**Guide: Setting up Auto Feed Script on Bioreactors**

**Step 1:** Select the unit you want to setup the script for, and then select the vessel widget.

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**Step 2:** When you are in the vessel widget, navigate to the “Scripting” tab.

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**Step 3:** In the “Scripting” tab, click on the “Open editor” button.

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**Step 4:** This will direct you to the Script editor. You will now acquire the necessary script.

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**Step 5:** Utilize the shortcut icon in the taskbar to access the relevant Confluence page.



**Step 6:** This shortcut should bring you to this confluence page.

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**Step 7:** In the “Page Tree,” expand the “Eppendorf BRX” tab and select “CCD Scripts and Templates.”

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**Step 8:** Access the “Script Generator Automator” tab from there.

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**Step 9:** Now you should see the “Feed Script Form”.

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**Step 10:** Fill out the “Feed Script Form” with details such as the amount of “Feed A” and “Feed B” to be syphoned into the bioreactor on specific days.

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**Step 11:** Ensure your experiment is named, note the “Experiment ID,” and input the bioreactor IDs under “BRX ID.”

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**Step 12:** Submit the form using the “Submit” button at the bottom of the “Feed Script Form.”

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**Step 13:** The form will appear under the download’s icon.

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**Step 14:** Click on the text (.txt) file to open it.

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**Step 15:** Make sure you select the whole text file and copy it to your clipboard (Ctrl + c).

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**Step 16:** Return to the script editor you opened (from step: 4) and paste the whole script from your clipboard (Ctrl + v).

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**Step 17:** When pasted hit the “Compile” button. The message output box should say: “Compilation successful”, you can now hit the “Confirm” button.

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**Step 18:** You are now ready to start the inoculation timer. Confirm the system status in the event box reads “Scale is busy” and “Automated feeding initialized for unit#: Unit 4.”

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