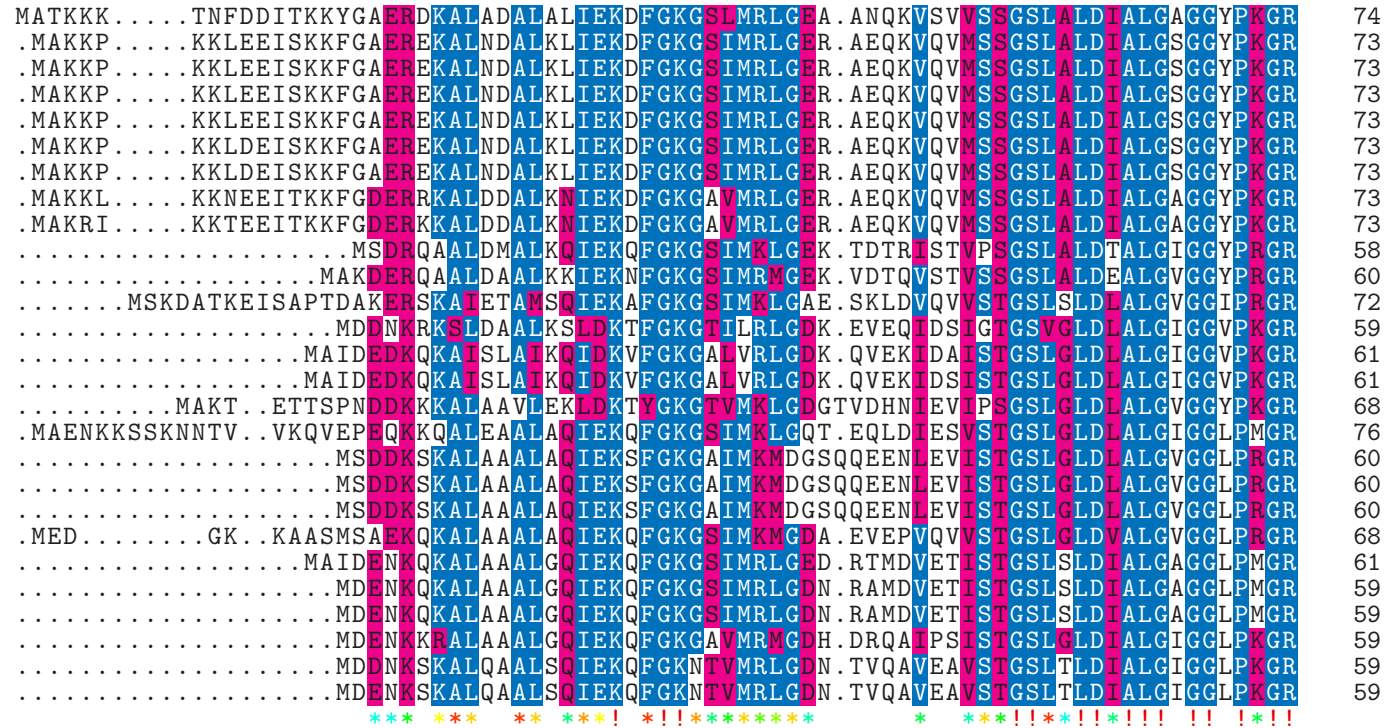


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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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consensus

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DEKIGGGSENAKYLAEHPFIDIDKQVRSKFLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE 385

DEKIGGGSENAKYLAEHPFIDIDKQVRSKFLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE 385

DEKIGGGSENAKYLAEHPFIDIDKQVRSKFLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIE 385

DEKIGGGSENAKYLADNPVFAEIDHQVRVQFLIDGEEAAE...SKKDE...VAQVDS..VNEEVTLDLGDELEIE 381

DEKIGGGSENAKYLADNPVFAEIDHQVRVQFLIDGEEAAE...SKKDE...AAQVDS..VNEEVTLDLGDELEIE 381

GEKIGGGSENAKRYLADHPQIFIDIRKVRVKFGLLEESEESAMAVASE...E.....TDDLALDLNGIEIE 377

GEKIGGGSENAKKFLADNPFIIDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE 382

EERLGGGRENAKQFLKENKDIMLMIQEIQIREHYGLDMNGVVQQQAEEETQEEL.EFE.E..... 348

SERIGGGRENAKNYLADHEDVEDEVRLKVR AAYCISDVPEEDLPTTEDE.QI.NILPDDST....EE..... 355

DERIGGKEKTIAYIAERPEMEQEIHDRVMAAIRAGNAGEAPALAPAPAA.PEAAEA..... 363

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DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEIMPLPDEP..LE.EME..... 347

DTKLGQGRD TVKELIKDNPELAEELEAKIREKLQAK..... 343

GTKIGGGKNNAVKWLKENPEIADKLEQEIRNTLASNPNSVLKATAS..D...SNDES DVEEFSEEE..L..... 372

GAKIGGGKNVRVWLKENPEISDEIDAKTRALNGVEMHITEGTQDE..TD.....GERPEE..... 348

GAKIGGGKNVRVWLKENPEISDEIDAKTRALNGVEMHITEGTQDE..TD.....GERPEE..... 348

GAKIGGGKNVRVWLKENPEISDEIDAKTRALNGVEMHITEGTQDE..TD.....GERPEE..... 348

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GDKIGGGKANACKYLKENPEIAKTLDKKLREMLLNPNEMQLI..AE.....TSSAADDVEFGAVPEEF..... 354

GSKIGGGKANAAKYLEENPEIQQEIEQQIRAKLLVTSANTKA..TP..VA.EDLADADI..... 347

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GNKIGGGKNNVIRYLEENPQIAEIERNIREQLLTGTNGAV..Q...I..EDEEPPDLLES..... 349

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NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349

consensus

 non conserved
 similar
 ≥ 50% conserved