

T-COFFEE, Version\_6.85 Tue Sep 9 14:03:25 WEST 2008

Cedric Notredame

CPU TIME:0 sec.

SCORE=77

\*

BAD AVG GOOD

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gi | 2132225 : 83  
gi | 7304901 : 82  
gi | 6978535 : 83  
gi | 5097870 : 83  
gi | 9168085 : 79  
gi | 1337789 : 74  
gi | 3256363 : 57  
cons : 77

gi | 2132225 1 ELGERQLVHVVKWAKALPGFRNLHVDDQMAVIQYSWMGLMVFAMGWRS 48  
gi | 7304901 1 ELGERQLVHVVKWAKALPGFRNLHVDDQMAVIQYSWMGLMVFAMGWRS 48  
gi | 6978535 1 ELGERQLVHVVKWAKALPGFRNLHVDDQMAVIQYSWMGLMVFAMGWRS 48  
gi | 5097870 1 ELGERQLVHVVKWAKALPGFRNLHVDDQMAVIQYSWMGLMVFAMGWRS 48  
gi | 9168085 1 ELGERQLVYVVVKWAKALPGFRNLHVDDQMSIQYSWMGLMIFAMGWRS 48  
gi | 1337789 1 ELGERQLVKVVVKWAKGLPGFRNLHVDDQMTVIQHTWMGVMVFALGWRS 48  
gi | 3256363 1 KSIHQQLVLFIEWAKSLPQFSFLAQADQAALLKGAASIIVLGVAYRS 48  
cons 1 : . : \* \* \* . : : \* \* \* . \* \* \* \* \* \* \* \* : : : : . : : : : : : \* \* 48

gi | 2132225 49 FTNVNSRMLYFAPDLVFN EYRMHKS RMYS - QCVRMRHLSQ --- EFGWL 92  
gi | 7304901 49 FTNVNSRMLYFAPDLVFN EYRMHKS RMYS - QCVRMRHLSQ --- EFGWL 92  
gi | 6978535 49 FTNVNSRMLYFAPDLVFN EYRMHKS RMYS - QCVRMRHLSQ --- EFGWL 92  
gi | 5097870 49 FTNVNSRMLYFAPDLVFN EYRMHKS RMYS - QCVRMRHLSQ --- EFGWL 92  
gi | 9168085 49 FTNVNSRMLYFAPDLVFN EYRMHKS RMYS - QCVRMRLQLSQ --- EFGWL 92  
gi | 1337789 49 YKNANARMLYFAPDLVFN DRRMHVSSMYE - HCVQMKHLSQ --- EFVLL 92  
gi | 3256363 49 ICLTVENTICLANDTL LP -- KEHATQVGD INCVVGR I IDE IVN PMRRL 94  
cons 49 : . . : : \* \* : : : : \* : : : . : \* \* : : : : : \* 96

gi | 2132225 93 QITPQEF LCMKALLLFSIIPVDGLKNQKFFDEL RMNYIKELDR II --- 137  
gi | 7304901 93 QITPQEF LCMKALLLFSIIPVDGLKNQKFFDEL RMNYIKELDR II --- 137  
gi | 6978535 93 QITPQEF LCMKALLLFSIIPVDGLKNQKFFDEL RMNYIKELDR II --- 137  
gi | 5097870 93 QITPQEF LCMKALLLFSIIPVDGLKNQKFFDEL RMNYIKELDR II --- 137  
gi | 9168085 93 QITPQEF LCMKALLLFSIIPVDGLKNQKLFDEL RMNYIKELDR II --- 137  
gi | 1337789 93 QVTQEEFLCMKALLLFSVIPVEGLKSQKYFDELRLTYINELDR LI --- 137  
gi | 3256363 95 NMDLIEYVALKAILFFNPV - VREINDQSPVENARYAFLRSLQRRC TDK 141  
cons 97 : : \* : : : \* \* : \* : \* : \* : \* : \* : \* : \* : \* : \* : \* : \* : \* 144

gi | 2132225 138 ACKRKNPTSCSRRFYQLTKLLDSVQPIA 165  
gi | 7304901 138 ACKRKNPTSCSRRFYQLTKLLDSVQPIA 165  
gi | 6978535 138 ACKRKNPTSCSRRFYQLTKLLDSVQPIA 165  
gi | 5097870 138 ACKRKNPTSCSRRFYQLTKLLDSVQPIA 165  
gi | 9168085 138 ACKRKNPTSCSRRFYQLTKVLDSVHP IA 165  
gi | 1337789 138 NYGRK -- TNCAMRFQQLTRLMDSLQPVV 163  
gi | 3256363 142 ALENMEDESMDCR SGKLLLLLLPSLQAIA 169  
cons 145 . . \* : \* : : \* : : : . 172

