ICTV Webinar Detailed Agenda

March 11, 2025

- 10:00 AM 11:30 AM U.S. CDT (Central Daylight Time)
- 15:00 16:30 UTC (Coordinated Universal Time)

Speaker List

- Elliot Lefkowitz, ICTV Data Secretary
- Evelien Adriaenssens, ICTV Vice President
- Steve Powell, Outreach Coordinator
- Logan Mims, Software Developer
- Don Dempsey, Software Developer
- Eden Black, Outreach and Programmer

Introduction

- Welcome (Elliot Lefkowitz)
 - Funding
 - o Aims of the supporting NIH/NIAID grant
- What is the ICTV? (Evelien Adriaenssens)
 - The charge and history of the ICTV
 - o Executive Committee
 - Voting Membership
 - Statutes
 - o The International Code of Virus Classification and Nomenclature
 - Thanks to those providing financial support the ICTV

Intro to ICTV website & overview of resources for finding virus information

- Official ICTV Resources (Elliot Lefkowitz)
 - o Master Species List
 - What it is and what it provides
 - How to search
 - o Virus Metadata Resource
 - What it is and what it provides
 - How to search
 - o Taxonomy database
 - Single Db drives everything
 - Data dumps available in GitHub
 - https://github.com/ICTV-Virus-Knowledgebase/ICTVdatabase
- Tour of Resources (Steve Powell)
 - o (using either slides or live website: show each item, with one line description)
 - o Taxonomy Browser
 - Visual Taxonomy Browser
 - o Proposals

- o Find the Species
- Taxon Name Etymology
- Virus Properties Table
- Virus Diagrams
- Report Chapters
- o Help Videos
- o Google Search at bottom

In-depth description of resources

- Taxonomy Browsers (introduction by Elliot Lefkowitz)
 - Taxonomy Browser (Steve Powell)
 - Description
 - How to access
 - How to use
 - Hierarchical display of the taxonomy in collapsing/expanding rows
 - Help features
 - Straight-forward example(s)
 - Example(s) showing limitations
 - Taxon Details (History of the Taxon) pages
 - Future enhancements
 - o Additional links (or tabs) to other resources
 - Virus properties
 - Report chapters
 - VMR entries
 - Historical Taxonomy Browser
 - Overview
 - User can select different MSL releases for a search
 - Visual Taxonomy Browser (Logan Mims)
 - Why it was developed
 - Provides a visual representation of the taxonomy
 - Uses the same taxonomy as the Taxonomy Browser
 - Description
 - Where to access on the website
 - How to use
 - o Search
 - o Display
 - Release, font size, zoom, export, search
 - Meaning of "unassigned" category
 - Demonstration
 - Straight forward example(s)
 - Display allows for side-by-side comparisons
 - Can easily visualize relatedness when comparing two species
 - Example(s) showing limitations

- Pop-up (links)
- Export formats
- Help features
- Future enhancements
 - Jira list
- Find the Species
 - Why Find the Species was developed (Elliot Lefkowitz)

smallpox example

- How to use Find the Species (Eden Black)
 - Introduce and explain the search strategies
 - Where to access
 - o From the home page and from the Taxonomy dropdown list
 - How was Find the Species engineered to query the data?
 - Give examples of search algorithms
 - Exact match (e.g., COVID, measles, mumps)
 - Explanation of result output column headings
 - All words (e.g., herpes simplex, Epstein Barr)
 - Any word (e.g., Crimean Congo, other)
 - Contains (e.g., HPV, immuno, pox)
 - Other examples
- What is Find the Species? (Don Dempsey)
 - (show navigation to page; bring up page as background before detailing)
 - How it works
 - (run a search with VMR *AND* NCBI common name hits, so can point to parts as he describes them)
 - Databases
 - Future enhancements
 - Custom records
 - Additional databases
 - Ask the attendees if they know of any

Other Resources (Elliot Lefkowitz)

- Taxon Name Etymology
- Virus Properties
- Virus Diagrams
- ICTV Report
- Help Videos

Future Plans (Elliot Lefkowitz)

- Demarcation criteria database
- Find the Virus
- API

Questions and answers