

MCP Workshop: Connect to an Existing Client (Postman)

In this guide, you'll learn how to connect your running MCP servers to a real-world client application. We'll use [Postman](#), which has built-in MCP support, to act as our client. This will allow you to explore your server's tools and resources interactively, just as an AI assistant would.

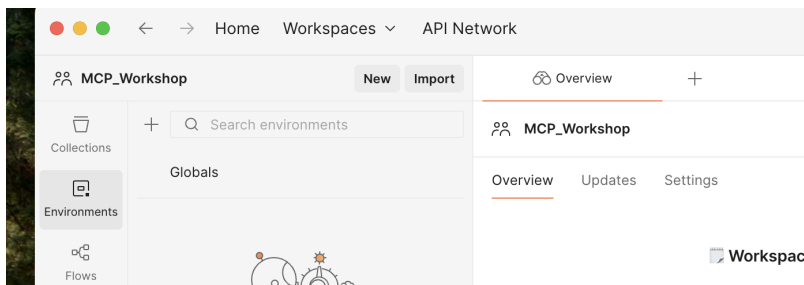
Prerequisites

If you haven't already, download and install the Postman desktop app.

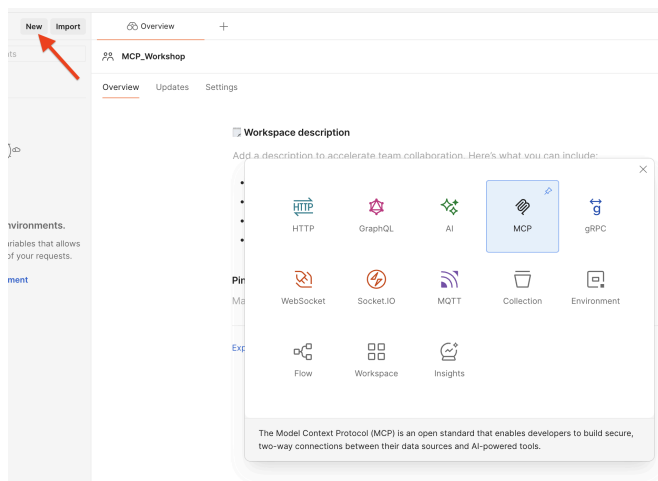
Download Link: <https://www.postman.com/downloads/>

Connect to the Servers

1. Let's connect Postman to the `SciLifeLab_agent_workshop/Section_2_MCP/existing_clients_and_servers/postman_mcp_basic_server.py` file.
2. **Open Postman:** Launch the Postman application and create a workspace if prompted (e.g., `MCP_Workshop`).



3. **Create New MCP Request:** In your workspace, click the **New** button.



4. **Rename the Request:** Click the default name (e.g., **Untitled Request**) and change it to something descriptive, like **My_MCP_Server**.



5. **Load Configuration:**

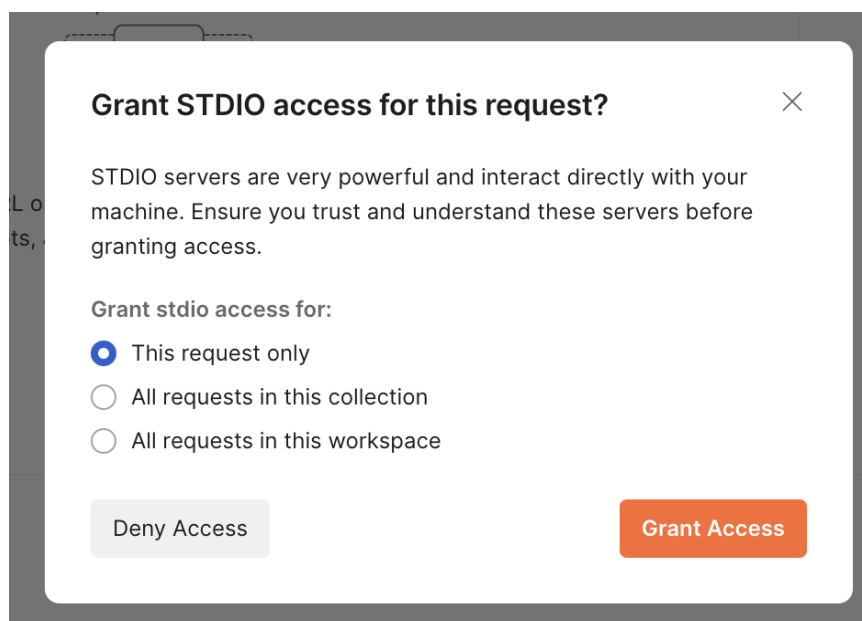
- Find the configuration file in your workshop materials:
SciLifeLab_agent_workshop/Section_2_MCP/existing_clients_and_servers/postman_mcp_basic_server_config.json
- Open this file in a text editor (e.g., VS Code or Notepad). Make sure to update it by adding the full path in the description.
- **Copy its entire contents** to your clipboard.

6. **Paste and Connect:**

- Return to Postman and paste the copied JSON content into the main configuration window (labeled **Enter command or paste JSON config**). Refer to the image under point 3 for guidance.
- Click the **Connect** button.

7. **Grant Access:** Postman will show a security pop-up asking for permission to connect

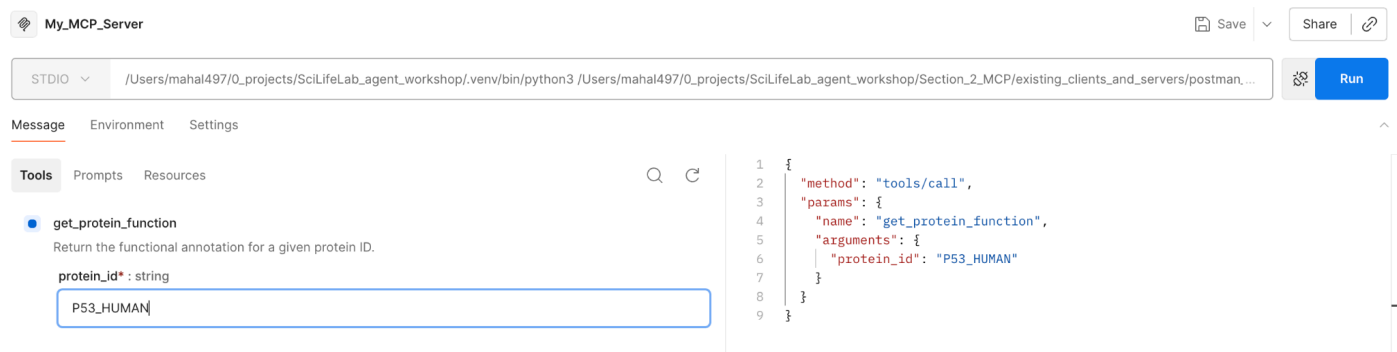
- Select **This request only**.
- Click **Grant Access**.



8. **Run a Tool:** You are now connected!

- Select the **Tools** tab in Postman.
- Click on the **get_protein_function** tool.

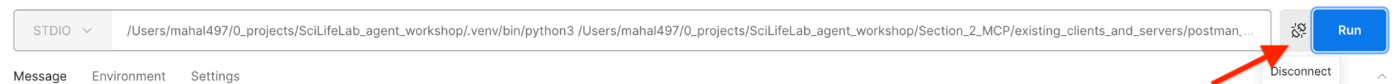
- In the `protein_id*` (required) field, type `P53_HUMAN`.
- Click the **RUN** button.



9. **Observe the Response:** Look at the **Response** panel at the bottom of the screen. You should see the JSON response from your server containing the protein's function!



10. **Disconnect:** Once you're done, click the **Disconnect** button at the top.



11. Now we'll use the `SciLifeLab_agent_workshop/Section_2_MCP/existing_clients_and_servers/postman_mcp_advanced_server.py` file for the advanced server. Repeat the process from step **one**, but in step **five**, use the `SciLifeLab_agent_workshop/Section_2_MCP/existing_clients_and_servers/postman_mcp_advanced_server_config.json` file instead.

12. **Explore!** You are now connected to the advanced server.
- Click the **Tools** tab. Notice all the new tools: `find_protein`, `generate_hypothesis`, `stream_analysis_log`, etc.
 - Click the **Resources** tab. You'll see the resources defined on the server, like `dataset://proteins`.

- **Try it:** Run the `find_protein` tool with the `protein_name` set to `p53`. Observe the "elicitation" response it returns, asking you to choose between human and mouse.

The screenshot shows the MCP client interface with the 'Response' tab selected. On the left, a list of messages shows 'tools / stream_analysis_log' at 16:46:40.457. The main panel displays a JSON response:

```

1 {
2   "content": [
3     {
4       "type": "text",
5       "text": "Initializing analysis...\nSequence loaded (393 aa)\nDetecting domains...\nTransactivation domain found (aa 1-42)\nDNA-binding domain found (aa 102-292)\nTetramerization domain found (aa 323-356)\nSearching for variants...\nSite R248Q found\nGenerating report...\nAnalysis complete."
6     },
7   ],
8   "structuredContent": {
9     "result": "Initializing analysis...\nSequence loaded (393 aa)\nDetecting domains...\nTransactivation domain found (aa 1-42)\nDNA-binding domain found (aa 102-292)\nTetramerization domain found (aa 323-356)\nSearching for variants...\nSite R248Q found\nGenerating report...\nAnalysis complete."
10    },
11   "isError": false
12 }

```

The screenshot shows the MCP client interface with the 'Notifications' tab selected. It displays a list of messages with 'info' icons:

- {"method":"notifications/message","params":{"level":"info","data":"Step 10/10: Analysis complete."}} 16:46:40.155
- {"method":"notifications/message","params":{"level":"info","data":"Step 9/10: Generating report..."}} 16:46:39.854
- {"method":"notifications/message","params":{"level":"info","data":"Step 8/10: Site R248Q found"}} 16:46:39.552
- {"method":"notifications/message","params":{"level":"info","data":"Step 7/10: Searching for variants..."}} 16:46:39.251
- {"method":"notifications/message","params":{"level":"info","data":"Step 6/10: Tetramerization domain found (aa 323-356)"}} 16:46:38.951
- {"method":"notifications/message","params":{"level":"info","data":"Step 5/10: DNA-binding domain found (aa 102-292)"}} 16:46:38.648
- {"method":"notifications/message","params":{"level":"info","data":"Step 4/10: Transactivation domain found (aa 1-42)"}} 16:46:38.347

The screenshot shows the MCP client interface with the 'Tools' tab selected. It lists the following tools:

- 2. register_client**
Register a client to receive notifications.
- 3. trigger_notification**
Trigger a notification to all registered clients.
- 4. find_protein**
Find proteins by name. Demonstrates elicitation by asking for clarification when multiple matches are found.
- generate_hypothesis** (selected)
Generate a research hypothesis for a protein. Demonstrates sampling by providing a prompt that should be completed by an LLM.

On the right, a JSON snippet shows the configuration for the 'generate_hypothesis' tool:

```

1 {
2   "method": "tools/call",
3   "params": {
4     "name": "generate_hypothesis",
5     "arguments": {
6       "protein_id": "P53_HUMAN"
7     }
8   }
9 }

```

Workshop Complete

You've successfully used a real-world client application to connect to and interact with your custom MCP servers. This demonstrates how any MCP-compliant application (like an AI assistant) can discover and use the tools you build.

Happy learning!