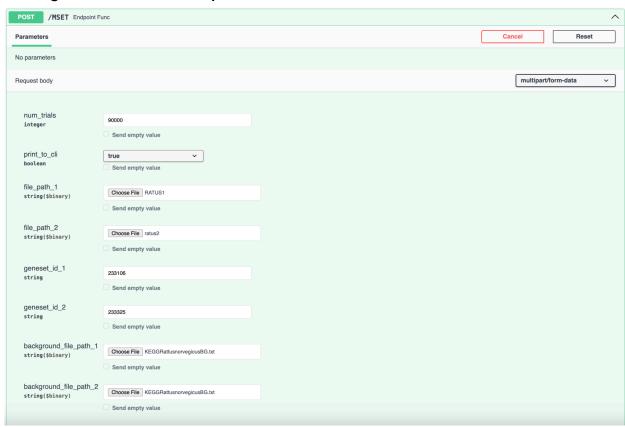
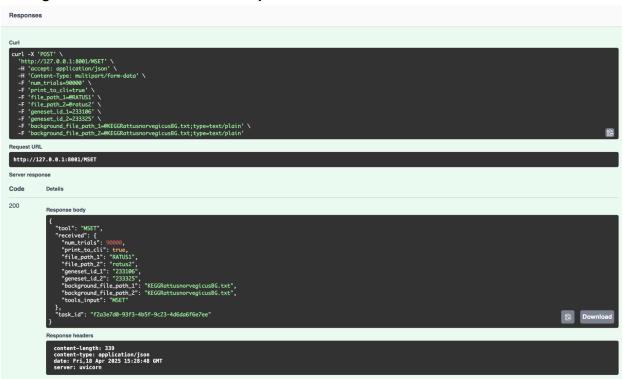
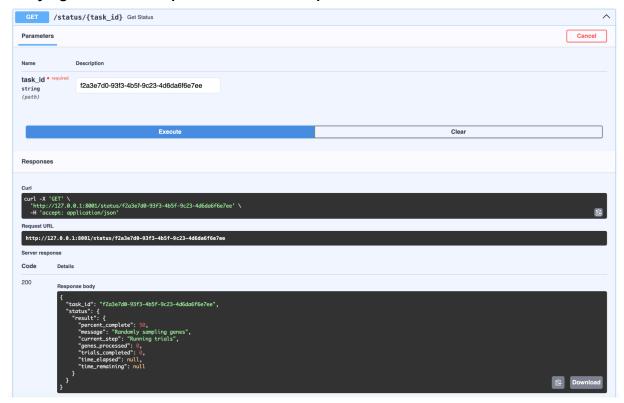
## Loading the MSET tool with inputs



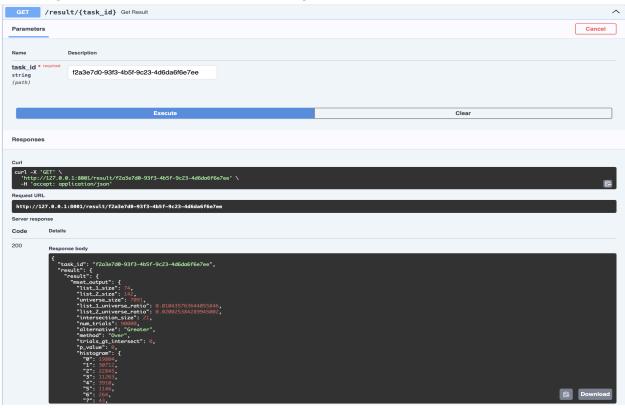
#### Getting the task id from the loaded input



#### Verifying the status endpoint to retrieve computation status



### Checking the result for the MSET tool through the fastapi endpoint



# **Error logging for invalid input**

```
[STATUS] 10% - Retrieving gene sets (Loading data)

Plugin result:
{'Error': 'Please provide geneset_id or file_path for group 1 and group 2'}

A Histogram data not found or not in expected format.
```