## **Annotation Exercise**

It will never really be this easy, but eventually you will be able to annotate your own data.



## Where we are

- 13:30-14:00 Primer Design to Amplify Microbial Genomes for Sequencing
- 14:00-14:15 Primer Design Exercise
- 14:15-14:45 Molecular Barcoding to Allow Multiplexed NGS
- 14:45-15:15 Processing NGS Data de novo and mapping assembly
- 15:15-15:30 Break
- 15:30-15:45 Assembly Exercise
- 15:45-16:15 Annotation
- 16:15-16:30 Annotation Exercise
- 16:30-17:00 Submitting Data to GenBank



# Use our asembled genome sequence

- Try annotating it at VIGOR website http://www.jcvi.org/vigor/submission.php
- Look at the .tbl file



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CGGGTGAACAACTTCTTTCTTCAGGTCCACTATGCCCGGATGCAGCGGGTATCCCAACA

TGGATTTGGATTGCCTACTCCCCCAGGAGTGGGTGCAGCACTTCTACCAAGAGGCAGCTC

CAGCACAATCTGATGTGGCTCTATTGAGATTTGTGAATCCAGACACGGGTAGGGTCCTGT TTGAGTGCAAACTTCATAAATCAGGCTATGTCACAGTGGCTCACACCGGTCAGCATGATT

TGCTGGATTGGCATCTGATGTCCTTGGCTCTGGACTTGGTTCCCTAATCAATGCTGGGGC TGGGGCCATCAACCAAAAGATTGATTTTGAAAAATAATAGAAAATTGCAGCAAGCTTCCTT

CCAGTTTAGCAGTAATCTACAACAGGCTTCCTTTCAACACGATAAAGAGATGCTCCAAGC ACAAATTGAGGCCACTAAAAAGTTGCAACAGGAAATGATGAAAGTCAAGCAGGCAATGCT

CAGGACTCCTACCCCCGCTCGGGGGCTCCTCCAACACATCTTCTAATGCTTCCACTGCTAC

TTCTATACATTCAAATCAAACTGTTTCAACGAGACTTGGTTCTACAGCTGGTTCTGGTAC CAATGTCTCGAGTCTCCCGTCAACTGCAAGGACTAGGAGTTGGGTTGAGGATCAAAACAG

AAATTTGTCACCTTTCATGAGGGGGGGCTCATAACATATCGTTTGTCACCCCACCATCTAG CAGATCCTCTAGCCAAGGCACAGTCTCAACCGTGCCTAAAGAAGTTTTGGACTCCTGGAC

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TGGCGCTTTCAACACGCGCAGGCAGCCTCTCTTCGCTCACATTCGTAGGCGAGGGGAGTC

Browse\_

Select virus type:

Norovirus

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Submit Clear

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### VIGOR - VIRAL GENOME ORF READER

Paste your sequences in FASTA format into the box

below or upload FASTA file to the right

#### → Run VIGOR

→ About VIGOR

→ Instructions & Test Samples

→ Coronavirus

→ Influenza virus

→ Measles & mumps virus

→ Metapneumovirus

→ Norovirus

→ Parainfluenza & Sendai virus

→ Respiratory syncytial virus (RSV)

→ Rhinovirus

→ Rotavirus

→ Rubella virus

→ SARS

Report a Bug

ACGGGTGTAA

Email address:

tstockwe@jcvi.org

→ Venezuelan equine → encephalitis virus & alphavirus

WELL MILL CALLS & DESCRIPTION

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→ Run VIGOR	Thank you for your submission. Your submission ID is <b>'dmzweb2_1898_1_WedAug211817402013'.</b> A notification email has been sent to <b>'tstockwe@jcvi.org'</b> You will receive another notification email once your job has been processed.				
→ Instructions & Test Samples					
<ul> <li>→ Coronavirus</li> <li>→ Coronavirus</li> <li>You will receive another notification email once your job has been processed.</li> <li>Back to submission portal</li> <li>Your results will be available at the following URL:</li> </ul>					
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→ Measles & mumps viruses					
→ Metapneumovirus					
→ Norovirus					
→ Parainfluenza & Sendai → viruses					
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→ Rubella virus	The job ID is 'dmzweb2_1898_1_WedAug211817402013'		
→ Metapneumovirus	~		
→ Norovirus			
→ Measles & mumps viruses			
Yellowfever & Japanese encephalitis viruses			
Respiratory syncytial virus			

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<pre>&gt;contig00001.1 ORF1 1699 Aa 3 5102 genome polyprotein MKMASNDASAAAVANSNNDTAKSSSDKMFSNMAVTFKRALGARPKQPPPREIPQRPPRP TPELVKKIPPPPNGEDEVVVSYSAKDGVSGLPELSTVRQPEETNTAFSVPPLNQRENRD AKEPLTGTILEMWDGEIYHYGLYVERGLVLGVHKPPAAISLAKVELTPLSLFWRPVYTPQ YLISPDTLKKLHGETFPYTAFDNNCYAFCCWVLDLNDSWLSRRMIQRTTGFFRPYQDMNR KPLPTMDDSKLKKVANIFLCALSSLFTRPIKDIIGKLRPLNINILASCDWTFAGIVESL ILLAELFGVFWTPPDVSAMIAPLLGDFELQGPEDLVVELVPVVMGGIGLVLGFTKEKIGR MLSSAASTLRACKDLGAYGLEILKLVMKWFFPKKEEANELAMVRSIEDAVLDLEAIENNH MTTLLKDKDSLATYMRTLDLEEEKARKLSTKSASPDIVGTINALLARIAAARSLVHRAKE ELSSRPRPVVLMISGRPGIGKTHLAREVAKRIAASLTGDQRVGLIPRNGVDHWDAYKGER VVLWDDYGMSNPIHDALRLQELADTCPLTLNCDRIENKGKVFDSDVIIITTNLANPAPLD YVVFAGSRRIDFLVYAEAPEVEKAKRDFFGQPDMWKNAFSSDFSHIKLALAPQGGFDKN GNTPHGKGVMKTLTTGSLIARASGLLHERLDEFELQGPALTTFNFDRNKVLAFRQLAAEN KYGLIDTMKVGRQLKDVKTMPDLKQALKNISIKKCQIVYSGCTYTLESDGKGNVKVDRIQ STSVQINNELAGALHHLRCARIRYYVKCVQEALYSIIQIAGAAFVTTRIIKRVNIQDLMS KPQVENTEEATNKDGCPKPKDDEEFVISSDDIKTEGKKGKNKTGRGKKHTAFSSKGLSDE EVDEYKHIREENGRYSIEEVLQDRDKYYEVEVAIARATEEDFCEEEEAXIRQRIFRPTRK QRKEERASLGLVTGSEIRKNPDDFKPKGKLWADDDRSVDYNEKLSFEAPPSIWSRIVNF GSGWGFWVSPSLFITSTHVIPQGAKEFFGVPIKQIQVHKSGEFCRLRFPKPIRTDVTGMI LEEGAPEGTVVTLLIKRSTGELMPLDARMGTHATMKIQGRTVGGGMGMLLTGSNAKSMDL GTTPGDCGCPIYKRGNDVVIGVHTAAARGGNTVICATQGGEGEATLEGGDSKGTYCGA PILGPGSAPKLSKTKFWRSSTAPLPPGTYEPAYLGGKDPRVKGGPSLQQVMRDQLKPT EPRGKPFKPSVLEAAKKTIINVLEQTIDPEKWSFAQACASLDKTTSSGHPHMRKNDCW NGESFTGKLADQASKANLMFEEGKNMTPVTTGALKDELVKTDKIYGKIKKRLLWGSDLAT MIKCARAFGGLMDELKAHCVTLPIRVGMNNNEDGPIIFEHSRYRYHYDDJSRNDSTQQ RAVLAAALEIMVKFSSEPHLAQVVAEDLLSPSVDVDGDFTISINEGLPSCTSDWNSI AHMLLICALSEVTNLSPDIIQANSLFSFYGDDELVSTDIKLDPEKLTAAKKSQLKFTR PDKTEGPLVISEDLDGLTFLRRTVTRDPAGWFGKLEQSSILRQMYWRGPNHEDPSESMI PHSQRPIQLMSLLGEAALHGPTFYSKISKLVIAELKEGGMDFYVPRQEPMFRWMRFSDLS TWEGDRNLAPSFVNEDGVE*</pre>		
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>contig00001.1.1 3 992 mat\_peptide protein p48
MKMASNDASAAAVANSNNDTAKSSSDKMFSNMAVTFKRALGARPKQPPPREIPQRPPRPP
TPELVKKIPPPPPNGEDEVVVSYSAKDGVSGLPELSTVRQPEETNTAFSVPPLNQRENRD
AKEPLTGTILEMWDGEIYHYGLYVERGLVLGVHKPPAAISLAKVELTPLSLFWRPVYTPQ

## Try your own favorite virus

- Get a complete unsegmented viral genome from GenBank – search "nucleotide" database for the virus you want with qualifier [organism] AND "complete genome"[title]
- Download the fasta record
- Try annotating it at VIGOR website
- Look at the .tbl file

