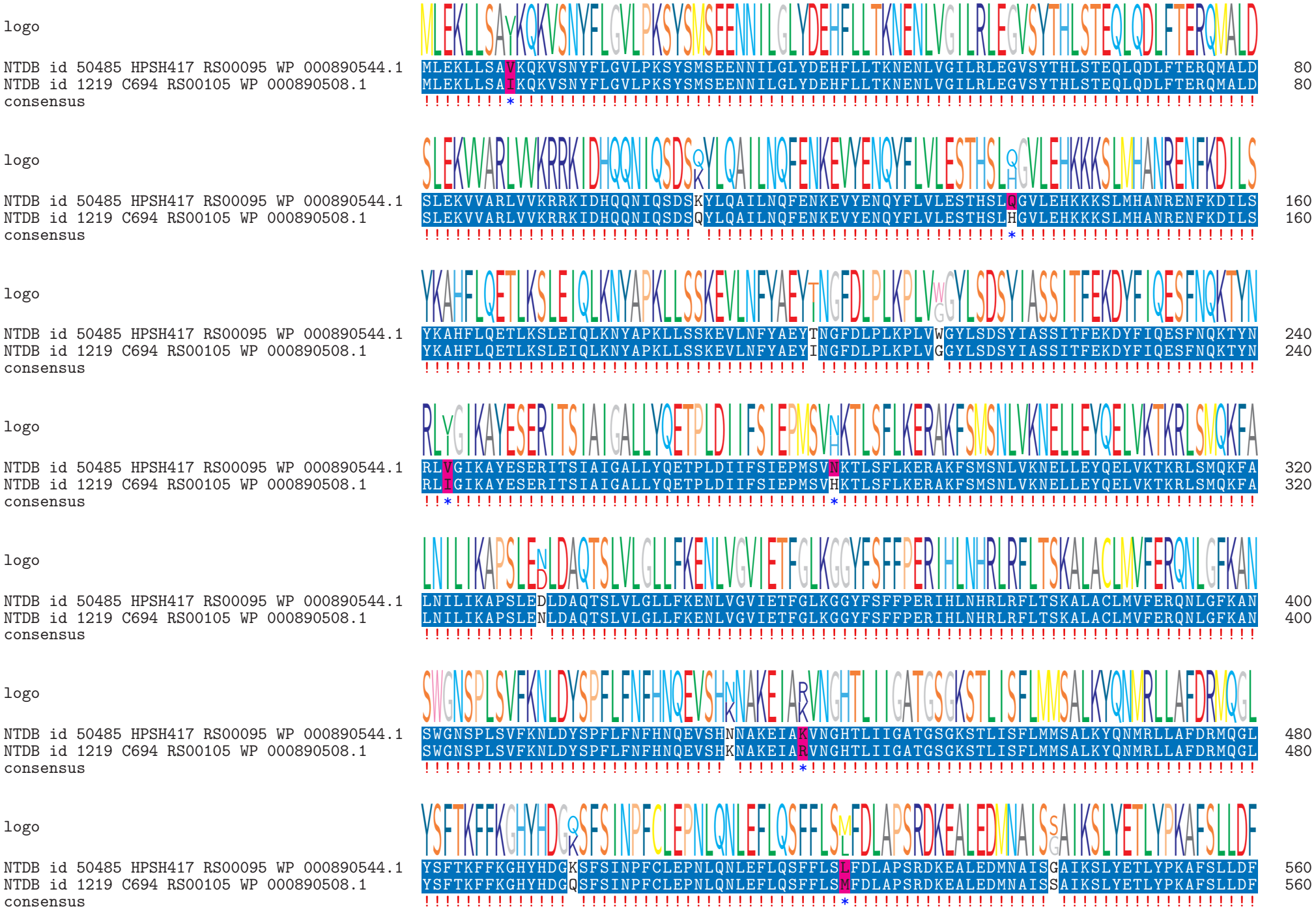
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consensus
logo
NTDB id 50485 HPSH417 RS00095 WP 000890544.1 SLEKVVARLVVKRRKIDHQQNIQSDSKYLQAILNQFENKEVYENQYFLVLESTHSLQGVLEHKKKSLMHANRENFKDILS 160
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consensus
logo
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consensus
logo
NTDB id 50485 HPSH417 RS00095 WP 000890544.1 RLVGIKAYESERITSIAIGALLYQETPLDIIFSIEPMSVNKTLSFLKERAKFSMSNLVKNELLEYQELVKTKRLSMQKFA 320
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consensus
logo
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NTDB id 1219 C694 RS00105 WP 000890508.1 LNILIKAPSLENLDAQTSLVLGLLFKENLVGVIETFGLKGGYFSFFPERIHLNHRLRFLTSKALACLMVFERQNLGFKAN 400
consensus
logo
NTDB id 50485 HPSH417 RS00095 WP 000890544.1 SWGNSPLSVFKNLDYSPFLFNFHNQEVSHNNAKEIAKVNGHTLIIGATGSGKSTLISFLMMSALKYQNMRLLAFDRMQGL 480
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consensus
logo
NTDB id 50485 HPSH417 RS00095 WP 000890544.1 YSFTKFFKGHYHDGKSFSINPFCLEPNLQNLEFLQSFFLSLFDLAPSRDKEALEDMNAISGAIKSLYETLYPKAFSLLDF 560
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consensus

logo

```
NTDB id 50485 HPSH417 RS00095 WP 000890544.1  KETLKRTSSNQLGLSLEPYLNNPLFNALNDAFNSNAFLNVINLDTITQNPKDLGLLAYYLFYKILEESRKNDSGFLVFLD  640
NTDB id 1219 C694 RS00105 WP 000890508.1      KETLKRTSSNQLGLSLEPYLNNPLFNALNDAFNSNAFLNVINLDTITQNPKDLGLLAYYLFYKILEESRKNDSGFLVFLD  640
consensus                                     !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

logo

```
NTDB id 50485 HPSH417 RS00095 WP 000890544.1  EFKSYVENDLLNAKINALITQARKANGVVVLALQDIYQLSGVKNAHSFLSNMGTLILYPQKNARELKHHFNVPLSETEIS  720
NTDB id 1219 C694 RS00105 WP 000890508.1      EFKSYVENDLLNTKINALITQARKANGVVVLALQDIYQLSGVKNAHSFLSNMGTLILYPQKNARELKHNFNVPLSETEIS  720
consensus                                     !!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!
```

logo

```
NTDB id 50485 HPSH417 RS00095 WP 000890544.1  FLENTPLYARQVLVKNLGNGNSNMIDVSLESLGRYLKIFNSDSSHVNKVKALQKDYPTEWREKLLKS   787
NTDB id 1219 C694 RS00105 WP 000890508.1      FLENTPLYARQVLVKNLGNGSSNMIDVSLESLGRYLKIFNSDSSHVSKVKALQKDYPTEWREKLLRS   787
consensus                                     !!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!*!
```

X non conserved

X similar

X ≥ 50% conserved