# **Databricks Workflows Exercise**

In this exercise you will create your own Databricks Workflow to orchestrate the ETL pipeline.



🦍 If you need any help with this exercise, please raise your hand and we will gladly assist you! 🥼



#### Step 1: Familiarize yourself with the ETL pipeline

The pipeline consists of three steps, represented by three Databricks notebooks.

- prepare\_bronze\_table.py
- 2. prepare silver table.py
- prepare\_gold\_table.py

These notebooks follow the medallion architecture where data flows through bronze, silver and gold tables. The first notebook loads the raw data (for the sake of this workshop, this is just a copy of another table). The second notebook cleans the data and the third notebook prepares the data for LLM processing.



### Step 2: Get familiar with the Workflow UI

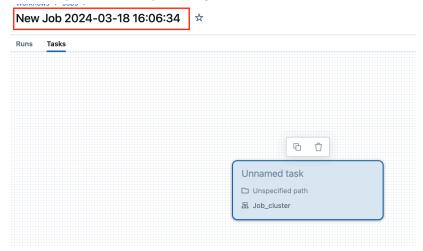
In the side panel of the workspace, navigate to **workflows**. This page has three sections:

- 1. **Jobs**: here, you will see all jobs that have been defined.
- 2. **Job runs**: here, you can see all previous runs of the jobs.
- 3. **Delta Live Tables**: here, you can see all DLT Pipelines that have been defined.

### Step 3: Build your own workflow

1. In the **Jobs** section, click on **Create job** on the top right, so start setting up the new job.

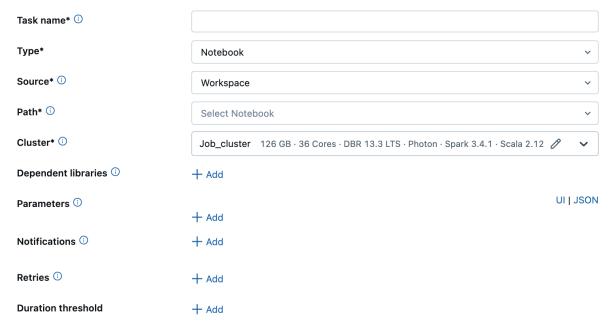
2. Give the job a name by clicking on the current name:



Name the workflow your-username>\_etl\_job. For example: amIdbricks-10\_etl\_job

## Step 4: Define the first task

Creating a job already created a first task with a configuration block:



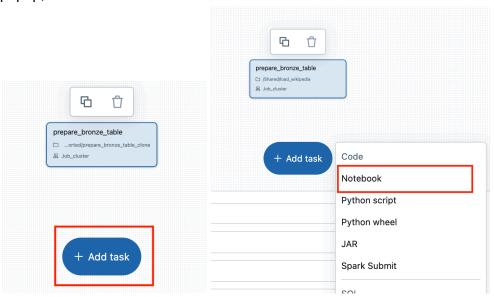
Now, we will configure this task:

- 1. First, set the Task name to prepare\_bronze\_table
- 2. Type: Notebook
- 3. Source: Workspace
- 4. Path: <your-home-folder>/notebooks/Workflows Exercise/prepare\_bronze\_table.py
- 5. Cluster: <the all purpose cluster you created in the previous exercise>
- 6. Leave all other settings blank / at their default settings

Complete the task definition by clicking on **Create task** in the bottom right corner.

### Step 4: Define the second task

You can add another task by clicking on the **Add task** button in the workflow lineage view. In the popup, select **Notebook.** 



Configure the task with the following settings:

- 1. Task name: prepare\_silver\_table
- 2. Type: Notebook
- 3. Source: Workspace
- 4. Path: <your-home-folder>/notebooks/Workflows Exercise/prepare\_silver\_table.py
- 5. Cluster: <the all purpose cluster you created in the previous exercise>
- 6. Depends on: prepare\_bronze\_table
- 7. Run if dependencies: All succeeded
- 8. Leave all other settings blank / at their default settings

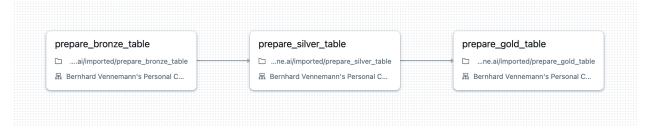
Complete the task definition by clicking on Create task in the bottom right corner.

Notice that this task is now dependent on the **prepare\_bronze\_table** task and only runs if the previous task was executed successfully.

### Step 4: Define the third task

Repeat the previous step to create a task **prepare\_gold\_table** based on the notebook **prepare\_gold\_table.py** 

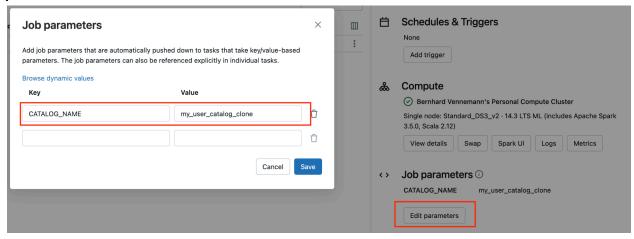
After completing this step, your workflow should look like this:



### Step 5: Set the Job parameter

The notebooks that make up the three tasks expect a parameter **CATALOG\_NAME** that defines which catalog should host the newly created bronze, silver and gold tables. Parameters can be defined on the task level and the job level. Since all tasks will share this parameter, we will define the **CATALOG\_NAME** parameter on the job level.

You can do so my clicking on the button **Edit parameters** in the right hand panel in the **Job parameters** section:



- 1. Define a new parameter with the Key **CATALOG\_NAME**
- 2. For the Value, provide the **name of the catalog that you created** in the previous exercise.
- 3. Confirm with Save

### Step 6: Run the workflow

Finally, we are ready to run the workflow.

- 1. Run the workflow by clicking on **Run now** in the top right corner of the screen.
- 2. Navigate to **Runs** (under the workflow name) and click on the current run (link in the column **Start time**)
- You will see the three tasks and the status of the run (starting with the first task and working its way through the pipeline until the gold table has been computed). This should take a couple of minutes.

- 4. You can click on the individual tasks to see how the notebooks are being executed.
- 5. When all tasks have been completed, navigate to your Catalog and inspect the newly created schemas and tables. To do so, click on **Catalog** in the side panel which brings you to the Catalog Explorer. Navigate to your catalog and have a look at the newly created schemas and tables. When you select a table by clicking on on its name, you can have a look at some **Sample Data** and other useful information, such as the **Lineage**.

