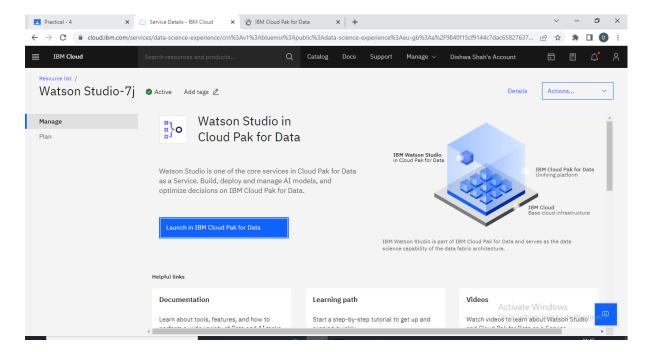
Institute of Computer Technology B. Tech Computer Science and Engineering Sub: Cognitive Computing (2CSE70E23)

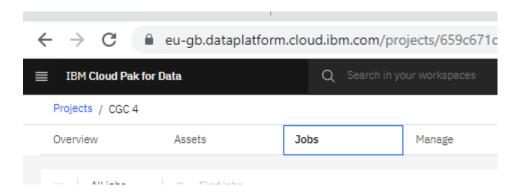
PRACTICAL 4

Jesal got a task from his team that he needs to deploy a machine learning model for the iris flower type. that can detect flower type as per the petal height and width. Use Watson Machine learning and Watson studio to train and deploy the model.

Go to IBM Cloud and search for Machine Learning. Go to Resource List and Launch in IBM Cloud Pak for Data.

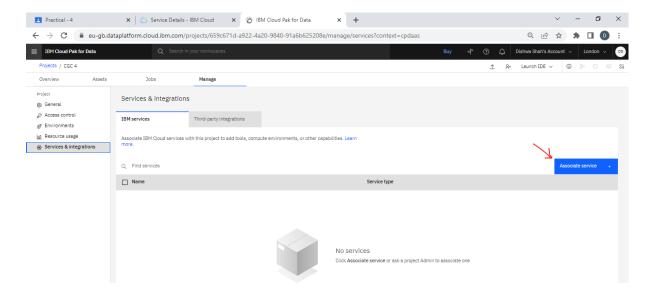


Create a new empty project and go to Jobs.

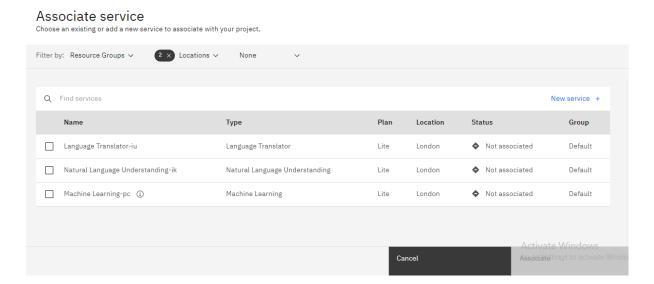


Jobs: Information regarding if there are any running jobs or not Manage: Project managing things

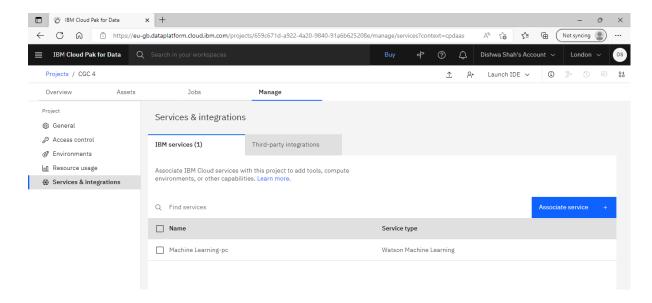
Go to Manage tab and go to Services and integrations and click on Associate services.



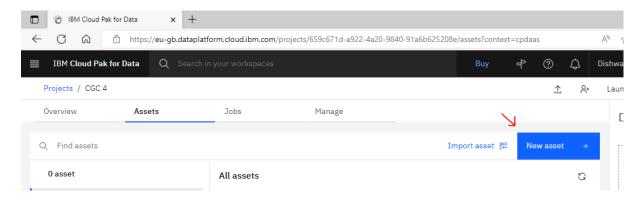
Select Machine Learning and Associate.



Now, click on the associated service.



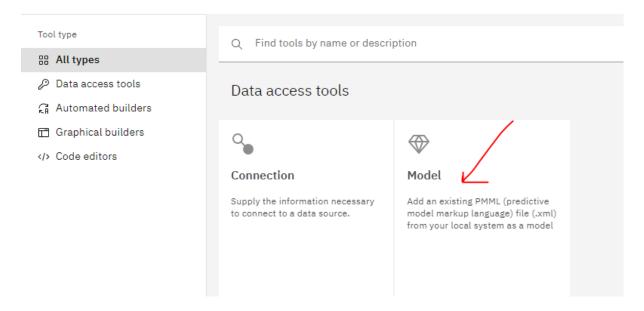
Go to Assets tab and create a New asset.



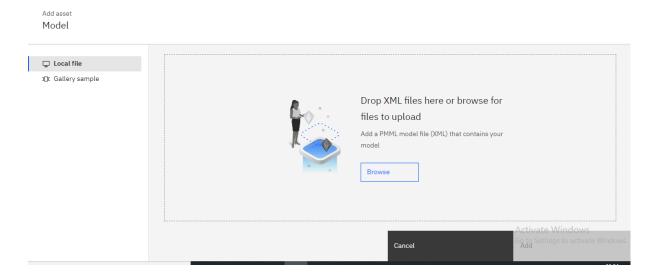
Select Model since we already have a pre-built model.

New asset

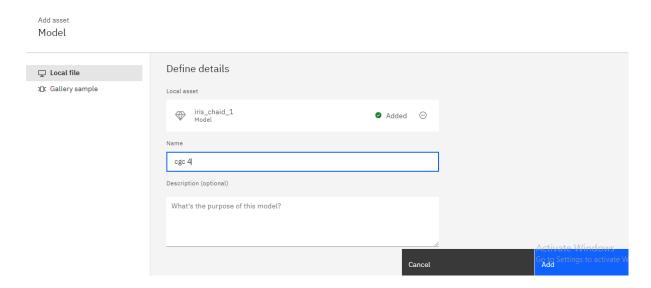
Select a tool based on what type of asset you want and how you want to work.



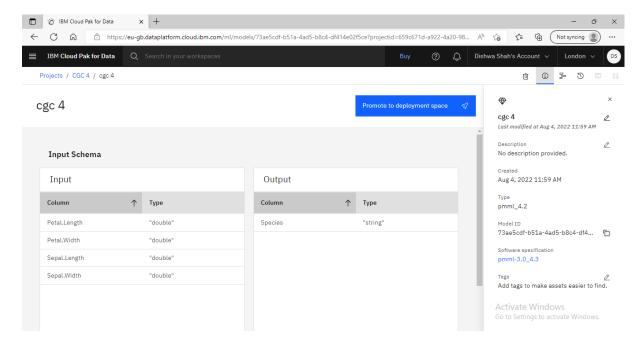
Browse the xml file and upload it.



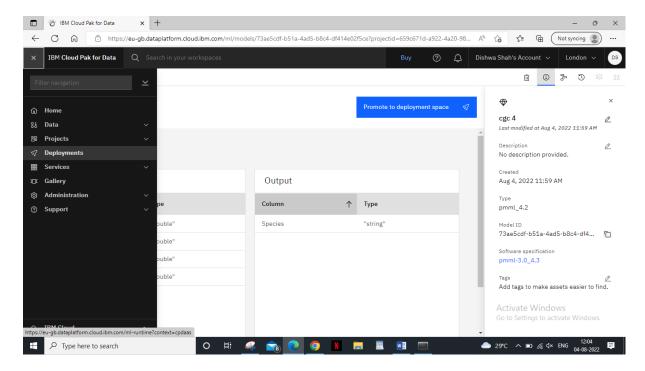
As we can see, the model is successfully validated. Give it a name and click on Add.



Now, we want to deploy the model. Watson Studio already has a pre-built deployment space for our model. Click on Promote to deployment space.



OR Click on the burger menu and select Deployments.



The deployment space is now ready.

The space is ready

Close this notification to resume your work. Click Deployments in the navigation pane to view and access the new space.

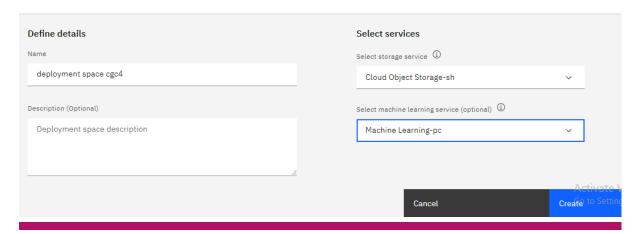
Step 1 of 1. Creating deployment space.

Close

Add details and select your ML model:

New deployment space

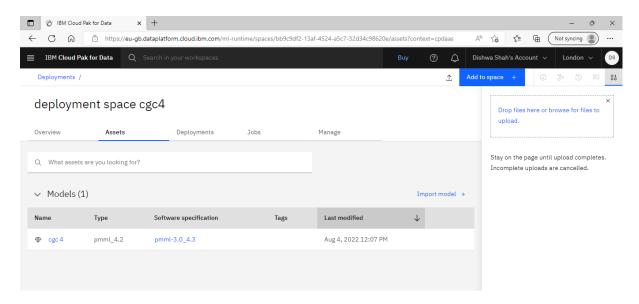
Use a space to collect assets in one place to create, run, and manage deployments



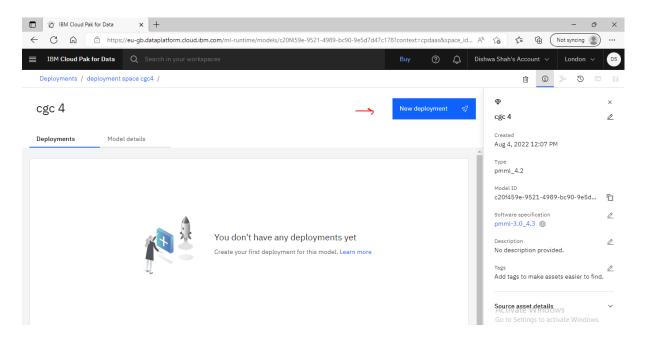
Click on the generated link to go to your deployment space:



Go to Assets and click on the model.

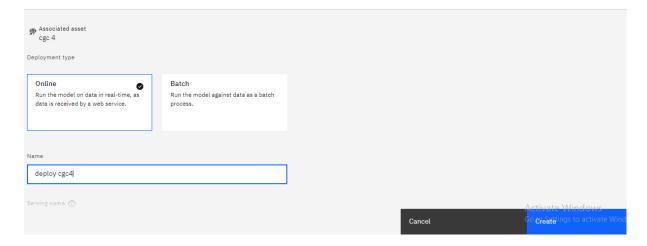


Click on New deployment.

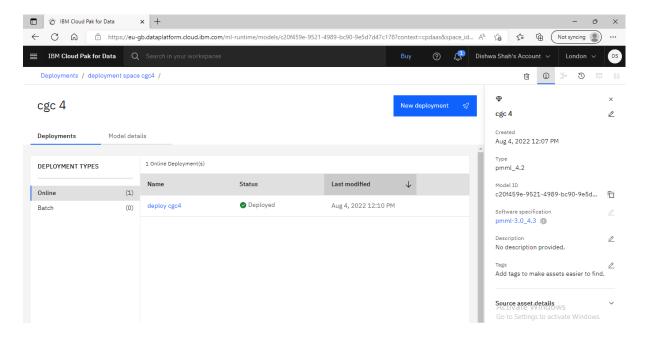


Select Online Deployment type.

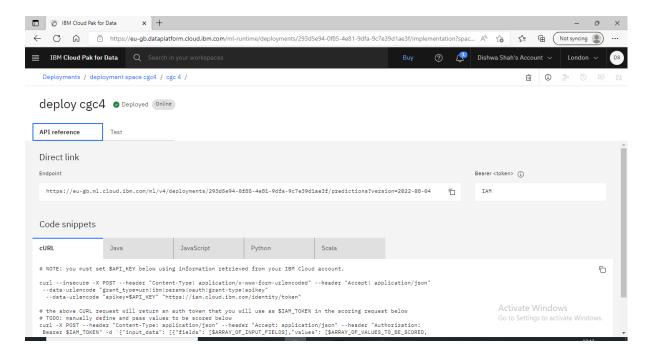
Create a deployment



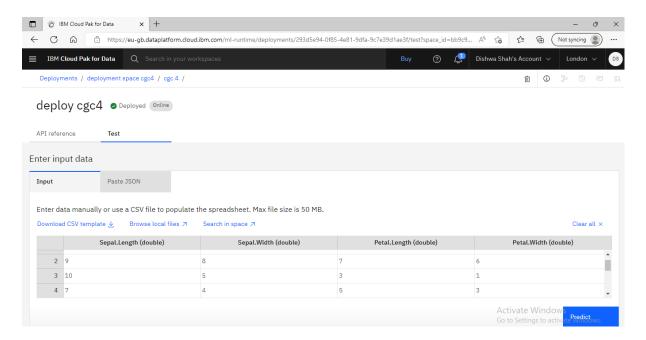
Click on the newly created deployment.



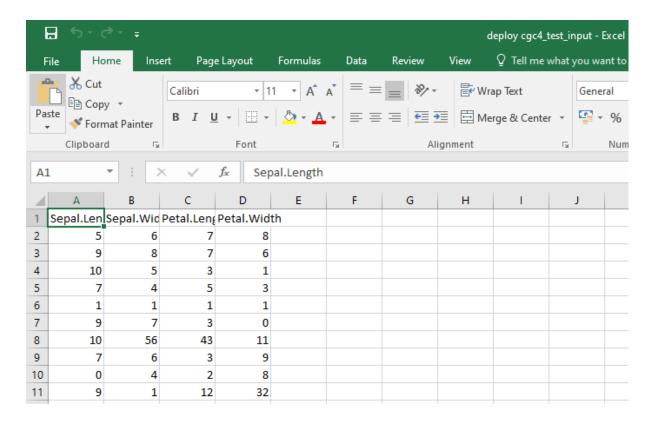
You can see the direct link here which you can use to connect to other tools. Also, various code snippets in different languages are available if you want to work in that language.



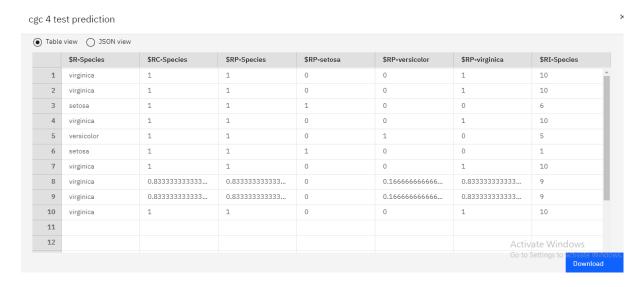
Go to Test and take 10 cases.



Click on Predict. You can download the CSV template.



Probability scores are displayed according to the species when you click on Predict. In the first column, it gives the predicted species.



You can also go to the JSON view if you want to work with it.

cgc 4 test prediction

You can download the JSON file as well if you want.

```
C:\Users\admin\Downloads\deploy cgc4_test_result.json - Notepad++
File Edit Search View Encoding Language Settings Tools Macro Run Plugins
🕞 🖆 🗎 🖺 🥦 😘 📤 | 🔏 🐚 🖿 🕇 🕦 🖊
 📑 abda 2.txt 🖾 🔚 deploy cgc4_test_result.json 🔀
       目(
   2
             "fields": [
   3
                  "$R-Species",
   4
                  "$RC-Species",
   5
                  "$RP-Species",
   6
                  "$RP-setosa",
   7
                  "$RP-versicolor",
   8
                  "$RP-virginica",
   9
                  "$RI-Species"
  10
             1,
  11
             "values": [
  12
  13
                       "virginica",
  14
                       1,
  15
                       1,
                       0,
  16
  17
                       0,
  18
                       "10"
  19
  20
                  1,
  21
                       "virginica",
  22
  23
  24
                       1,
  25
                       0,
  26
  27
                       "10"
  28
  29
                  1,
  30
                       "setosa",
  31
  32
                       1,
  33
                       1,
JSON file
```

This is the predicted output generated:

```
"$RP-virginica",
        "$RI-Species"
"values": [
                "virginica",
                1,
                1,
                0,
                0,
                1,
                "10"
        ],
                "virginica",
                1,
                1,
                0,
                0,
                1,
                "10"
        ],
                "setosa",
                1,
                1,
                1,
                0,
                0,
                "6"
        ],
        [
                "virginica",
                1,
                1,
                0,
                0,
                1,
                "10"
        ],
                "versicolor",
                1,
                1,
                0,
                1,
                0,
                "5"
```

```
],
[
      "setosa",
       1,
       1,
       1,
      0,
      0,
      "1"
],
[
      "virginica",
       1,
       1,
      0,
      0,
       1,
      "10"
],
[
      "virginica",
      0.8333333333333333,
      0.8333333333333333,
      0,
      0.166666666666667,
      0.8333333333333333,
      "9"
],
      "virginica",
      0.8333333333333333,
      0,
      0.166666666666666667,
      ],
      "virginica",
       1,
       1,
      0,
      0,
       1,
      "10"
]
```

]

}