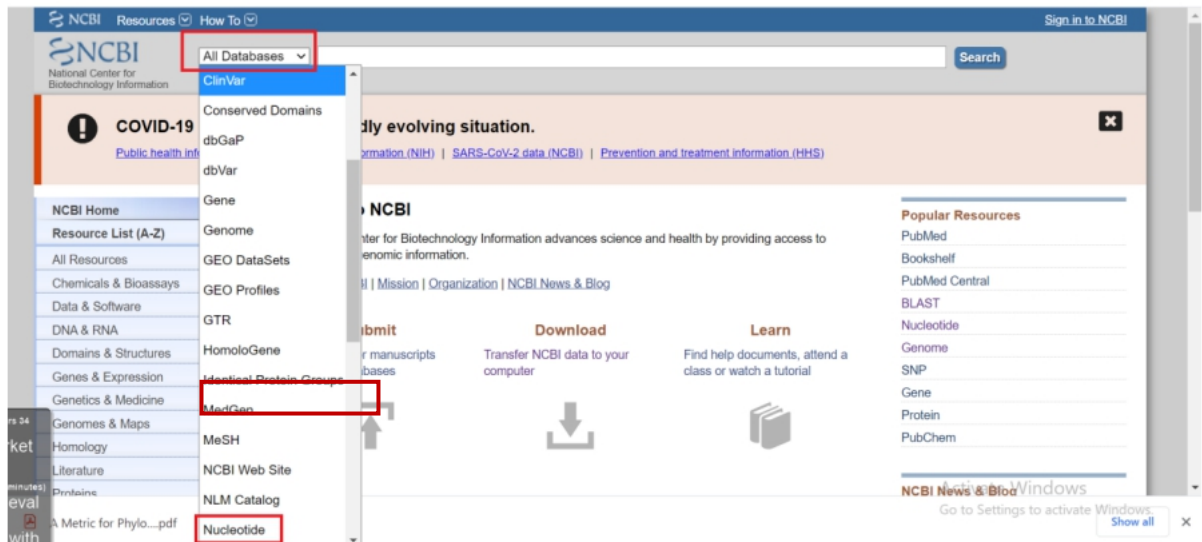
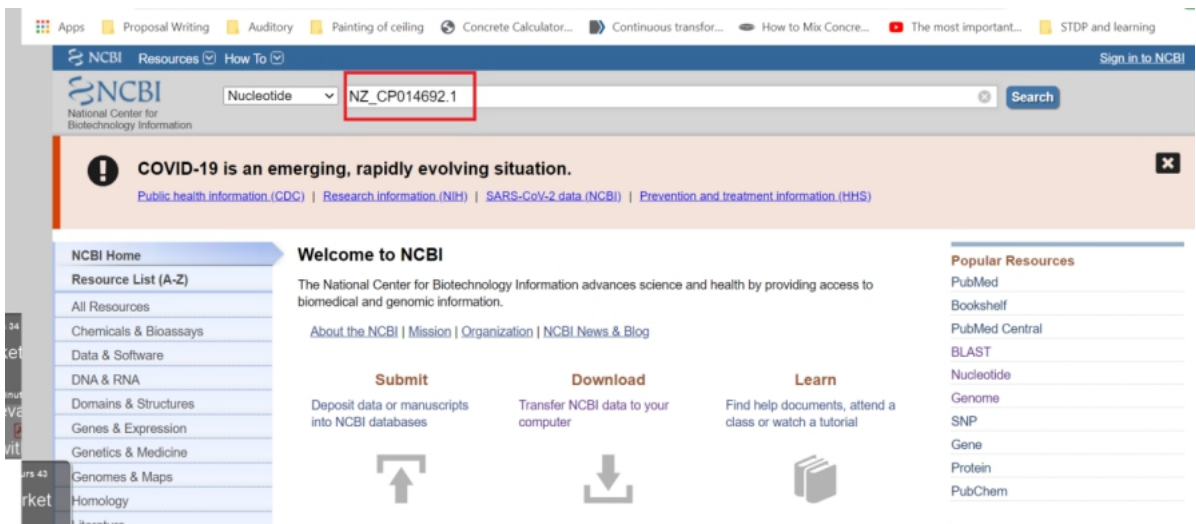


## How to download the Genome Nucleotide Sequences.

1. Go to <https://www.ncbi.nlm.nih.gov>
2. Select "Nucleotide" .



3. Enter the Accession code of the species and click "search" . E.g., NZ\_CP014692.1



4. Click on "FASTA" .

### Acetobacter aceti strain TMW2.1153 chromosome, complete genome

NCBI Reference Sequence: NZ\_CP014692.1

[FASTA](#) [Graphics](#)

Go to: ☐

LOCUS NZ\_CP014692 3725037 bp DNA circular CON 06-SEP-2020  
DEFINITION Acetobacter aceti strain TMW2.1153, complete genome.  
ACCESSION NZ\_CP014692  
VERSION NZ\_CP014692.1  
DBLINK BioProject: [PRJNA224116](#)  
BioSample: [SAMN04396917](#)  
Assembly: [GCF\\_002005445.1](#)

5. Click on 'send to' , select as follows and click 'Create File' . This will download the FASTA file of the gene sequence of 'NZ\_CP014692' .

FASTA ▾

**Acetobacter aceti strain TMW2.1153 chromosome, complete genome**

NCBI Reference Sequence: NZ\_CP014692.1

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>NZ\_CP014692.1 Acetobacter aceti strain TMW2.1153, complete genome.  
TCATGAAGCGCTTTCCGAACGGCGGCACAGAAAGCGGCGTCCGTATCAGCCCTTCGGCGCCGTCTATGC  
GGTATTTCGGCGGTTGGCGGCGAGCGCTTTCTGACGGAATGGCTGCGCGCGCTCGTCGGGACCATACC  
CTTCAGGCGGAGACCGCCCTGCACAGGTTGCTTGAGCAGAACATGGATGCAGGCGTGACCGTGATCCAC  
CGGACACGATGCACTCCCTGTCCGCCCTCCGTCTTTCCGCCGCGCCGCTCCGACACGCTTTTCATCCCT  
TGTTCCGCAAGGGACGAGTGTGGTTTACTGAAAGCGCTCTAGGCAACACGGCGGCACGAAACACCCCTT  
TCTCTGACAGCATCATGATCAACGCTTACAGCCAGCCATACAGCGCGCTGGTCTGGAAACACGTACCGG  
CGGTCAAGACAATTGTCTATCTCTGAGCGGATCAGGAAAAAATCCGGATGGCGTCATCCCTGAACAGCT  
TTTCGGCGTGTCCATTCTTGTCTTGATACCGGTGAAAGGTGAAATGCGGCAAGTGCGAGGCGCTGGT  
TAATGCCCTGTGGCTGCCCTGGCGATTTCATTGAAACGATCTGATTCCTCTTGATGAGCCGATGCTGATC  
CTGTCTTCGGCAGGCGTTGAGGCAAGCATGGCGTCTGAAAGCCGCTTCGGCGGGCGTCTTCGCATCA  
TCCAGCTCGGCGTCCGCGTCTGAAAGCGCATCGGACCTGGATCTGATCCTGCTGATGCCGCGAGGACGA  
CTACCCGGAAGCCCGCAGACGCTGCTGATCTGAACATGCCGCTGAACGGCGCGGATATCTTACGGCCGCT  
ATTCTCTGATACGACACAAGGACCGTTTCGCTCTGCTGGGAGGACCGACCCGCTCACTTCGGCTTCA  
GCAGTCACGATGCAATGGTCTGCTGCTCCCGCGCCGACGGCTGGCCGACGACGACACAGAGTGC  
GCTGCTGCGGCTTCGGCGGCGGATGCTGATGAAATTCGCGGCGGCTGCTGAGAGCT

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