```
logo
NTDB id 310098 CH602 RS03715 WP 014550482.1  MIYFTMFLLGGILGIALWFYLSGFITRLQQEIYATYIELFPQNFLSFQPSVSSIQQKKCGHILRYFLNVGVEFIFLQIAF   80
NTDB id 1357 HI 0296 AAC21960.1              MIYFTMFLLGGILGIALWFYLSGFITRLQQNIYAIYVELFPQNRSPFQPHFASIQQKKCGHILRYFFSIGVGFIFLQIAF   80
NTDB id 1368 NTHI RS01940 WP 011271970.1     MIYFTMFLLGGILGIALWFYLSGFITHLQQEIYATYVELFPQNSSPFQPHFASIQQKKCGHILRYFFSIGVGFIFLQIAF   80
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!*!!!*!*!!!!!! **!!!***!!!!!!!!!!!!!!!***!!*!!!!!!!!

logo
NTDB id 310098 CH602 RS03715 WP 014550482.1  KDSTFAIWLGLTLLILWTISYLDWHYQLISTTLCLWLLTIGLFGADNNFSLLTLSESIKSAASFFIVFYAIYWIAKFYYG  160
NTDB id 1357 HI 0296 AAC21960.1              KDSIFTVWIGLTLIILWTISYLDWHYQLISTTPCLWLLTLGLFGADNNFSLLTLSESIKSAASFFIVFYVIYWLAKFYYG  160
NTDB id 1368 NTHI RS01940 WP 011271970.1     KDSIFTVWIGLTLIILWTISYLDWHYQLISTTPCLWLLTLGLFGADNNFSLLTLSESIKSAASFFIVFYAIYWIAKCYYR  160
consensus                                    !!!*!**!*!!!!*!!!*!!!!!!!!!!!!!!!*!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!*!!*!!*

logo
NTDB id 310098 CH602 RS03715 WP 014550482.1  KEAFGRGDYWLAMALGSFIHLETLPHFLLLASVLGICFSLIHKKKKEFIPFAPFMNLSAVIIYFVKYYGY  230
NTDB id 1357 HI 0296 AAC21960.1              KEAFGRGDYWLAMALGSFIHLETLPHFLLLASVLGICFSLIHRKKKEFLPFAPFMNLSAVIIYFVKYYGY  230
NTDB id 1368 NTHI RS01940 WP 011271970.1     KEAFGRGDYWLAMALGSFIHLETLPHFLLLASVLGICFSLIHKKKKEFIPFAPFMNLSAIIIYLVKYYGY  230
consensus                                    !!!!!!!!!!!!!!!!!!!!
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