```
logo
NTDB id 89051 Pcarn RS11420 WP 261833990.1  MSMDNTAAKVEDLAQNSDEMEVASVELE.........RVVEEPEVKEEFDASAKNLDATQLYLGEIGFSPLLTAEEEVLYARRAL   76
NTDB id 1183 GTF74 RS11590 WP 000116737.1   MSVSNTVTKVEEFDFEDEALEVLETDAELTSDEELVAVEGASEDVREEFDASAKSLDATQMYLSEIGFSPLLTAEEEVLYARRAL   85
consensus                                   !!* !!  !!!*     * *!!   * !*********     **!*!!!!!!!! !!!!!*!! !!!!!!!!!!!!!!!!!!!!!
```

```
logo
NTDB id 89051 Pcarn RS11420 WP 261833990.1  RGDEAARKRMIESNLRLVVKISRRYSNRGLALLDLIEEGNLGLIRAVEKFDPERGFRFSTYATWWIRQTIERALMNQTRTIRLPI   161
NTDB id 1183 GTF74 RS11590 WP 000116737.1   RGDEAARKRMIESNLRLVVKISRRYSNRGLALLDLIEEGNLGLIRAVEKFDPERGFRFSTYATWWIRQTIERALMNQTRTIRLPI   170
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

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logo
NTDB id 89051 Pcarn RS11420 WP 261833990.1  HVVKELNIYLRTARELSQKLDHEPTAEEIAQQLDKPVDDVSKMLRLNERVSSVDTPIGGDGDKALLDIIPDINNSDPEVSTQDED   246
NTDB id 1183 GTF74 RS11590 WP 000116737.1   HVVKELNIYLRTARELSQRLDHEPTPEEIALELDRPVDDVTKMLRLNERISSVDTPIGGDGDKALLDILPDSHNADPEFSTQDDD   255
consensus                                   !!!!!!!!!!!!!!!!!!*!!!!!!*!!!!  !!*!!!!!*!!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!! *! !!! !!!!*!
```

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logo
NTDB id 89051 Pcarn RS11420 WP 261833990.1  IKHSLIDWLDELNPKQKEVLARRFGLLGYEPSTLEEVGREINLTRERVRQIQVEGLRRLREILLKQGLNMESLFNFDSE.   325
NTDB id 1183 GTF74 RS11590 WP 000116737.1   IRESLLNWLDELNPKQKEVLARRFGLLGYEPSTLEEVGREINLTRERVRQIQVEGLRRLREILVKQGLNMEALFNVEYDN   335
consensus                                   !**!!* !!!!!!!!!!!!!!!!!!!!
```