```
logo                                          MKHWFFLIILFFMNCSWGQDPFDKTQRNRSQFDNAQTVMEQTEIISSDVPNNLCGADENRQAAEIPLNALKLVGVVISKD

NTDB id 1151768 LSO74 RS03465 WP 005656464.1  MKYWFFLIILFFMNCSWGQDPFDKTQRNRSQFDNAQIVMEQTEIISSDVPNNLCGADENRQAAEIPLNALKLVGVVISKD   80
NTDB id 1346 HI 0436 AAC22095.1               MKHWFFLIILFFMNCSWGQDPFDKTQRNRSQFDNAQTVMEQTEIISSDVPNNLCGADENRQAAEIPLNALKLVGVVISKD   80
NTDB id 1372 NTHI RS02650 WP 011272047.1      MKHWFFLIILFFMNCSWGQDPFDKTQRNRSQFDNAQTVMEQTEIISSDVPNNLCGADENRQAAEIPLNALKLVGVVISKD   80
consensus                                     !!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo                                          KAFALLQDQGLQYYSVLEGVDVAQEGYIVEKINQNNVQFMRKLGEQCDSSEWKKLSF

NTDB id 1151768 LSO74 RS03465 WP 005656464.1  KAFALLQDQALQVYSVLEGVDVAQEGYIVEKINQNNVQFMRKLGEQCDSSEWKKLSF   137
NTDB id 1346 HI 0436 AAC22095.1               KAFALLQDQGLQVYSVLEGVDVAQEGYIVEKINQNNVQFMRKLGEQCDSSEWKKLSF   137
NTDB id 1372 NTHI RS02650 WP 011272047.1      KAFALLQDQGLQIYSVLEGVDVAQEGYIVEKINQNNVQFMRKLGEQCDSSEWKKLSF   137
consensus                                     !!!!!!!!!*!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved