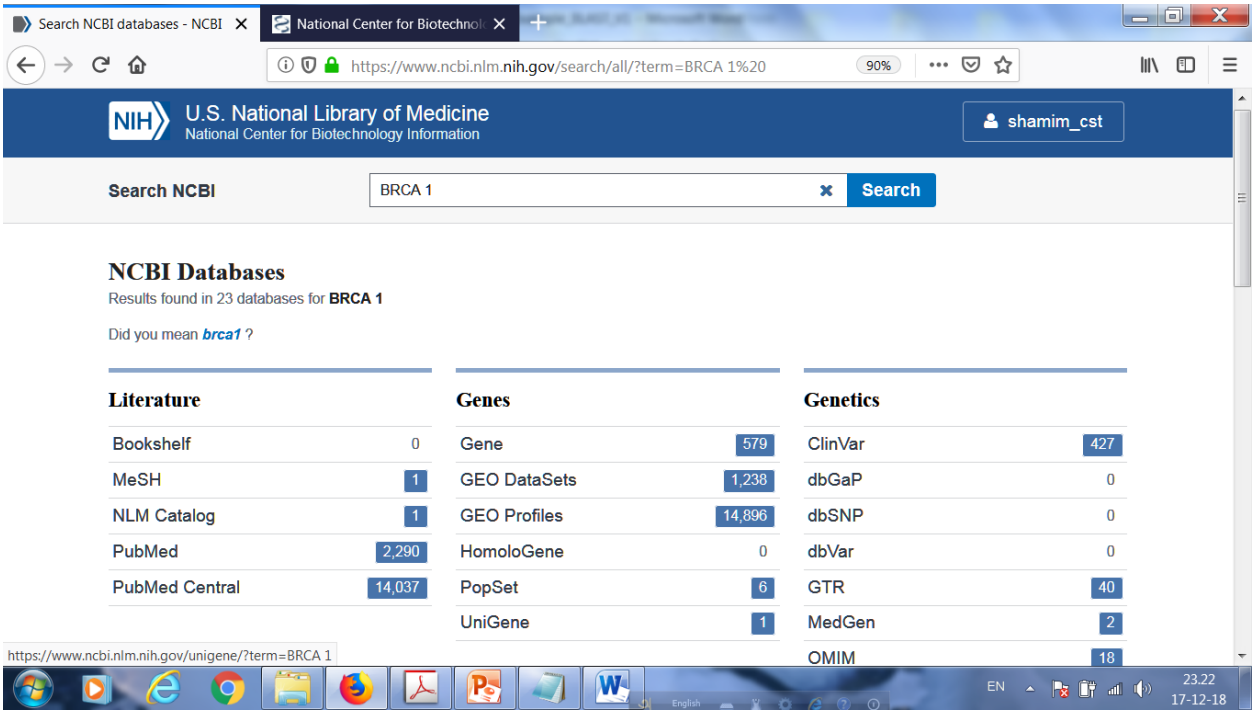

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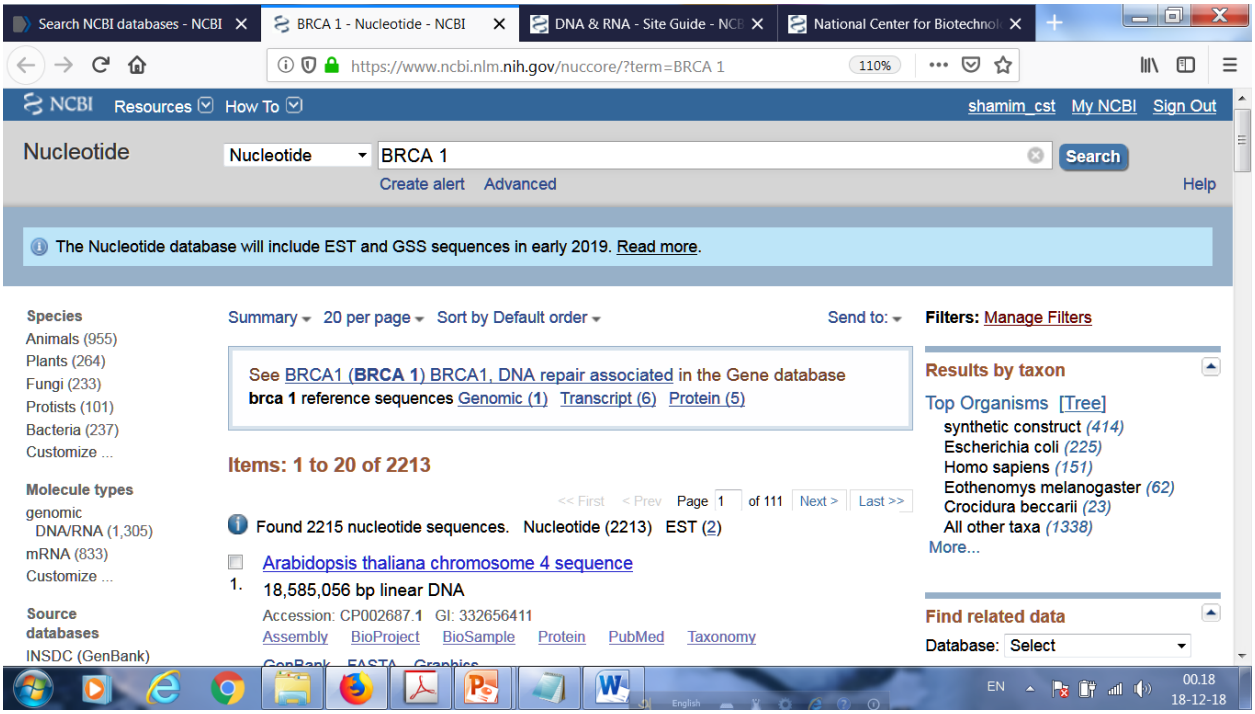

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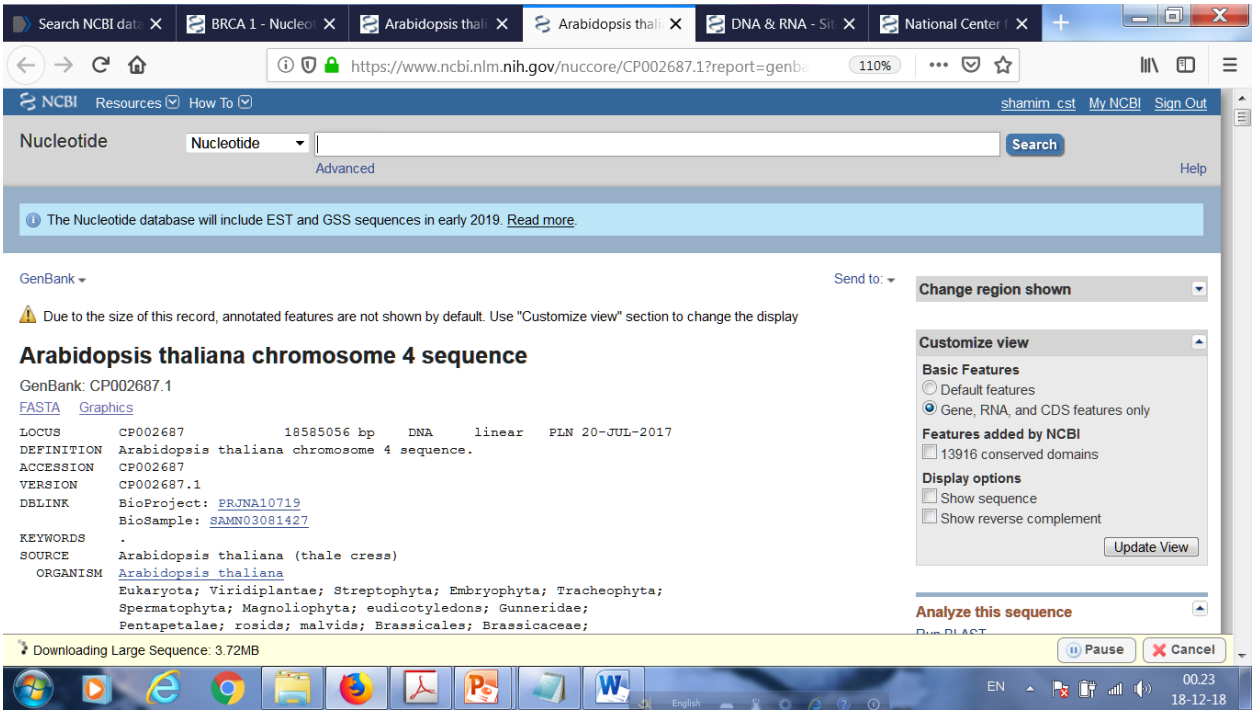

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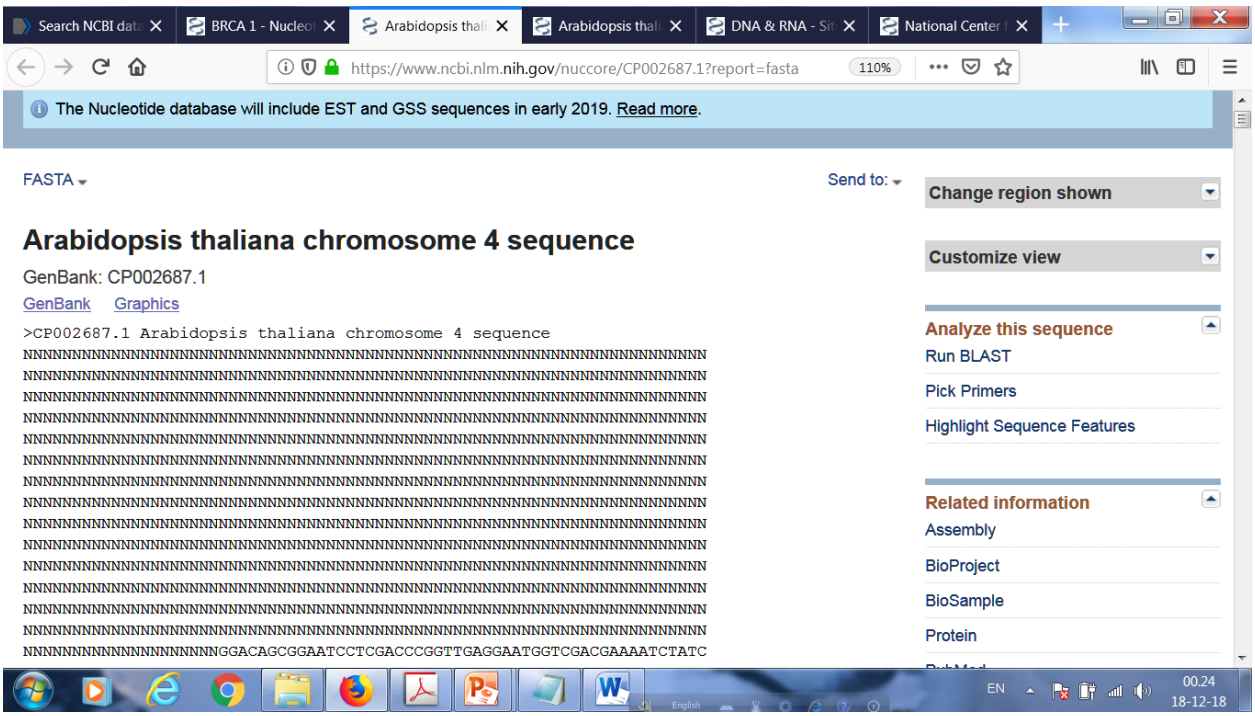
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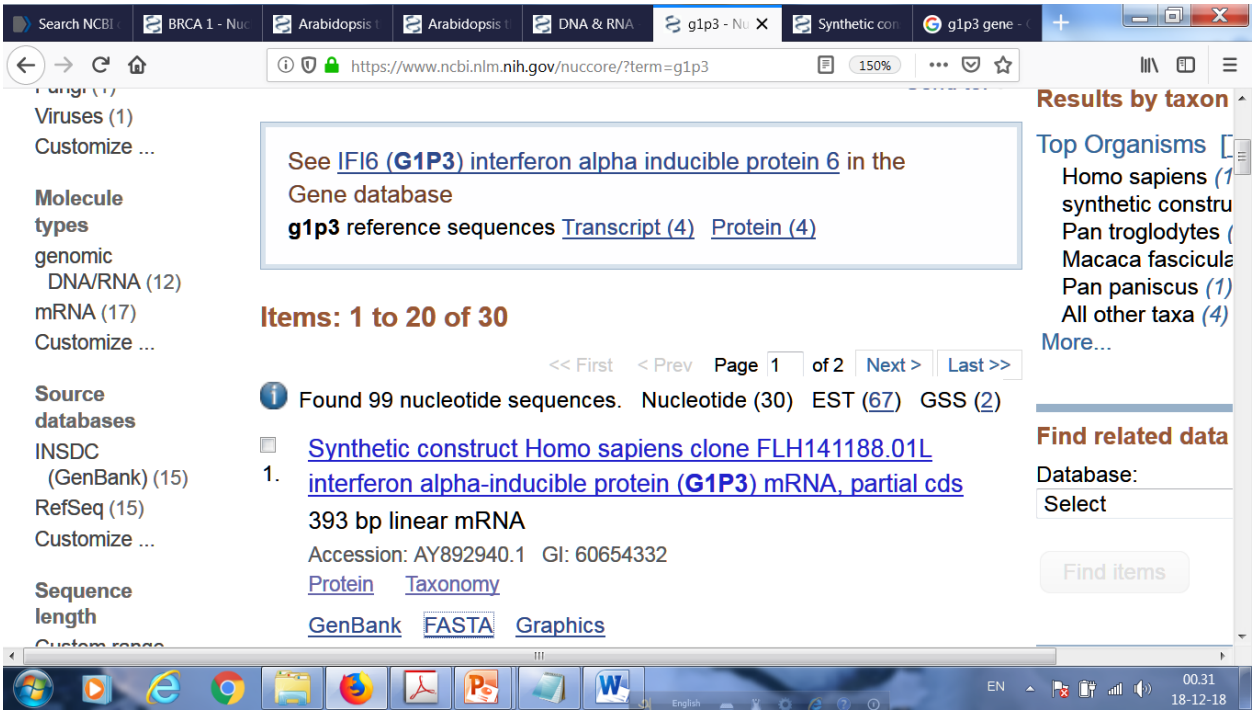
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See IFI6 (G1P3) interferon alpha inducible protein 6 in the Gene database
g1p3 reference sequences Transcript (4) Protein (4)
Items: 1 to 20 of 30
<< First < Prev Page 1 of 2 Next > Last >>
Found 99 nucleotide sequences. Nucleotide (30) EST (67) GSS (2)
1. Synthetic construct Homo sapiens clone FLH141188.01L interferon alpha-inducible protein (G1P3) mRNA, partial cds
393 bp linear mRNA
Accession: AY892940.1 GI: 60654332
Protein Taxonomy
GenBank FASTA Graphics
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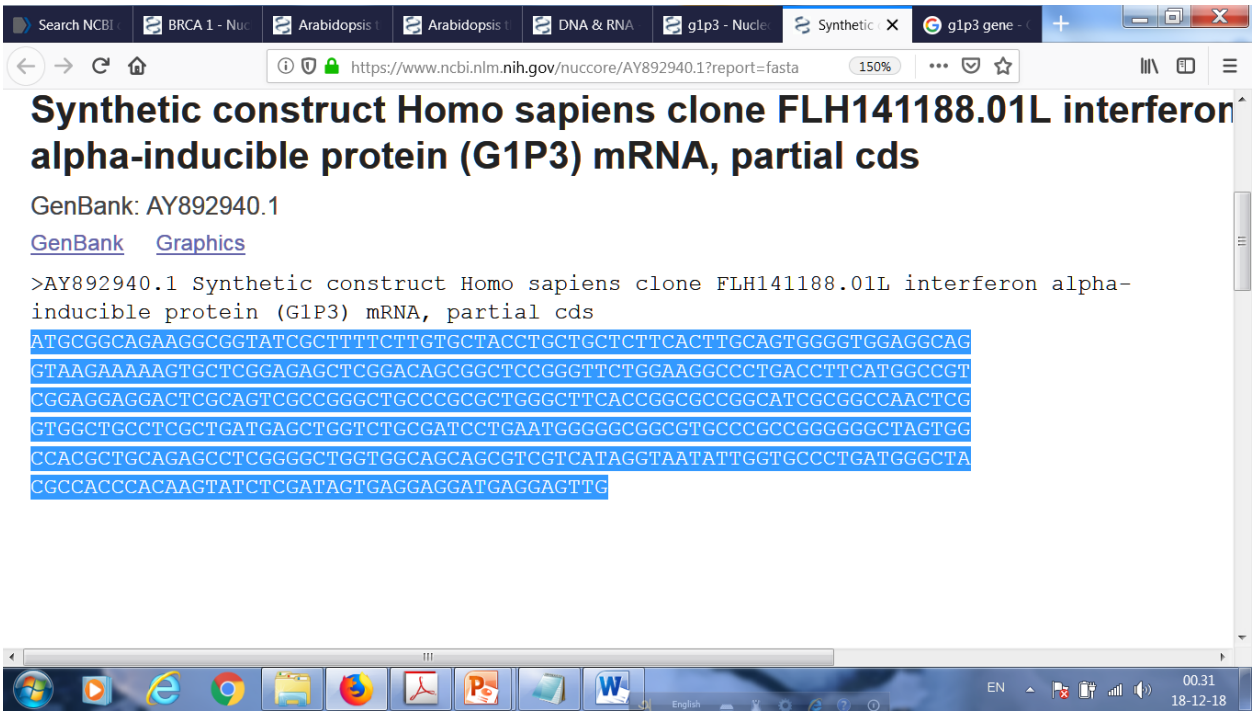
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>AY892940.1 Synthetic construct Homo sapiens clone FLH141188.01L interferon alpha-inducible protein (G1P3) mRNA, partial cds
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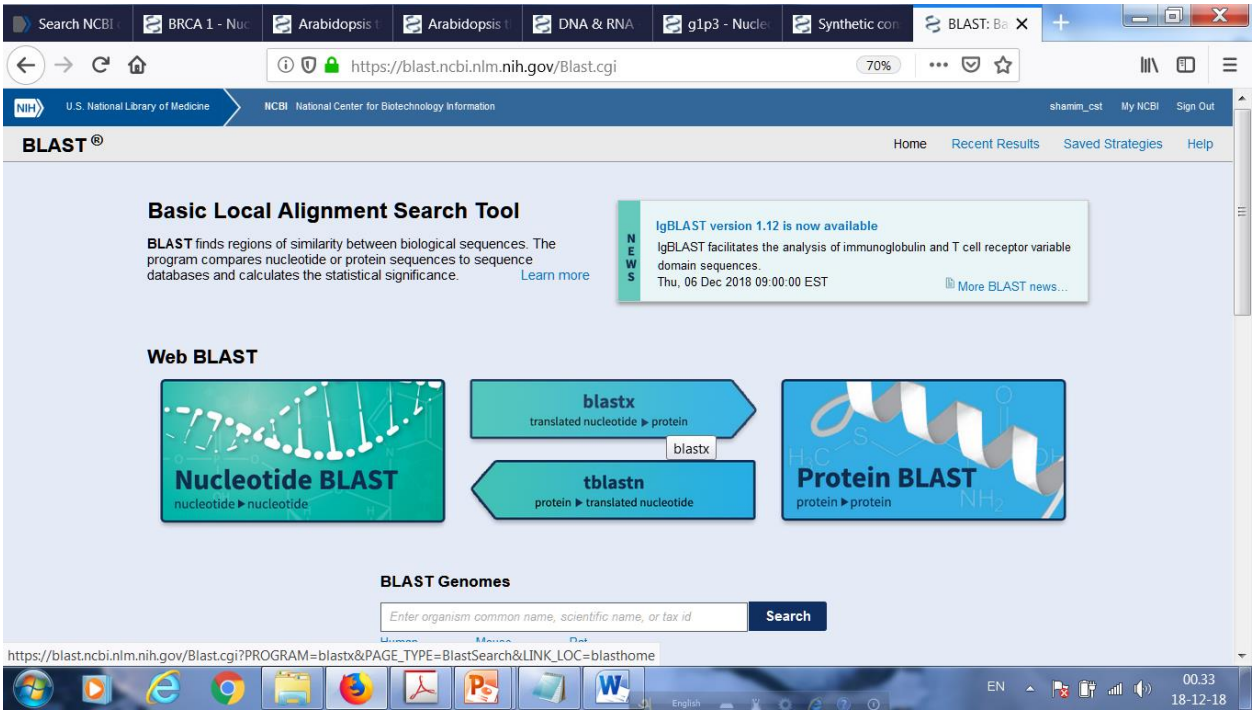

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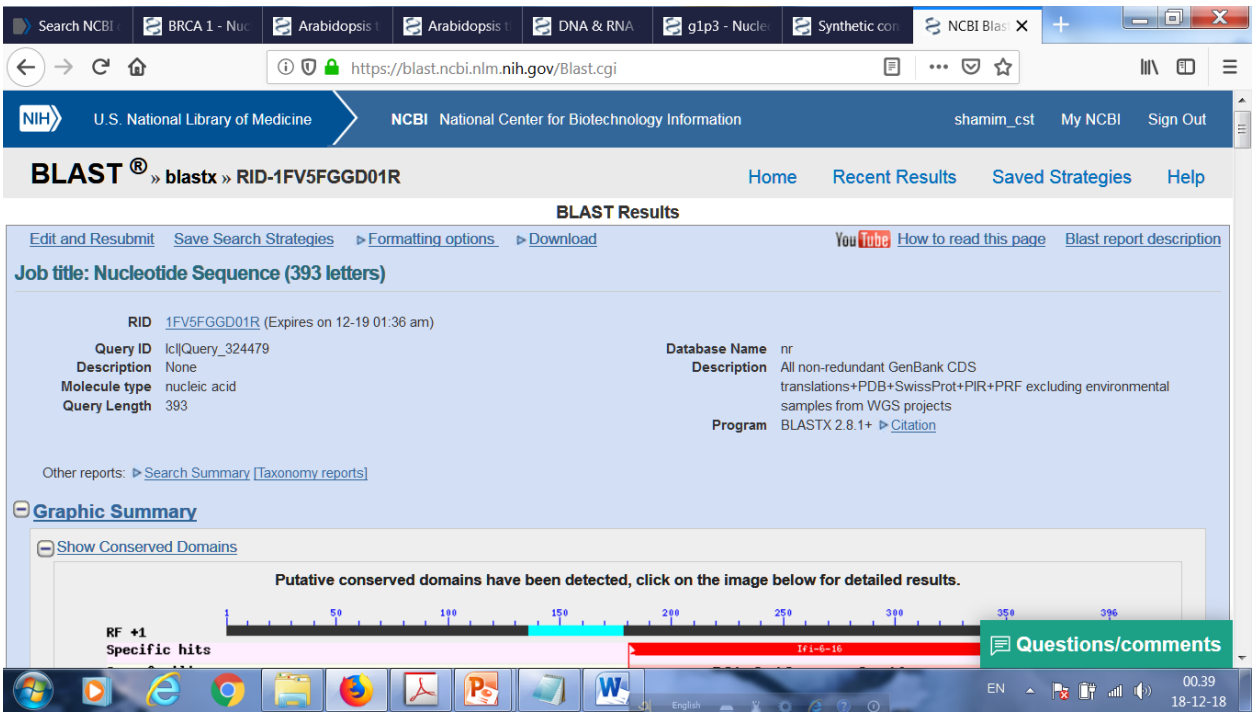
Gip3 sequence found from FASTA



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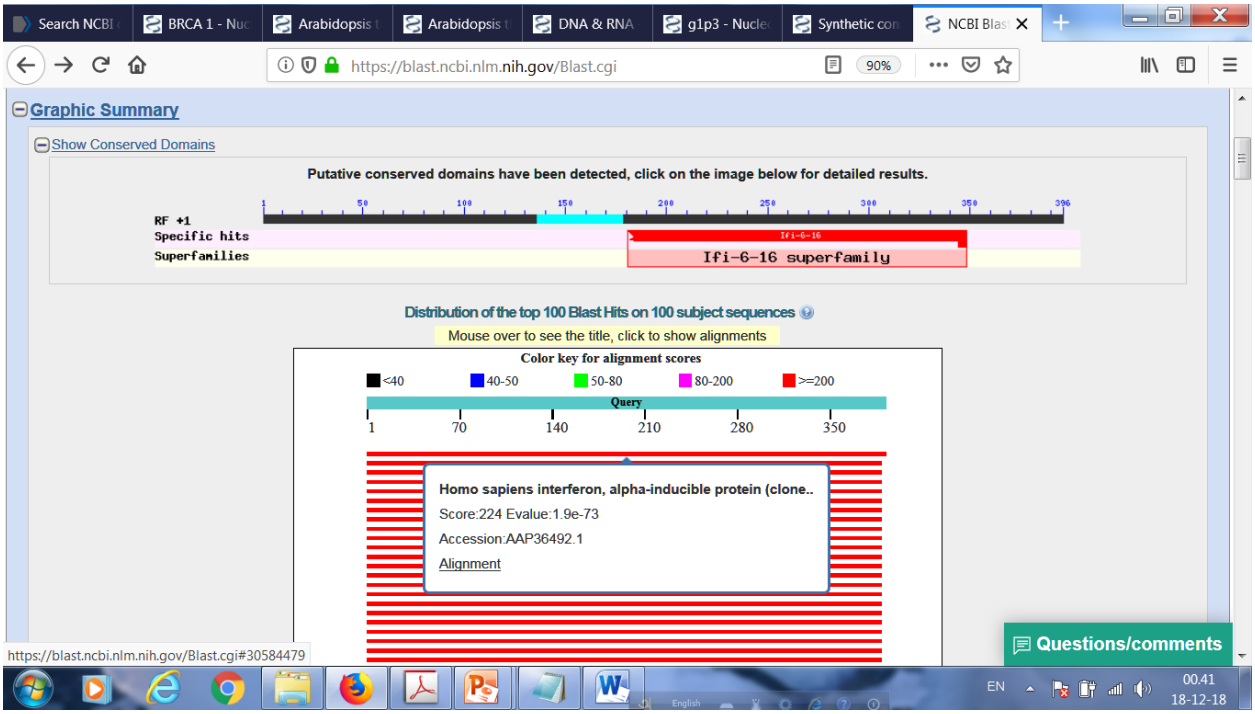

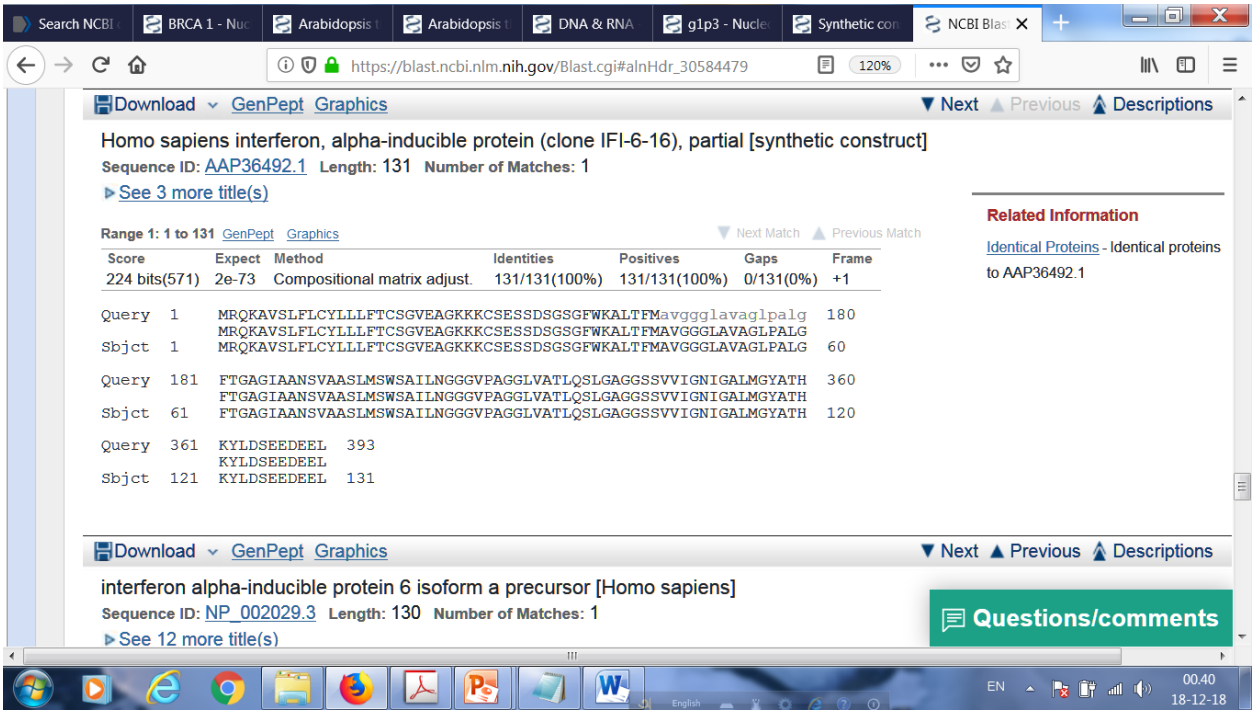

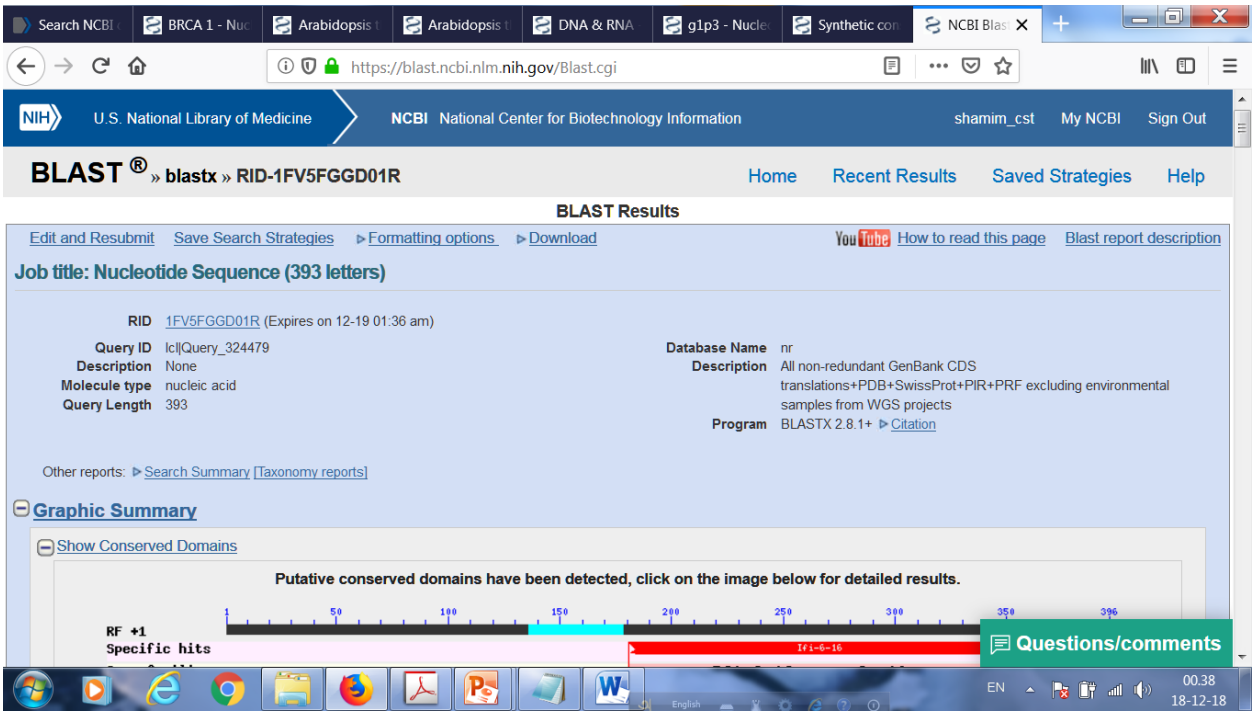
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Query ID lcl|Query_324479
Description None
Molecule type nucleic acid
Query Length 393
Database Name nr
Description All non-redundant GenBank CDS translations+PDB+SwissProt+PIR+PRF excluding environmental samples from WGS projects
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Other reports: Search Summary [Taxonomy reports]
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Putative conserved domains have been detected, click on the image below for detailed results.
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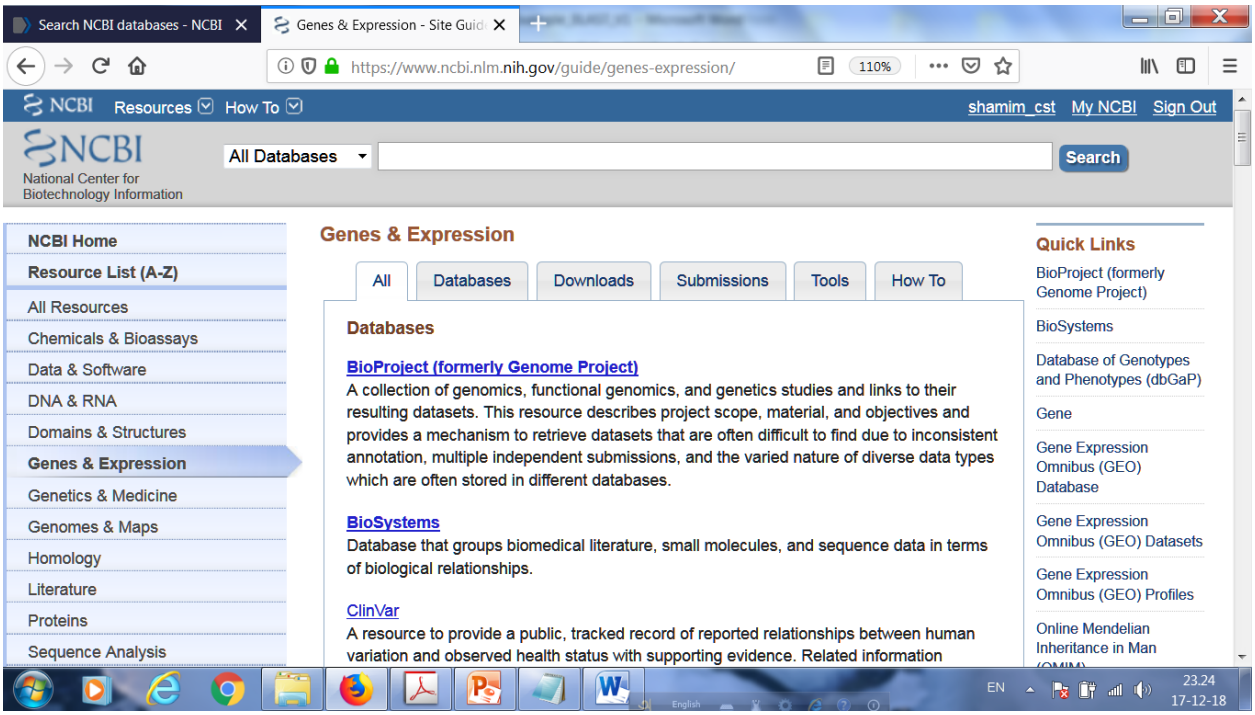
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Database that groups biomedical literature, small molecules, and sequence data in terms of biological relationships.
ClinVar
A resource to provide a public, tracked record of reported relationships between human variation and observed health status with supporting evidence. Related information
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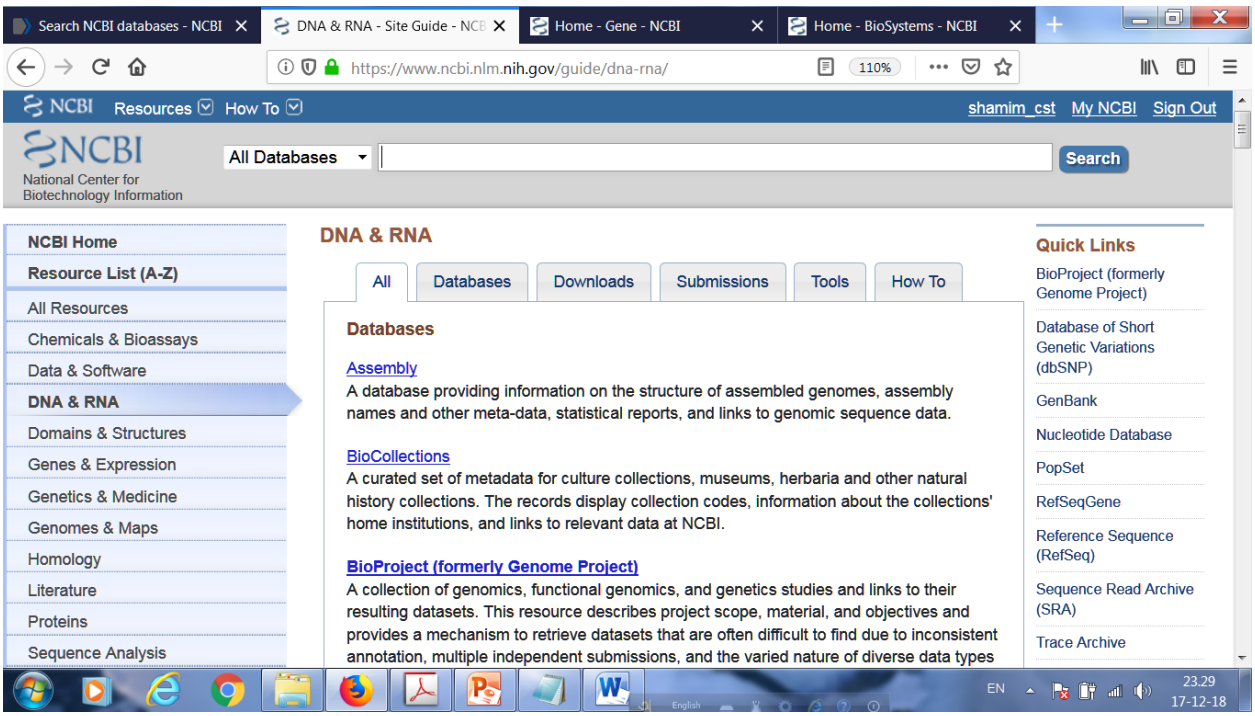
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A curated set of metadata for culture collections, museums, herbaria and other natural history collections. The records display collection codes, information about the collections' home institutions, and links to relevant data at NCBI.
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Nucleotide Database
PopSet
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Reference Sequence (RefSeq)
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Trace Archive
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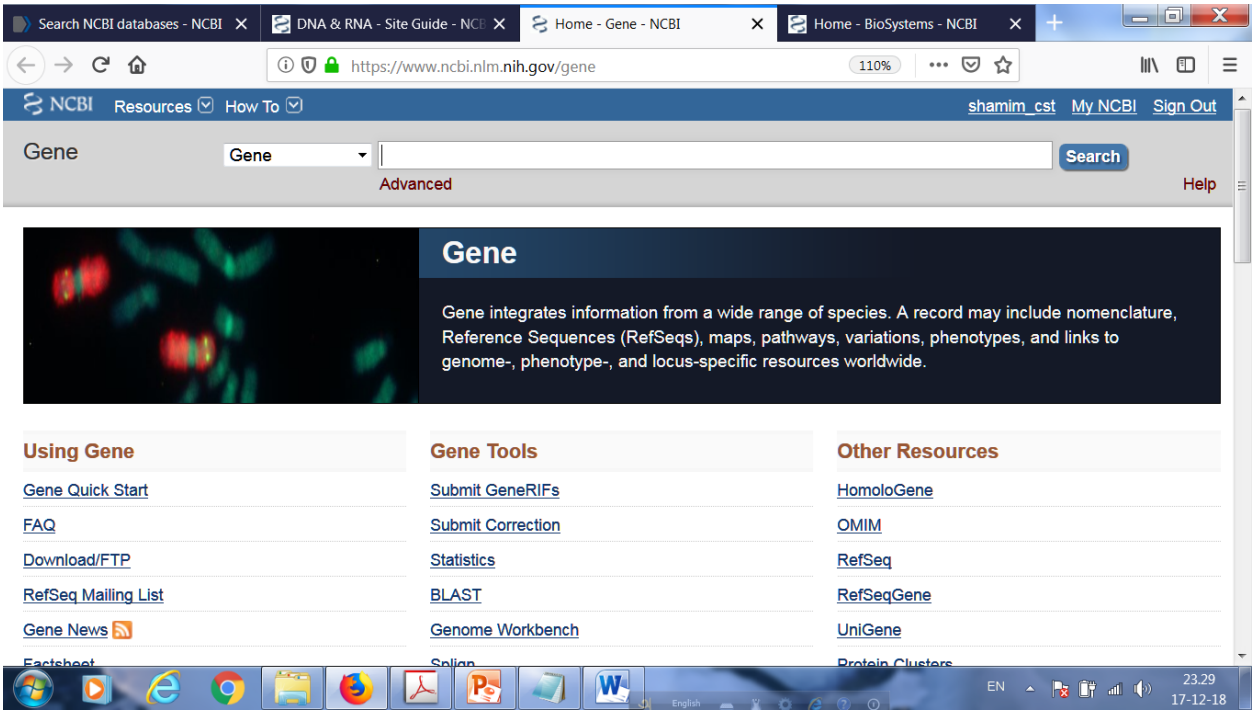
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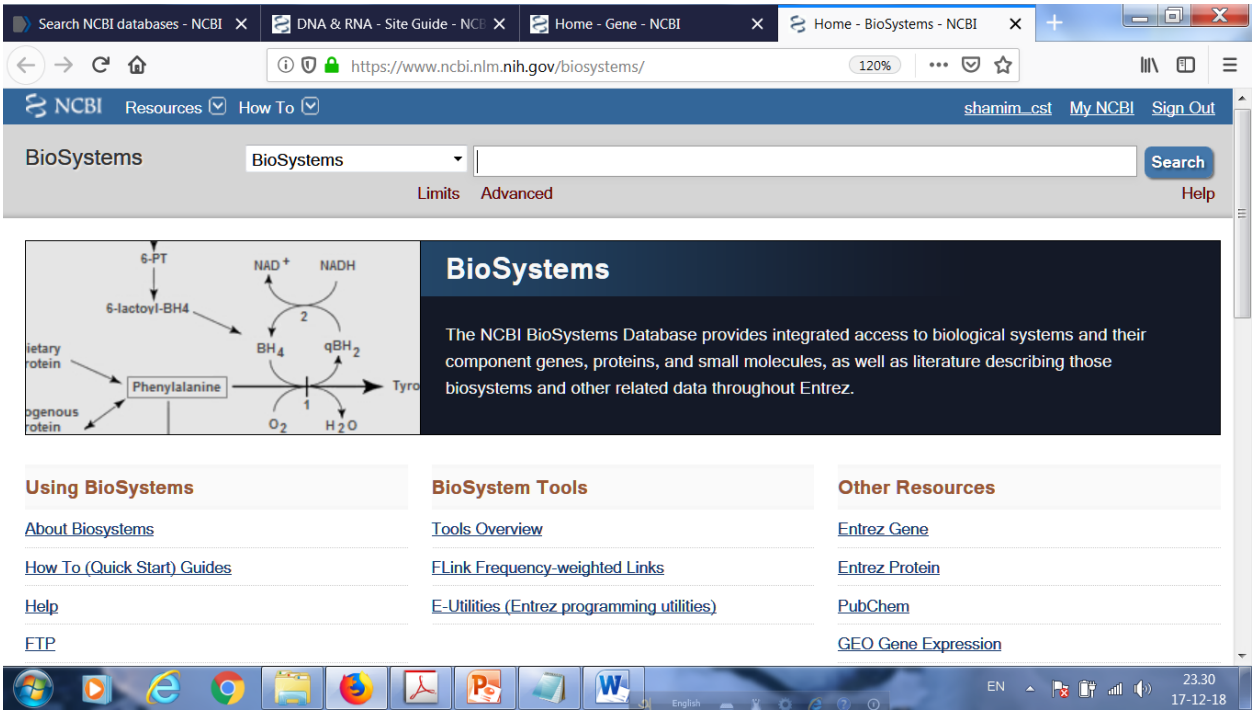
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FLink Frequency-weighted Links
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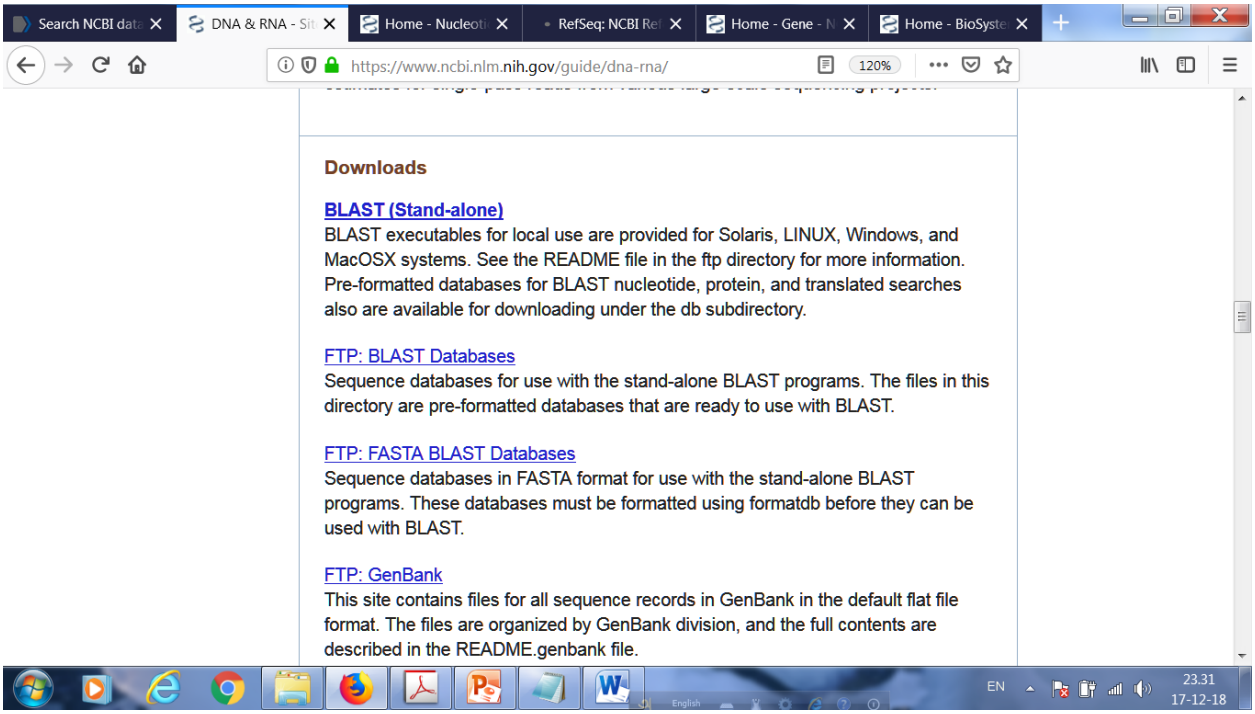
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BLAST executables for local use are provided for Solaris, LINUX, Windows, and MacOSX systems. See the README file in the ftp directory for more information. Pre-formatted databases for BLAST nucleotide, protein, and translated searches also are available for downloading under the db subdirectory.
FTP: BLAST Databases
Sequence databases for use with the stand-alone BLAST programs. The files in this directory are pre-formatted databases that are ready to use with BLAST.
FTP: FASTA BLAST Databases
Sequence databases in FASTA format for use with the stand-alone BLAST programs. These databases must be formatted using formatdb before they can be used with BLAST.
FTP: GenBank
This site contains files for all sequence records in GenBank in the default flat file format. The files are organized by GenBank division, and the full contents are described in the README.genbank file.
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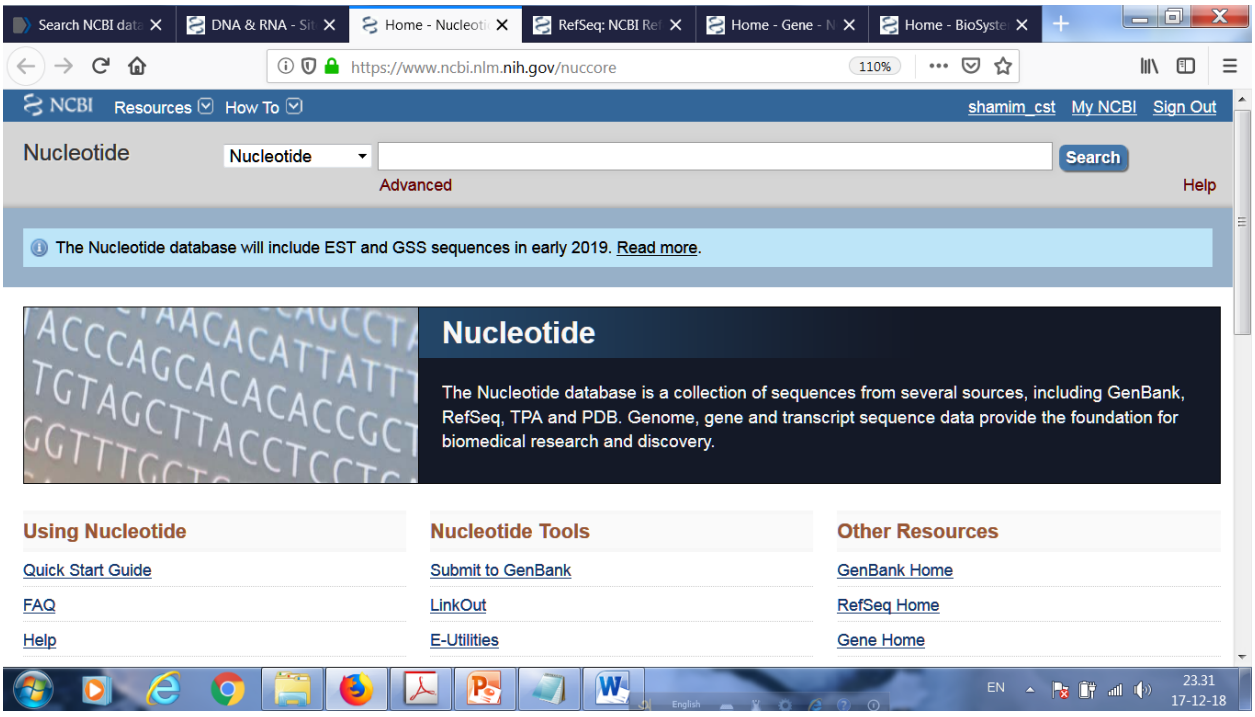
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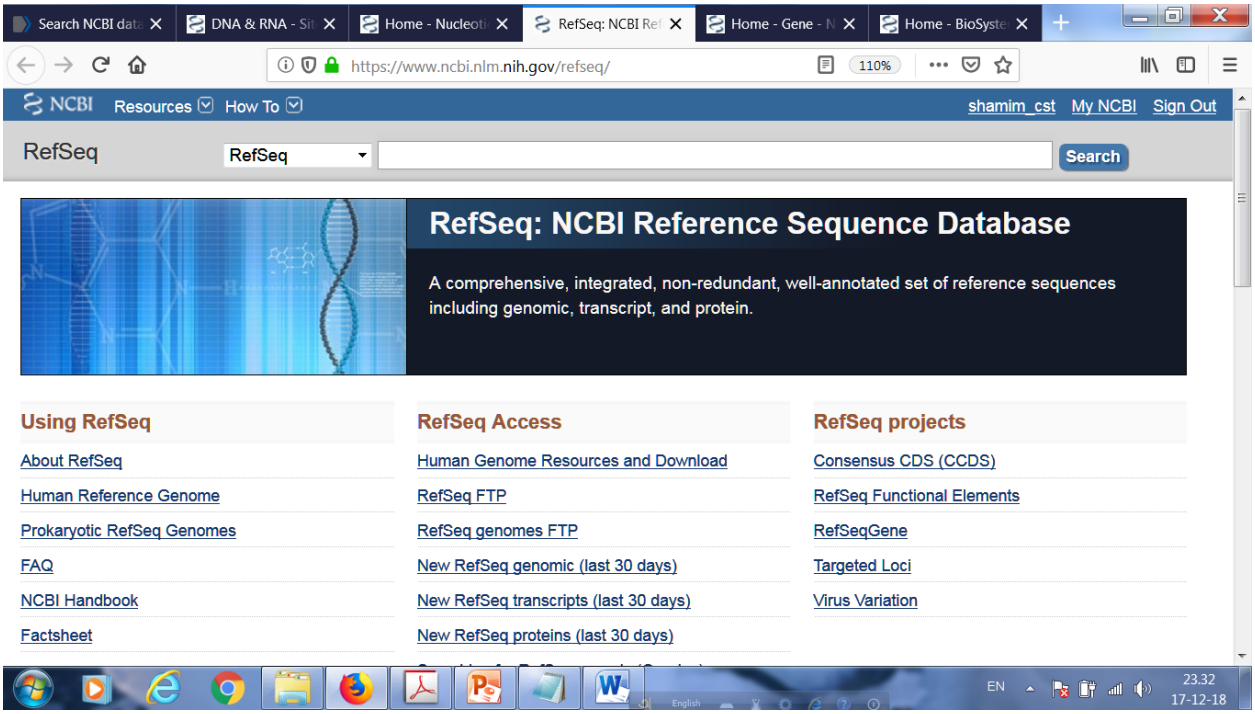
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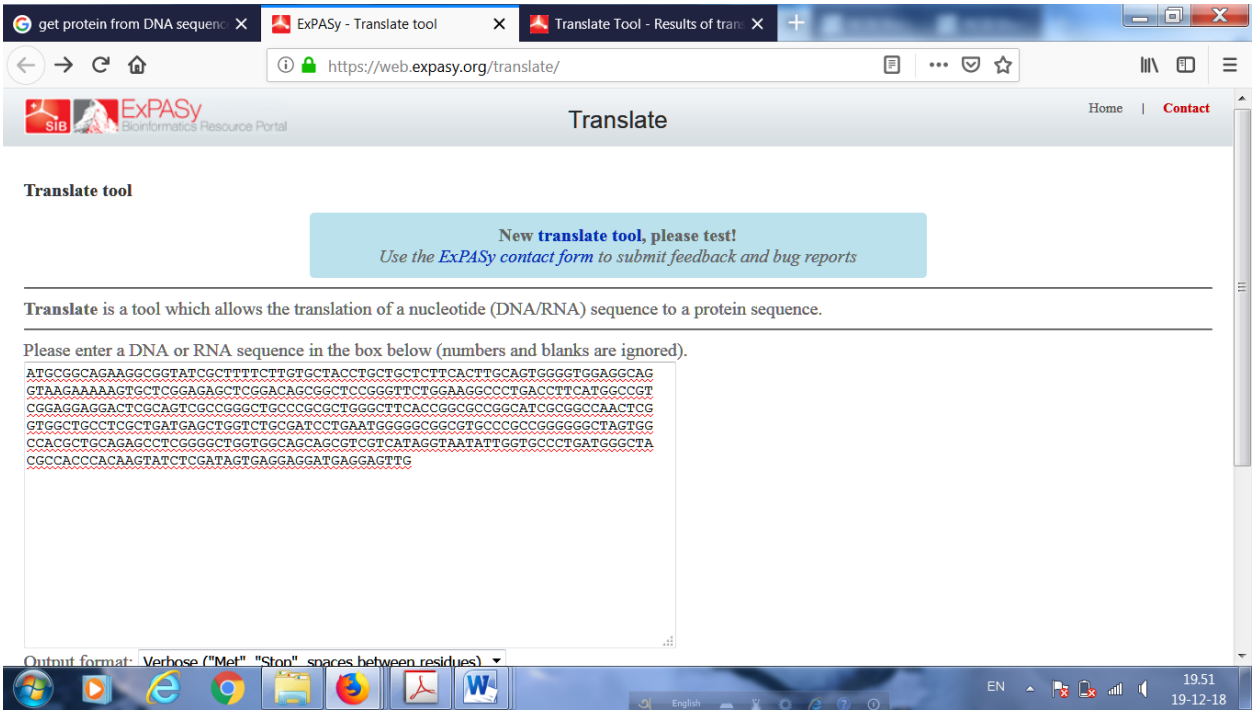
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GTAAGAAAAAGTGCTCGGAGAGCTCGGACAGCGGCTCCGGGTTCTGGAAGGCCCTGACCTTCATGGCCGT
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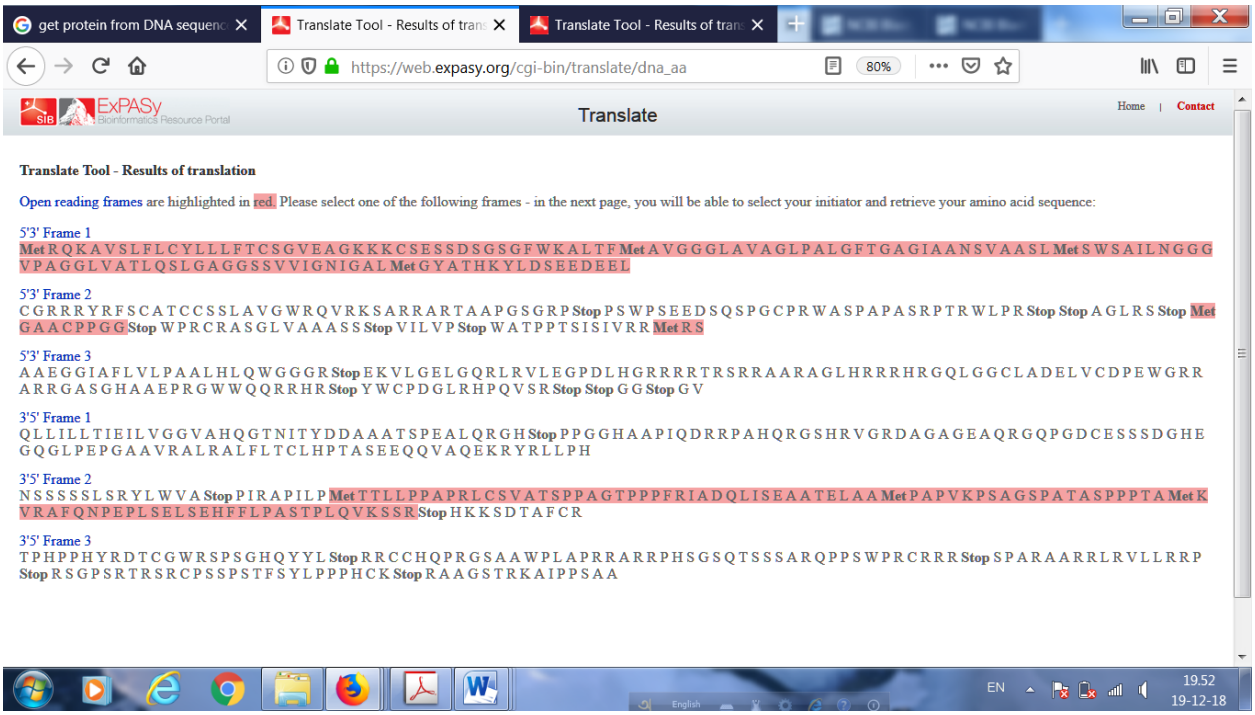
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C G R R R Y R F S C A T C C S S L A V G W R Q V R K S A R R A R T A A P G S G R P Stop P S W P S E E D S Q S P G C P R W A S P A P A S R P T R W L P R Stop Stop A G L R S Stop Met
G A A C P P G G Stop W P R C R A S G L V A A A S S Stop V I L V P Stop W A T P P T S I S I V R R Met R S
5'3' Frame 3
A A E G G I A F L V L P A A L H L Q W G G G R Stop E K V L G E L G Q R L R V L E G P D L H G R R R R T R S R R A A R A G L H R R R H R G Q L G G C L A D E L V C D P E W G R R
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3'5' Frame 1
Q L L I L L T I E I L V G G V A H Q G T N I T Y D D A A A T S P E A L Q R G H Stop P P G G H A A P I Q D R R P A H Q R G S H R V G R D A G A G E A Q R G Q P G D C E S S S D G H E
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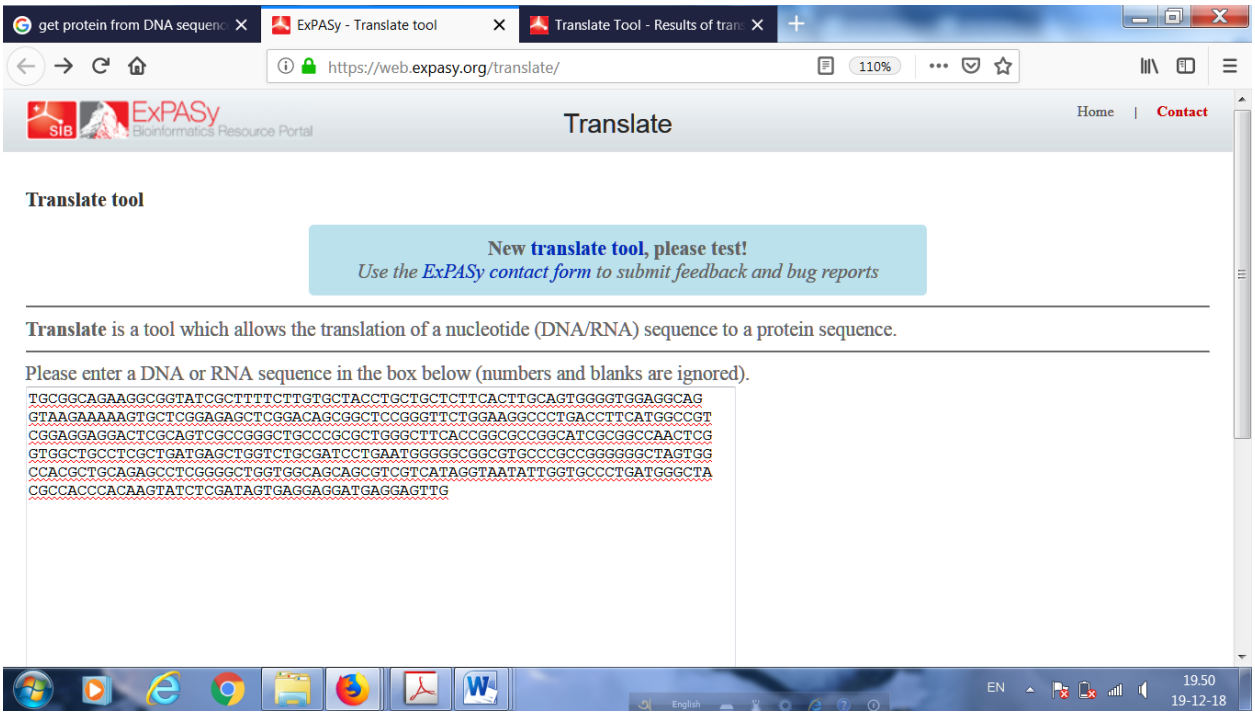

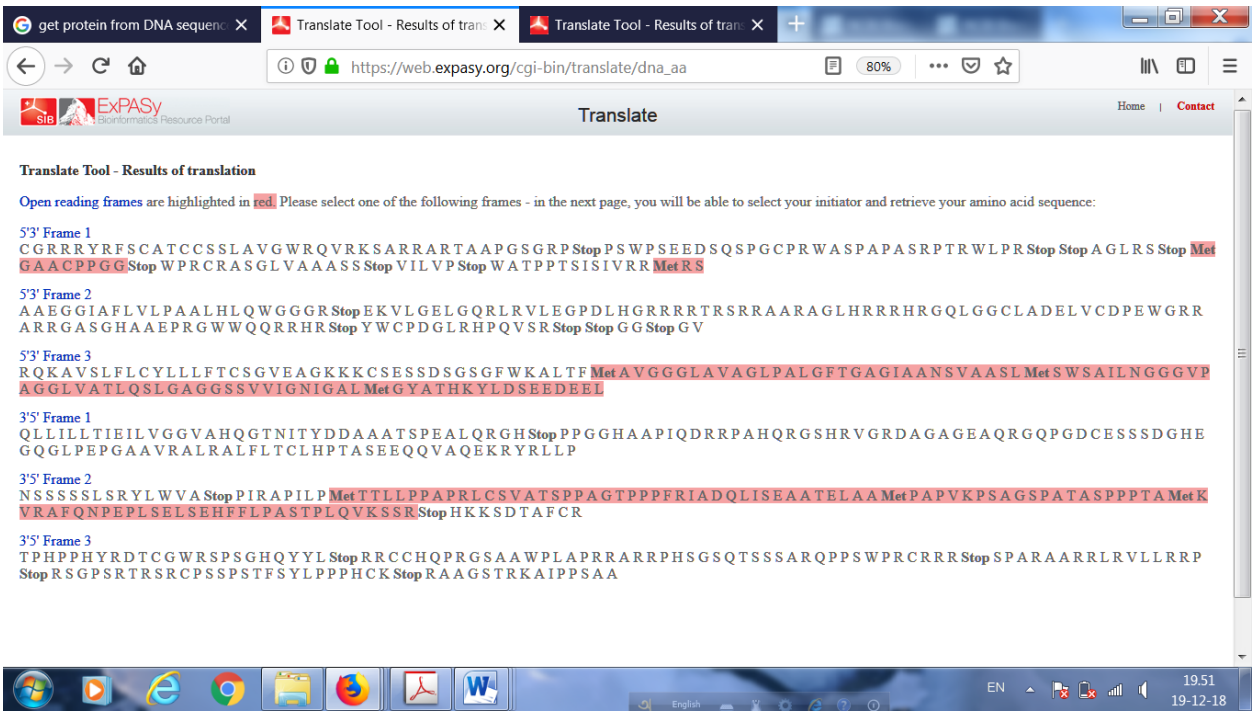
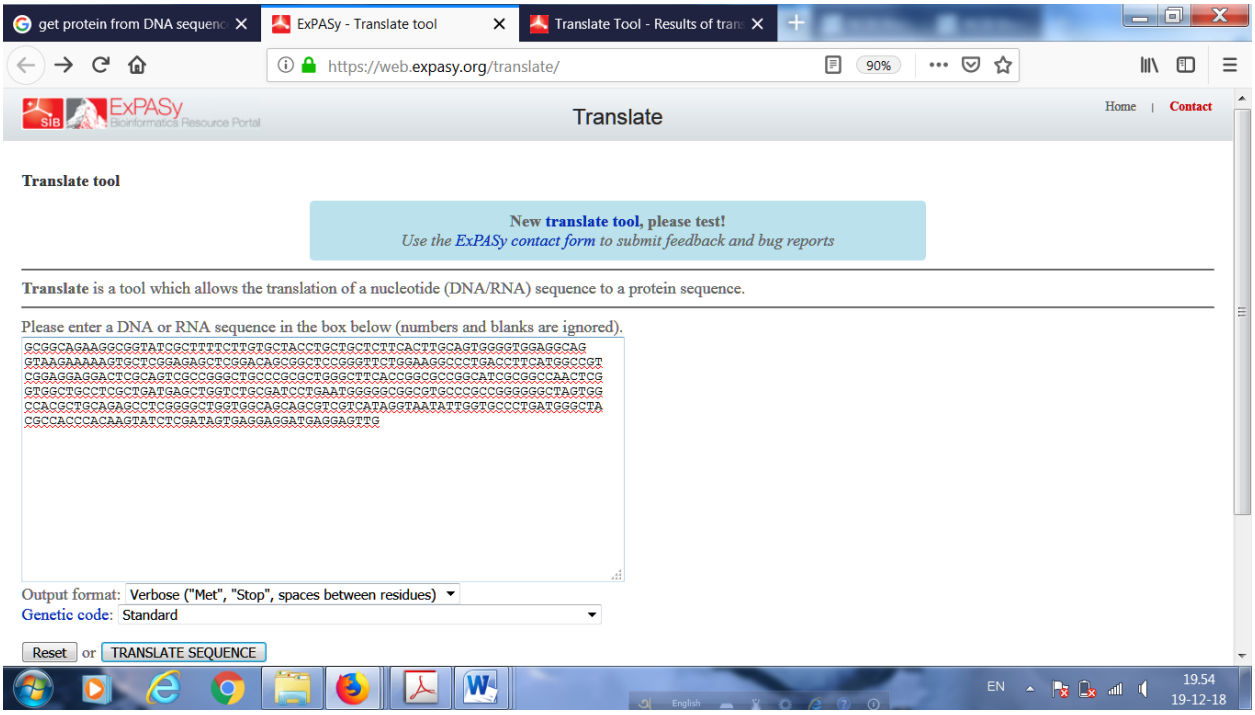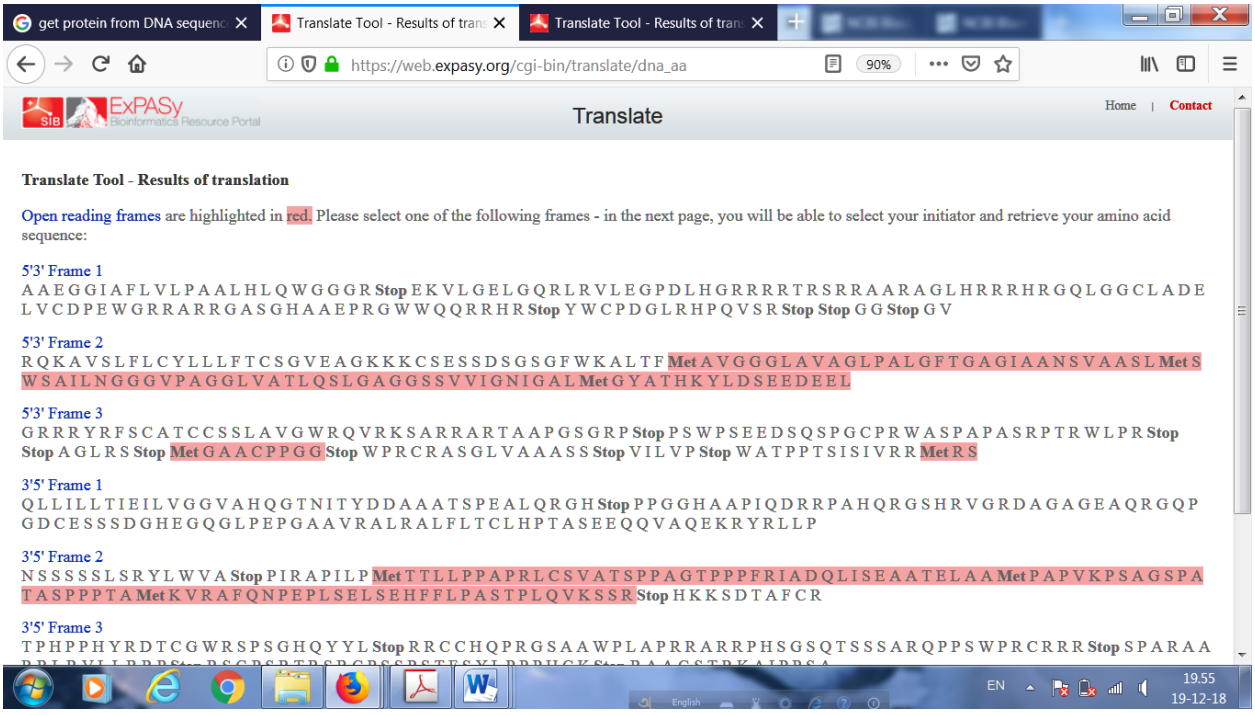
After deleting second Nucleotide