

NCBI/BLAST/blastn suite/Formatting Results - 81BAD9UH01N

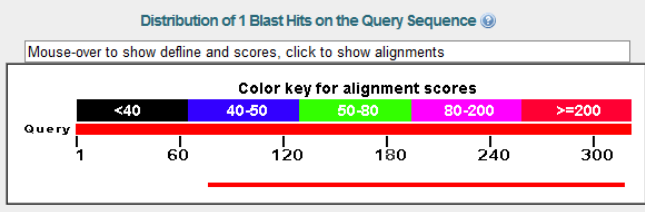
Your search is limited to records matching entrez query: AP002768.3.
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DQ188100:Homo sapiens transfection product...

Query ID	gi 75914674 gb DQ188100.1	Database Name	nr
Description	Homo sapiens transfection product sequence	Description	All GenBank+EMBL+DBJ+PDB sequences (but no EST, STS, GSS, environmental samples or phase 0, 1 or 2 HTGS sequences)
Molecule type	nucleic acid	Program	BLASTN 2.2.25+ Citation
Query Length	321		

Other reports: [Search Summary](#) [Taxonomy reports](#) [Distance tree of results](#)

Graphic Summary



Descriptions

Legend for links to other resources: [U](#) UniGene [E](#) GEO [G](#) Gene [S](#) Structure [M](#) Map Viewer [P](#) PubChem BioAssay

Sequences producing significant alignments:

Accession	Description	Max score	Total score	Query coverage	E value	Max ident	Links
AP002768.3	Homo sapiens genomic DNA, chromosome 11q, clone:RP11-1	446	446	75%	3e-127	100%	E

Alignments

Select All [Get selected sequences](#) [Distance tree of results](#)

```
>|dbj|AP002768.3| E D Homo sapiens genomic DNA, chromosome 11q, clone:RP11-673F18,
complete sequence
Length=186084

Score = 446 bits (241), Expect = 3e-127
Identities = 241/241 (100%), Gaps = 0/241 (0%)
Strand=Plus/Minus

Query 78      TC|CCCTCACCATAGTCGATGGGCTCAGTTCCTCGGAAGCATAGTATGGTGGACAAGTCA 137
Sbjct 116181   TC|CCCTCACCATAGTCGATGGGCTCAGTTCCTCGGAAGCATAGTATGGTGGACAAGTCA 116122

Query 138     AGGGCTCTGGTCTTGTGTACACTTTTATCAGCCAGCAGAAAATATCCAGAAGATTCCTA 197
Sbjct 116121   AGGGCTCTGGTCTTGTGTACACTTTTATCAGCCAGCAGAAAATATCCAGAAGATTCCTA 116062

Query 198     TGGGAGAGGACTCAGCCTGAAGCATCAGCATTGAGCAGCCAGGCTGCCACAGCCTTGAC 257
Sbjct 116061   TGGGAGAGGACTCAGCCTGAAGCATCAGCATTGAGCAGCCAGGCTGCCACAGCCTTGAC 116002

Query 258     AGATTGATTCCTGCAGCCTGGGAGCTCAGCCTCCCTTGGTGTCTGAAGGAGCTCAACAGCC 317
Sbjct 116001   AGATTGATTCCTGCAGCCTGGGAGCTCAGCCTCCCTTGGTGTCTGAAGGAGCTCAACAGCC 115942

Query 318     A 318
Sbjct 115941   A 115941
```

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