

NCBI/BLAST/blastn suite/Formatting Results - 81BRUD8X01S

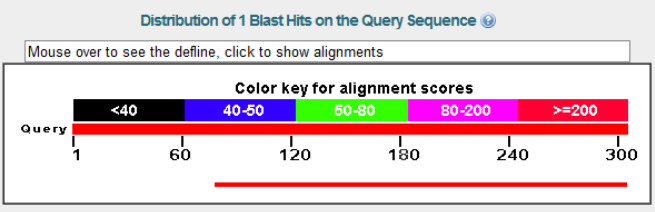
Your search is limited to records matching entrez query: AP002768.3.
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DQ188101:Homo sapiens transfection product...

Query ID [gi|75914675|gb|DQ188101.1](#) **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 306

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-i	420	420	74%	2e-119	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 = 420 bits (227), Expect = 2e-119
Identities = 227/227 (100%), Gaps = 0/227 (0%)
Strand=Plus/Minus

Query 80      TCCTCCCTCACCATAGTCGATGGGCTCAGTCCCTGGAAGCATAGTATGGTGGACAAGTCA 139
Sbjct 116181    TCCTCCCTCACCATAGTCGATGGGCTCAGTCCCTGGAAGCATAGTATGGTGGACAAGTCA 116122

Query 140     AGGGCTCTGGTCTTGTGTACACTTTTATCAGCCAGCAGAAAAATATCCAGAAGATTCCTA 199
Sbjct 116121    AGGGCTCTGGTCTTGTGTACACTTTTATCAGCCAGCAGAAAAATATCCAGAAGATTCCTA 116062

Query 200     TGGGAGAGGACTCAGCCTGAAGCAICAGCATTGAGCAGCCAGGCTGCCACAGCCTTGAC 259
Sbjct 116061    TGGGAGAGGACTCAGCCTGAAGCAICAGCATTGAGCAGCCAGGCTGCCACAGCCTTGAC 116002

Query 260     AGATTGATTCGAGCCTGGGAGCTCAGCTCCCTTGGTGTCTGAAG 306
Sbjct 116001    AGATTGATTCGAGCCTGGGAGCTCAGCTCCCTTGGTGTCTGAAG 115955
```

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