Test Alignment

```
1aboA      .NLFVALYDFVASGDNTLSITKGE.......KLRVLGY..      30
1ycsB      KGVIYALWDYEPQNDDELPMKEGD.......CMTIIHR..      31
1pht       GYQYRALYDYKKEREEDIDLHLGDILTVNKGSLVALGFSD      40
1vie       ........DRVRKKSGA..AWQGQIVGWYCTNLTPEGYAV      30
1ihvA      NFRVY....YRDSRD...PVWKGP......AKLLWKGEGA      27
consensus       **  **      *         *          *     *
```
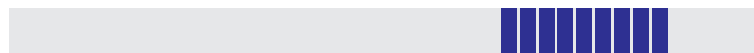
TC

```
1aboA      ...NHNGE..WC..EAQTKNGQGWVPSNYITPVN......      57
1ycsB      ...EDEDEIEWW..WARLNDKEGYVPRNLLGLYP......      60
1pht       GQEARPEEIGWLNGYNETTGERGDFPGTYVEYIGRKKISP      80
1vie       ESEAHPGSVQIY.............PVAALERIN......      51
1ihvA      VVIQDNSDIKVV.............PRRKAKIIRD.....      49
consensus       **  *              *    *          *
```
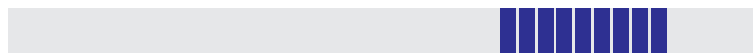
TC

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Reference Alignment

```
1aboA       .NLFV.ALYDFVASGDNTLSITKGEKLRV.......LGYN       31
1ycsB       KGVIY.ALWDYEPQNDDELPMKEGDCMTI.......IHRE       32
1pht        .GYQYRALYDYKKEREEDIDLHLGDILTVNKGSLVALGFS       39
1vie        .........DRVRKKSG..AAWQGQIVGW.........YC       20
1ihvA       ......NFRVYYRDSRD..PVWKGPAKLL.........WK       23
consensus         **  **              *
```

```
1aboA       HNG.......EWCEA..QTKNGQGWVPSNYITPVN.....       57
1ycsB       DEDEI.....EWWWA..RLNDKEGYVPRNLLGLYP.....       60
1pht        DGQEARPEEIGWLNGYNETTGERGDFPGTYVEYIGRKKIS       79
1vie        TNLTP....EGYAVESEAHPGSVQIYPVAALERIN.....       51
1ihvA       GEG.......AVVIQ...DNSDIKVVPRRKAKIIRD....       49
consensus             *          * **      *
```

```
1aboA       .    57
1ycsB       .    60
1pht        P    80
1vie        .    51
1ihvA       .    49
consensus
```

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