Biological Database Search and Retrieval of Nucleotide Sequence

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NCBI Nucleotide

(<u>https://www.ncbi.nlm.nih.gov/nucleotide/</u>)

- The National Centre for Biotechnology Information Nucleotide database has sequences from
- GenBank (<u>https://www.ncbi.nlm.nih.gov/genbank/</u>)
- RefSeq (<u>https://www.ncbi.nlm.nih.gov/refseq/</u>)
- TPA (<u>https://www.ncbi.nlm.nih.gov/genbank/tpa/</u>)
- PDB (<u>www.rcsb.org/pdb</u>).

<u>Step 1</u>

Open any web browser and type/enter <u>pubmed.ncbi.nlm.nih.gov</u> in the address bar and Click enter ↓



Google

pubmed.ncbi.nlm.nih.gov homepage will open



Step 2

Click on dropdown menu(All Databases) and select Nucleotide database by clicking on Nucleotide

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New Features Version 3.5.0 of Genome Workbench NCBI's sequence annotation

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<u>Step 3</u>

Enter the name of the gene in the search bar (MTHFR in this example) and click on search

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https://www.ncbi.nlm.nih.gov/nuccore/?term=MTHFR page will open

<u>Step 4</u>

Click on the FASTA to obtain the nucleotide sequence of the *Mus musculus* MTHFR gene (Sequence no. 2 i.e. sequence of your interest in this example)

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https://www.ncbi.nlm.nih.gov/nuccore/?term=MTHFR

<u>https://www.ncbi.nlm.nih.gov/nuccore/AH007491.2?report=fasta</u> page will open and the nucleotide sequence of the *Mus musculus* MTHFR gene, complete cds will open.

<u>Step 5</u> Click on send to



https://www.ncbi.nlm.nih.gov/nuccore/AH007491.2?report=fasta

<u>Step 6</u>

First click on file under Choose destination section and then

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<u>Step 6</u>

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References/Acknowledgements

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