

Screenshot Summary

- 67525_screenshot_1.mp4
 - 2.2.1 SCREEN: selecting CaseOLAP LIFT tool and inspecting biomedical literature and graph for analysis. **00:00-00:35**
- 67525_screenshot_2.mp4
 - 2.2.2 Open CaseOLAP LIFT protocol site, scroll through and highlight steps 1 through 3 being completed. **00:00-00:13**
- 67525_screenshot_3.mp4
 - 2.3.1. SCREEN: typing the Git clone command into the terminal and downloading the Know2BIO repository. **00:00-00:12**
- 67525_screenshot_4.mp4
 - 2.3.2. SCREEN: Run `create_edge_files.py` in the terminal and display progress of file downloads and graph extraction. **00:00-00:14**
- 67525_screenshot_5.mp4
 - 2.4.1. SCREEN: typing and running the `prepare_kgs.py` command in the terminal. **00:00-00:42**
- 67525_screenshot_6.mp4
 - 2.5.1. SCREEN: Command-line execution showing input files and final merged output being saved. **00:00-00:28**
 - 2.5.2. SCREEN: comprehensive graph created. **00:29-00:47**
- 67525_screenshot_7.mp4
 - 2.7.1. SCREEN: Talent typing and executing the filter script on the terminal. **00:00-00:34**
 - 2.7.2. SCREEN: Running the command and terminal output being obtained. **00:34-01:04**
- 67525_screenshot_8.mp4
 - 2.8.1. SCREEN: executing the prediction script using the command line. **00:00-00:34**
 - 2.8.2. SCREEN: Display of output path as the script runs. **00:34-00:53**
- 67525_screenshot_9.mp4
 - 2.9.1. SCREEN: opening a terminal and connecting to the RUGGED Docker container. **00:00-00:28**
- 67525_screenshot_10.mp4
 - 2.9.2. SCREEN: reattaching to the previously running container. **00:00-00:05**
 - 2.9.3. SCREEN: typing `cd /workspace/RUGGED` to enter the correct directory. **00:05-00:12**
 - 2.9.4. SCREEN: Terminal showing the current path as `/workspace/RUGGED` after directory navigation. **00:12-00:15**
- 67525_screenshot_11.mp4
 - 2.10.1. SCREEN: Display of active services required by RUGGED. **00:00-00:16**
- 67525_screenshot_12.mp4

- 2.10.2. SCREEN: Terminal execution of the `python rugged.py` command and the system launching RUGGED. **00:00-00:10**
- 2.11.1. SCREEN: User entering a query starting with 'query "What are the currently prescribed drugs classified as beta blockers, antiarrhythmic drugs, and antifibrotic drugs?"' and receiving a list of drug responses related to the disease. **00:00-00:38**
- 67525_screenshot_13.mp4
 - 2.13.1. SCREEN: User entering a search query and system returning a list of scientific articles, titles, and abstract snippets. **00:00-00:58**
 - 2.15.1. SCREEN: Typing “summarize” and saving a generated summary of the conversation to a text file. **00:58-01:28**
- 67525_screenshot_14.mp4
 - 2.14.1. SCREEN: Dialogue view showing user adjusting their line of questioning, with the system adapting prior responses to maintain contextual relevance. **00:00-00:15**
- 67525_screenshot_15.mp4
 - 2.14.2. SCREEN: Terminal showing edited Cypher query being entered and adjusted results retrieved from Neo4j. **00:00-00:35**
- 67525_screenshot_16.mp4
 - 2.15.2. SCREEN: scrolling through the summary file, making edits for clarity, and finalizing it. **00:00-00:18**
- 67525_screenshot_17.mp4
 - 2.16.1. SCREEN: Opening the log directory in a file browser or using command line to view log contents. **00:00-00:15**
- 67525_screenshot_18.mp4
 - SCREEN: Talent typing a predictive inquiry and RUGGED returning possible new drug targets with supporting links or scores. **00:00-00:27**