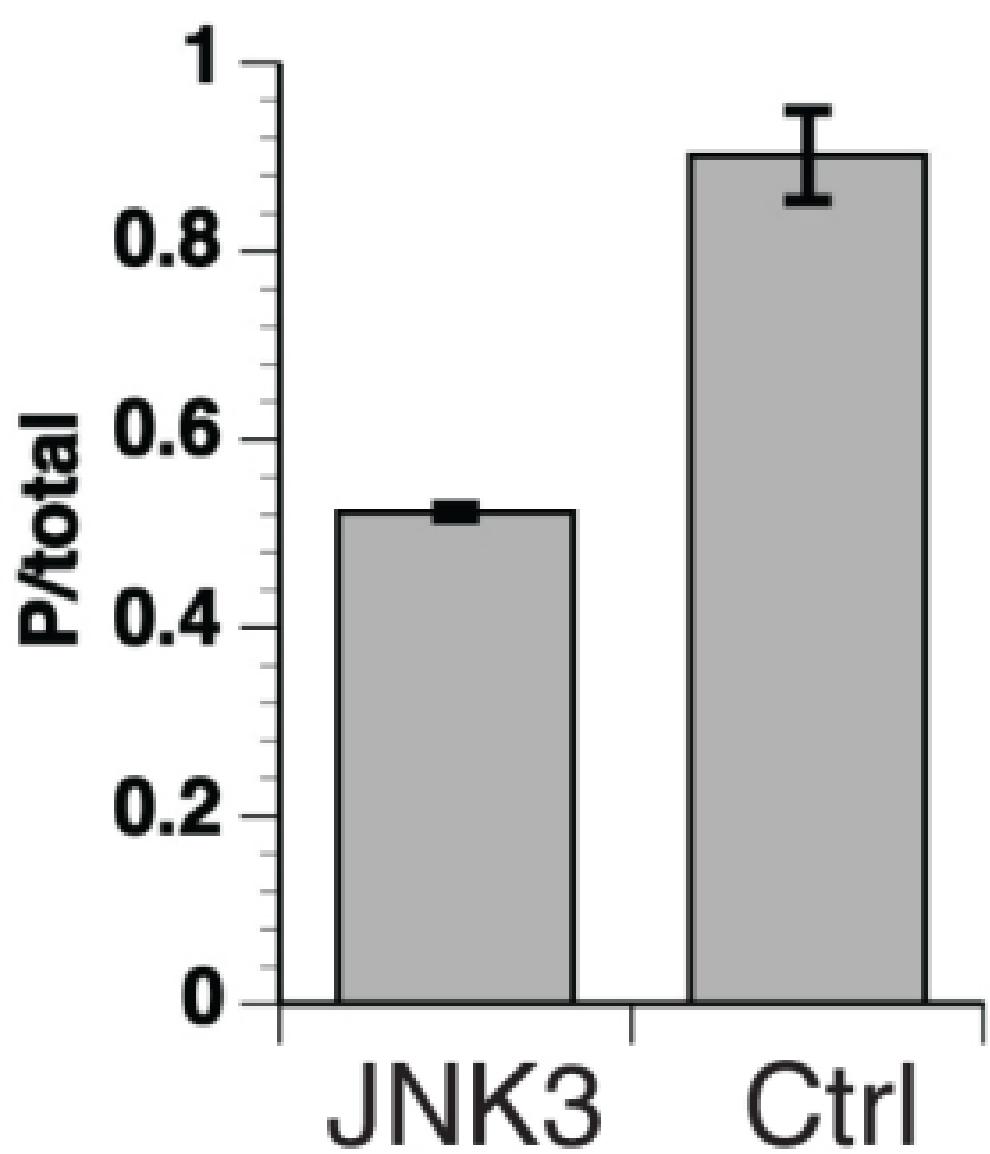


A**JNK3: Full ms RT: 58.44****JNK1: Full ms RT: 58.44**

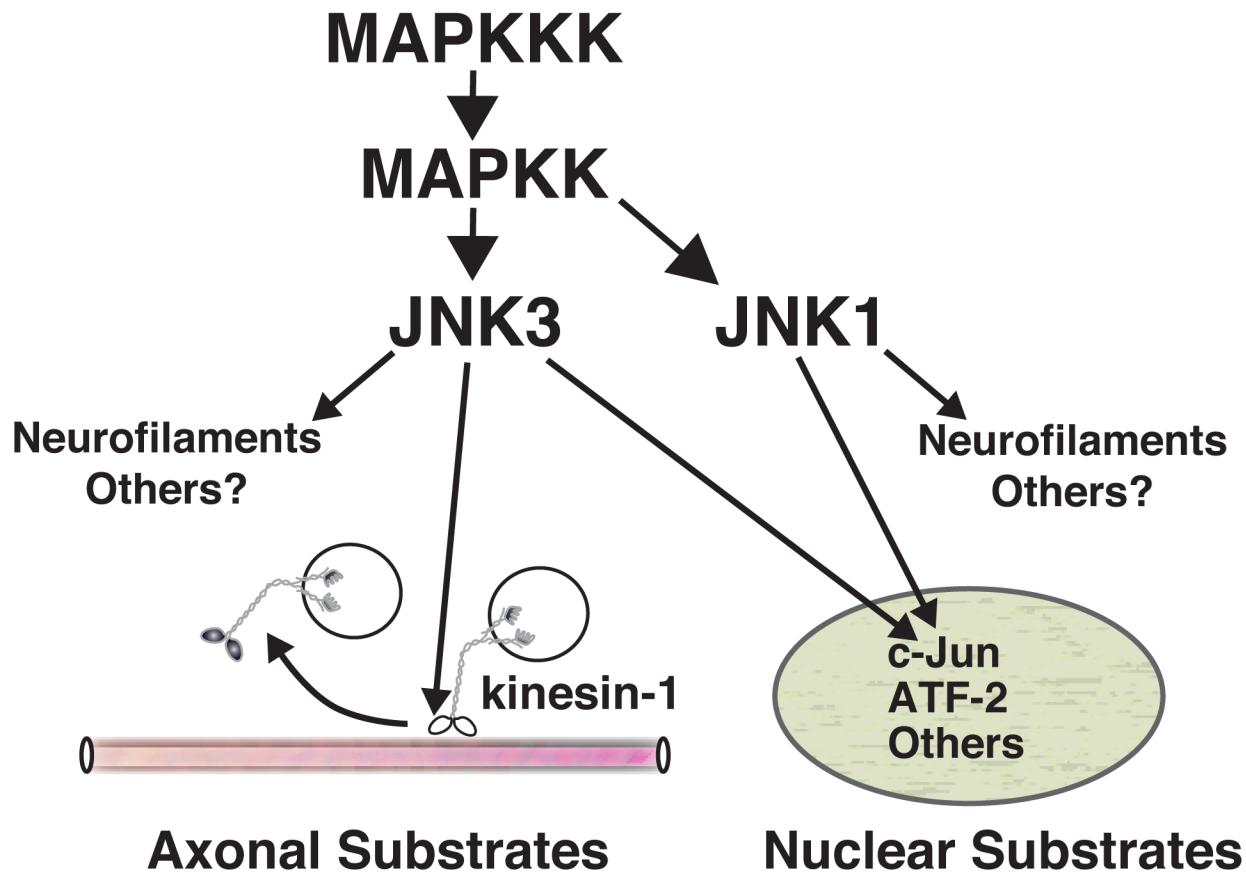
A

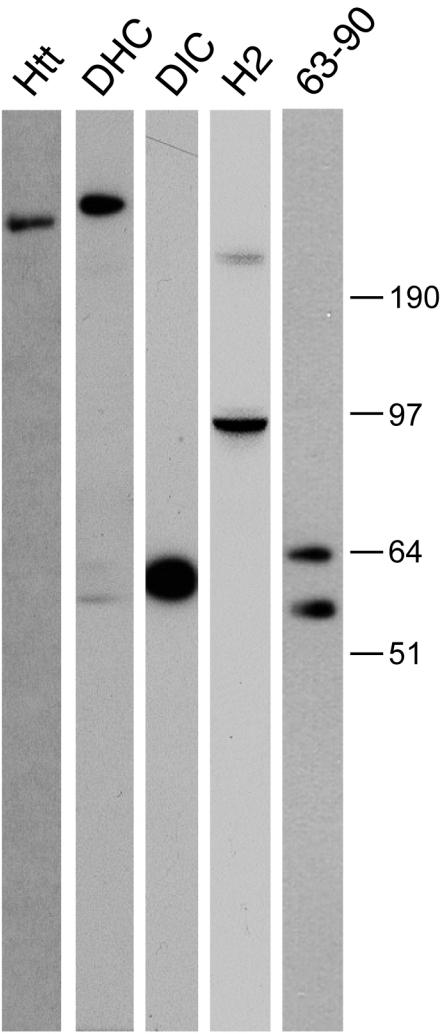
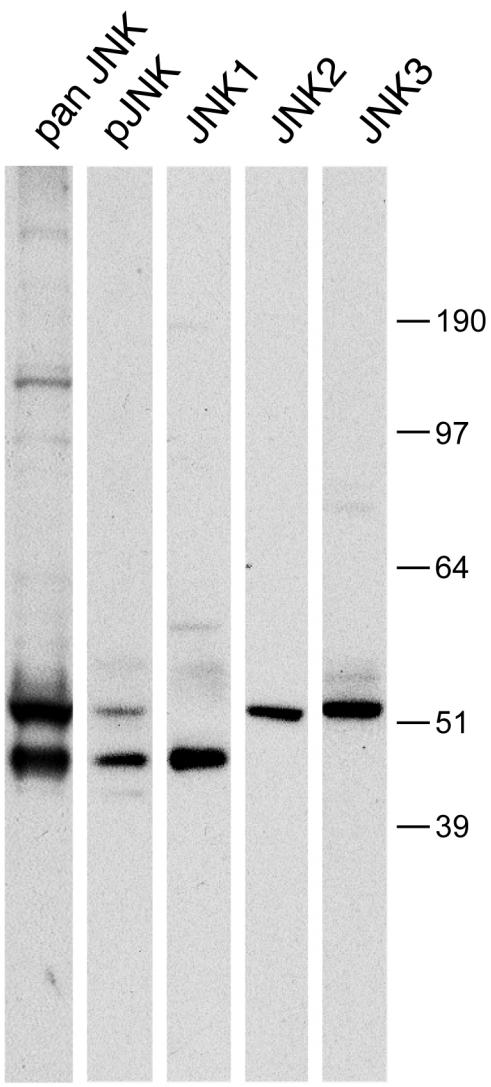
Homo 173-FVSS^{*}SPEEVMDVIDEGK-188
Mus 173-FVSSSPEEVMDVIDEGK-188
Loligo 173-FVSSSPEEVMEVIDEGK-188

B



PolyQ-Htt



A**B****C**