<u>GKIKYKSEQAENYRKLILATAEDPRVILL</u>KLSDRLDNVKTLWVFREEKRKKIAKETMEIY SPOT AQUAE HNKTRSIKEANTISKMFFAMTHDIRIIIIKLADKLHNMTTLSYLPKNRUDRIAKDCLSTY SPOT BORBU KERDKKEA QAENERKMIMAMVODIRVILIKLADRTHNMRTLGSLRPDKRRRIARETLEIY SPOT ECOLI KERTROEAQVENERKMILAMTROIRVVLIKLADRTHNMRTLGSLRPDKRRRIAKETLEIY SPOT HAEIN LKNKKENLNLKSFVNIAINSQQEINVMVLKLADRLDNIASIEFLPIEKQKVIAKETLELY SPOT MYCGE LNRKKEDLNLKSLVNI AMSSQQEVNALVLKLADRLDNISSIEFLAVEKOKI I AKETLELY SPOT_MYCPN AKENRTOIKAQYLRKLYLSMAKDIRVIIVKLADRLHNLKTIGYLKPERQOIIARESLEIY SPOT SPICI NESSTTEHQAENERRMELAMAKDIRVIVVKLADRLHNMRTLDALSPEKORRIARETKDIF SPOT SYNY3 APLAHRLGVWSIKNELEDWAFKYLYPEEYEKVRNFVKESRKNLEE SPOT_AQUAE VPIAERLGISSLKTYLEDLSFKHLYPKDYKEIKNELSETKIEREK SPOT_BORBU SPLAHRLGIHHIKTELEELGFEALYPNRYRVIKEVVKAARGNRKE SPOT ECOLI CPLAHRLGIEHIKNELEDLSFQAMHPHRYEVLKKLVDVARSNRQD SPOT_HAEIN AKIAGRIGMYPVKTKLADLSFKVLDLKNYDNTLSKINKOKVFYDN SPOT MYCGE AKIAGRIGMYPYKTOLADLSFKYLDPKNFNNTLSKINOOKVFYDN SPOT MYCPN SAIAHRLGMKAVKQETEDISFKIINPVQYNKIVSLLESSNKEREN SPOT SPICI APLANRLGIWREKWELEDLSEKYLEPDSYRKIQSLVVEKRGDRES SPOT SYNY3