ML Lineage Workflows

August 2, 2024

• Required Versions:

- Snowflake: 8.27 or higher and Enterprise Edition or higher

Snowflake ML Python: 1.6.0 or higher
Snowpark Python: 1.21.0 or higher

• Last Updated: 8/1/2024

ML Lineage workflows

This notebook showcases various machine learning workflows, delving into the lineage of each process. It highlights essential features of Snowflake's ML, including Snowflake Feature Store, Dataset, ML Lineage, Snowpark ML Modeling and Snowflake Model Registry.

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1. Set up environment

1.1 Connect to Snowflake

Let's start with setting up our test environment. We will create a session and a schema. The schema FS_DEMO_SCHEMA will be used as the Feature Store. It will be cleaned up at the end of the

demo. You need to fill the connection_parameters with your Snowflake connection information. Follow this **guide** for more details about how to connect to Snowflake.

```
[]: from snowflake.snowpark import Session, context, exceptions
     try:
         # Retrieve active session if in Snowpark Notebook
         session = context.get_active_session()
     except exceptions.SnowparkSessionException:
         # ACTION REQUIRED: Need to manually configure Snowflake connection if using
      \hookrightarrow Jupyter
         connection_parameters = {
             "account": "<your snowflake account>",
             "user": "<your snowflake user>",
             "password": "<your snowflake password>",
             "role": "<your snowflake role>",
             "warehouse": "<your snowflake warehouse>",
             "database": "<your snowflake database>",
             "schema": "<your snowflake schema>",
         }
         session = Session.builder.configs(connection_parameters).create()
         print(session)
     assert session.get_current_database() != None, "Session must have a database_
      ⇒for the demo."
     assert session.get_current_warehouse() != None, "Session must have a warehouse_
      ⇔for the demo."
```

<snowflake.snowpark.session.Session: account="ax_test_qa3", role="ACCOUNTADMIN",
database="LINEAGE_DEMO_DB", schema="PUBLIC", warehouse="AX_XL">

```
[]: # The schema where Feature Store lives.
FS_DEMO_SCHEMA = "FEATURE_STORE"
# The schema model lives.
MODEL_DEMO_SCHEMA = "MODEL_SCHEMA"

# Make sure your role has CREATE SCHEMA privileges or USAGE privileges on the_
schema if it already exists.
session.sql(f"CREATE OR REPLACE SCHEMA {FS_DEMO_SCHEMA}").collect()
session.sql(f"CREATE OR REPLACE SCHEMA {MODEL_DEMO_SCHEMA}").collect()
```

[]: [Row(status='Schema MODEL_SCHEMA successfully created.')]

1.2 Select your example

We have prepared some examples that you can find in our open source repo. Each example contains the source dataset, feature view and entity definitions which will be used in this demo. ExampleHelper (included in snowflake-ml-python) will setup everything with simple APIs and you don't have to worry about the details.

load_example() will load the source data into Snowflake tables. In the example below, we are using the "wine_quality_features" example. You can replace this with any example listed above. Execution of the cell below may take some time depending on the size of the dataset.

```
[]: from snowflake.ml.feature_store.examples.example_helper import ExampleHelper
     example_helper = ExampleHelper(session, session.get_current_database(),_
       →FS_DEMO_SCHEMA)
     source tables = example helper.load example('wine quality features')
     session.table(source_tables[0]).limit(5).to_pandas()
[]:
        WINE_ID
                  FIXED_ACIDITY
                                  VOLATILE_ACIDITY
                                                     CITRIC_ACID
                                                                   RESIDUAL_SUGAR
     0
              1
                             7.4
                                               0.70
                                                             0.00
                                                                               1.9
     1
              2
                            7.8
                                               0.88
                                                             0.00
                                                                               2.6
     2
              3
                            7.8
                                               0.76
                                                                               2.3
                                                             0.04
     3
              4
                            11.2
                                               0.28
                                                             0.56
                                                                               1.9
     4
              5
                             7.4
                                               0.70
                                                             0.00
                                                                               1.9
        CHLORIDES
                    FREE_SULFUR_DIOXIDE
                                          TOTAL_SULFUR_DIOXIDE
                                                                              PH \
                                                                  DENSITY
     0
            0.076
                                      11
                                                              34
                                                                   0.9978
                                                                            3.51
            0.098
     1
                                      25
                                                              67
                                                                   0.9968
                                                                            3.20
     2
            0.092
                                      15
                                                              54
                                                                   0.9970
                                                                            3.26
     3
            0.075
                                      17
                                                              60
                                                                   0.9980
                                                                            3.16
     4
            0.076
                                      11
                                                              34
                                                                   0.9978
                                                                           3.51
        SULPHATES
                    ALCOHOL
                              QUALITY
     0
             0.56
                        9.4
                                    5
             0.68
                                    5
     1
                        9.8
     2
                                    5
             0.65
                        9.8
     3
             0.58
                        9.8
                                    6
     4
             0.56
                        9.4
                                    5
```

2. Feature View Lineage

Create a new feature store and register and entities and feature views. More details on feature store APIs can be found here). For the detailed workflow of feature store refer to the notebook here

```
[]: from snowflake.ml.feature_store import (
    FeatureStore,
    FeatureView,
    Entity,
```

```
CreationMode
fs = FeatureStore(
    session=session,
    database=session.get_current_database(),
    name=FS_DEMO_SCHEMA,
    default_warehouse=session.get_current_warehouse(),
    creation_mode=CreationMode.CREATE_IF_NOT_EXIST,
)
all_entities = []
for e in example_helper.load_entities():
    entity = fs.register_entity(e)
    all_entities.append(entity)
all_feature_views = []
for fv in example_helper.load_draft_feature_views():
    rf = fs.register_feature_view(
        feature_view=fv,
        version='1.0'
    all_feature_views.append(rf)
fs.list_feature_views().select('name', 'version', 'desc', 'refresh_freq').show()
```

Query Lineage Query the upstream lineage of the feature views we just created.

```
[]: for fv in all_feature_views:
    print("Upstream Lineage of feature view '" + fv.name + "'")
    print(fv.lineage(direction='upstream'))
```

LineageNode.lineage() is in private preview since 1.5.3. Do not use it in production.

```
Lineage.trace() is in private preview since 1.16.0. Do not use it in production.

Upstream Lineage of feature view 'EXTRA_WINE_FEATURES'
[LineageNode(
   name='LINEAGE_DEMO_DB.FEATURE_STORE.WINEDATA',
   version='None',
   domain='table',
   status='ACTIVE',
   created_on='2024-08-01 22:44:14'
)]

Upstream Lineage of feature view 'WINE_FEATURES'
[LineageNode(
   name='LINEAGE_DEMO_DB.FEATURE_STORE.WINEDATA',
   version='None',
   domain='table',
```

3. Training Data Lineage

status='ACTIVE',

)]

Next step in ML workflows will be generating training data that is needed to train the model. There are 2 ways to generate training data. 1. Using feature views. 2. Using source tables directly.

3.1 Training Data from Feature views

created_on='2024-08-01 22:44:14'

Lets explore the workflow of creating training data sets using the feature views.

Retrieve some metadata columns that are essential when generating training data.

```
[]: label_cols = example_helper.get_label_cols()
    timestamp_col = example_helper.get_training_data_timestamp_col()
    excluded_cols = example_helper.get_excluded_cols()
    join_keys = [key for entity in all_entities for key in entity.join_keys]
```

Create a spine dataframe that's sampled from source table.

```
O 6 544
1 5 978
```

```
2
             5
                     679
3
             5
                    1459
             5
4
                        5
. .
             6
                     508
507
508
             6
                     624
509
             5
                    1132
             5
510
                     511
             5
511
                    1568
```

[512 rows x 2 columns]

3.1.1 Dataset as training data Snowflake Dataset generated from feature views created above. Dataset is a readonly objects helps in reproducability of the ML model.

Query Lineage Query Upstream lineage of the dataset we just generated.

```
[]: my_dataset.lineage(direction="upstream")
[]: [LineageNode(
       name='LINEAGE DEMO DB.FEATURE STORE.WINEDATA',
       version='None',
       domain='table',
       status='ACTIVE',
       created_on='2024-08-01 22:44:14'
     ),
     FeatureView( name=EXTRA WINE FEATURES, entities=[Entity(name=WINE,
     join_keys=['WINE_ID'], owner=None, desc=Wine ID column.)],
     _feature_df=<snowflake.snowpark.dataframe.DataFrame object at 0x1711bc3d0>,
     _timestamp_col=None, _desc=Static feature view about wine quality which never
    refresh., _infer_schema_df=<snowflake.snowpark.dataframe.DataFrame object at
     Ox1711fd910>, _query=SELECT "WINE_ID", "SULPHATES", "ALCOHOL" FROM
     "LINEAGE_DEMO_DB".FEATURE_STORE.winedata, _version=1.0,
     _status=FeatureViewStatus.STATIC, _feature_desc=OrderedDict([('SULPHATES', ''),
     ('ALCOHOL', '')]), _refresh_freq=None, _database=LINEAGE_DEMO_DB,
     _schema=FEATURE_STORE, _warehouse=None, _refresh_mode=None,
```

```
lineage node name=LINEAGE DEMO DB.FEATURE STORE.EXTRA WINE FEATURES,
     _lineage_node_domain=feature_view, _lineage_node_version=1.0,
     _lineage_node_status=None, _lineage_node_created_on=None,
     _session=<snowflake.snowpark.session.Session: account="ax_test_qa3",
     role="ACCOUNTADMIN", database="LINEAGE_DEMO_DB", schema="MODEL_SCHEMA",
     warehouse="AX XL">),
     FeatureView(_name=WINE_FEATURES, _entities=[Entity(name=WINE,
     join keys=['WINE ID'], owner=None, desc=Wine ID column.)],
     _feature_df=<snowflake.snowpark.dataframe.DataFrame object at 0x1711a1550>,
     _timestamp_col=None, _desc=Managed feature view about wine quality which
    refreshes everyday., _infer_schema_df=<snowflake.snowpark.dataframe.DataFrame
     object at 0x1711a12b0>, _query=SELECT "WINE_ID", "FIXED_ACIDITY", "CITRIC_ACID",
     "CHLORIDES", "TOTAL_SULFUR_DIOXIDE", "PH", ("FIXED_ACIDITY" * "CITRIC_ACID") AS
     "MY NEW FEATURE" FROM "LINEAGE DEMO DB".FEATURE STORE.winedata, _version=1.0,
     _status=FeatureViewStatus.ACTIVE, _feature_desc=OrderedDict([('FIXED_ACIDITY',
     ''), ('CITRIC_ACID', ''), ('CHLORIDES', ''), ('TOTAL_SULFUR_DIOXIDE', ''),
     ('PH', ''), ('MY_NEW_FEATURE', '')]), _refresh_freq=1 day,
     _database=LINEAGE_DEMO_DB, _schema=FEATURE_STORE, _warehouse=AX_XL,
     _refresh_mode=INCREMENTAL, _refresh_mode_reason=None, _owner=ACCOUNTADMIN,
    _lineage_node_name=LINEAGE_DEMO_DB.FEATURE_STORE.WINE_FEATURES,
     _lineage_node_domain=feature_view, _lineage_node_version=1.0,
    _lineage_node_status=None, _lineage_node_created_on=None,
     session=<snowflake.snowpark.session.Session: account="ax test qa3",
     role="ACCOUNTADMIN", database="LINEAGE_DEMO_DB", schema="MODEL_SCHEMA",
     warehouse="AX XL">)]
    Query the downstream of feature views we used to create the dataset.
[]: for fv in all_feature_views:
         print("Downstream Lineage of feature view '" + fv.name + "'")
         print(fv.lineage(direction='downstream'))
    Downstream Lineage of feature view 'EXTRA_WINE_FEATURES'
    [Dataset(
      name='LINEAGE_DEMO_DB.FEATURE_STORE.MY_DATASET',
      version='4.0',
    )]
    Downstream Lineage of feature view 'WINE_FEATURES'
    [Dataset(
      name='LINEAGE DEMO DB.FEATURE STORE.MY DATASET',
      version='4.0',
    )]
```

_refresh_mode_reason=None, _owner=ACCOUNTADMIN,

3.1.2 Tables as Training data Alternatively, you can create a regular table as a dataset from feature views. The downside is that tables are mutable, so reproducibility cannot be guaranteed.

```
[]: my_table_data = fs.generate_dataset(
    name="my_table_dataset",
    spine_df=spine_df,
    features=all_feature_views,
    version="4.0",
    spine_timestamp_col=timestamp_col,
    spine_label_cols=label_cols,
    exclude_columns=excluded_cols,
    desc="This is the dataset joined spine dataframe with feature views",
    output_type="table"
)
```

Query Lineage You can also explore lineage of generated table from the feature view. You filter the results to see just table entities.

```
[]: for fv in all_feature_views:
         print("Downstream Lineage of feature view '" + fv.name + "'")
         print(fv.lineage(direction='downstream', domain_filter=["table"]))
    Downstream Lineage of feature view 'EXTRA_WINE_FEATURES'
    [LineageNode(
      name='LINEAGE_DEMO_DB.FEATURE_STORE.MY_TABLE_DATASET_4',
      version='None',
      domain='table',
      status='ACTIVE',
      created_on='2024-08-01 22:44:28'
    )]
    Downstream Lineage of feature view 'WINE_FEATURES'
    [LineageNode(
      name='LINEAGE_DEMO_DB.FEATURE_STORE.MY_TABLE_DATASET_4',
      version='None',
      domain='table',
      status='ACTIVE',
      created_on='2024-08-01 22:44:28'
    )]
```

3.2 Training Data from source tables

We will explore the workflow of creating training dataset directly from source tables instead of feature views.

3.2.1 Dataset from source table as training data Create the dataset from a source table. Lineage works in a similar way even when its trained with source view or a stage.

```
name="my_dataset_from_table",
    version="v1",
    input_dataframe=session.table(source_tables[0]),
)
```

Query Lineage Query the upstream lineage of the dataset we just created.

4. Model Lineage

Now let's train a simple random forest model, and evaluate the prediction accuracy.

Let's create a registry to save the trained models. All models need to be logged into the registry for their lineage to be tracked.

```
[]: from snowflake.ml.registry import Registry

registry = Registry(
    session=session,
    database_name=session.get_current_database(),
    schema_name=MODEL_DEMO_SCHEMA,
)
```

4.1 Model trained in snowflake ecosystem

Lets define a training function that uses Random forest to build the model

```
max_depth=3, n_estimators=20, random_state=42
)
rf.fit(train)
predictions = rf.predict(test)
output_label_names = ['OUTPUT_' + col for col in label_cols]
mse = snowml_metrics.mean_squared_error(
    df=predictions,
    y_true_col_names=label_cols,
    y_pred_col_names=output_label_names
)
accuracy = 100 - snowml_metrics.mean_absolute_percentage_error(
    df=predictions,
    y_true_col_names=label_cols,
    y_pred_col_names=output_label_names
)
print(f"MSE: {mse}, Accuracy: {accuracy}")
return rf
```

4.1.1 Model trained using Dataset Convert dataset to a snowpark dataframe and examine all the features in it.

```
[]: training_data_df = my_dataset.read.to_snowpark_dataframe()
assert training_data_df.count() == sample_count
# drop rows that have any nulls in value.
training_data_df = training_data_df.dropna(how='any')
training_data_df.to_pandas()
```

```
[]:
          QUALITY SULPHATES ALCOHOL FIXED_ACIDITY CITRIC_ACID CHLORIDES \
                5
                                   9.4
                        0.56
                                                   7.4
                                                               0.00
                                                                          0.076
     0
                5
                        0.68
                                   9.8
                                                   7.8
                                                               0.00
     1
                                                                          0.098
                5
     2
                        0.64
                                   9.5
                                                   7.6
                                                               0.29
                                                                          0.075
     3
                5
                        0.70
                                  11.1
                                                   7.9
                                                               0.40
                                                                          0.062
                7
     4
                        0.76
                                  10.7
                                                  11.8
                                                               0.49
                                                                          0.093
                                                               0.00
                                                                          0.081
     507
                4
                         0.46
                                   9.6
                                                   8.1
     508
                5
                        0.64
                                   9.7
                                                   6.7
                                                               0.08
                                                                          0.064
     509
                7
                        0.68
                                  11.4
                                                  13.3
                                                               0.75
                                                                          0.084
     510
                6
                         0.48
                                  10.5
                                                   5.6
                                                               0.78
                                                                          0.074
     511
                        0.63
                                  10.2
                                                  10.0
                                                               0.31
                                                                          0.090
          TOTAL_SULFUR_DIOXIDE
                                   PH MY_NEW_FEATURE
     0
                                                 0.000
                             34 3.51
     1
                             67 3.20
                                                 0.000
```

2	66	3.40	2.204
3	20	3.28	3.160
4	80	3.30	5.782
	•••	•••	•••
507	24	3.38	0.000
508	34	3.33	0.536
509	43	3.04	9.975
510	92	3.39	4.368
511	62	3.18	3.100

[512 rows x 9 columns]

Train the random forest model using Snowpark-ML and the dataset, then log the model in the registry.

```
[]: random_forest_model = train_model_using_snowpark_ml(training_data_df)

model_version = registry.log_model(
    model_name="MODEL_TRAINED_ON_DATASET",
    version_name="v1",
    model=random_forest_model,
    comment="Model trained with feature views, dataset",
)
```

feature cols: ['MY_NEW_FEATURE', 'PH', 'TOTAL_SULFUR_DIOXIDE', 'CITRIC_ACID',
'CHLORIDES', 'SULPHATES', 'FIXED_ACIDITY', 'ALCOHOL']

The version of package 'snowflake-snowpark-python' in the local environment is 1.20.2, which does not fit the criteria for the requirement 'snowflake-snowpark-python'. Your UDF might not work when the package version is different between the server and your local environment.

The version of package 'numpy' in the local environment is 1.24.4, which does not fit the criteria for the requirement 'numpy==1.24.3'. Your UDF might not work when the package version is different between the server and your local environment.

The version of package 'scikit-learn' in the local environment is 1.3.2, which does not fit the criteria for the requirement 'scikit-learn==1.3.0'. Your UDF might not work when the package version is different between the server and your local environment.

/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/site-

packages/sklearn/base.py:348: InconsistentVersionWarning: Trying to unpickle estimator DecisionTreeRegressor from version 1.3.0 when using version 1.3.2. This might lead to breaking code or invalid results. Use at your own risk. For more info please refer to:

https://scikit-learn.org/stable/model_persistence.html#security-maintainability-limitations

```
warnings.warn(
```

/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/site-packages/sklearn/base.py:348: InconsistentVersionWarning: Trying to unpickle

estimator RandomForestRegressor from version 1.3.0 when using version 1.3.2. This might lead to breaking code or invalid results. Use at your own risk. For more info please refer to:

https://scikit-learn.org/stable/model_persistence.html#security-maintainability-limitations

```
warnings.warn(
```

The version of package 'numpy' in the local environment is 1.24.4, which does not fit the criteria for the requirement 'numpy==1.24.3'. Your UDF might not work when the package version is different between the server and your local environment.

The version of package 'scikit-learn' in the local environment is 1.3.2, which does not fit the criteria for the requirement 'scikit-learn==1.3.0'. Your UDF might not work when the package version is different between the server and your local environment.

```
MSE: 0.25267477340674793, Accuracy: 99.92349333548655
```

/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/contextlib.py:113:
UserWarning: `relax_version` is not set and therefore defaulted to True.
Dependency version constraints relaxed from ==x.y.z to >=x.y, <(x+1). To use
specific dependency versions for compatibility, reproducibility, etc., set
`options={'relax_version': False}` when logging the model.
 return next(self.gen)</pre>

Query Lineage Query the upstream of the model we just trained.

```
[]: ds = model_version.lineage(direction="upstream")
ds
```

The model can also be explored as part of the downstream path of the dataset used to train the model.

```
[]: ds[0].lineage(direction="downstream")
```

4.1.2 Model trained using source tables Train the random forest model using Snowpark-ML and the source tables, then log the model in the registry.

Lineage works in a similar way even when its trained with source view or a stage.

feature cols: ['VOLATILE_ACIDITY', 'RESIDUAL_SUGAR', 'TOTAL_SULFUR_DIOXIDE',
'CITRIC_ACID', 'PH', 'CHLORIDES', 'SULPHATES', 'DENSITY', 'FIXED_ACIDITY',
'ALCOHOL', 'FREE_SULFUR DIOXIDE']

/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/site-

packages/sklearn/base.py:348: InconsistentVersionWarning: Trying to unpickle estimator DecisionTreeRegressor from version 1.3.0 when using version 1.3.2. This might lead to breaking code or invalid results. Use at your own risk. For more info please refer to:

https://scikit-learn.org/stable/model_persistence.html#security-maintainability-limitations

```
warnings.warn(
```

/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/site-

packages/sklearn/base.py:348: InconsistentVersionWarning: Trying to unpickle estimator RandomForestRegressor from version 1.3.0 when using version 1.3.2. This might lead to breaking code or invalid results. Use at your own risk. For more info please refer to:

https://scikit-learn.org/stable/model_persistence.html#security-maintainability-limitations

```
warnings.warn(
```

WARNING:snowflake.snowpark.session:The version of package 'numpy' in the local environment is 1.24.4, which does not fit the criteria for the requirement 'numpy==1.24.3'. Your UDF might not work when the package version is different between the server and your local environment.

WARNING:snowflake.snowpark.session:The version of package 'scikit-learn' in the local environment is 1.3.2, which does not fit the criteria for the requirement 'scikit-learn==1.3.0'. Your UDF might not work when the package version is different between the server and your local environment.

MSE: 0.39049050644031114, Accuracy: 99.90829288453038

/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/contextlib.py:113:
UserWarning: `relax_version` is not set and therefore defaulted to True.
Dependency version constraints relaxed from ==x.y.z to >=x.y, <(x+1). To use specific dependency versions for compatibility, reproducibility, etc., set `options={'relax_version': False}` when logging the model.

return next(self.gen)

Query Lineage Query the upstream of the model we just trained.

```
[]: table = model_version.lineage(direction="upstream")
table
```

The model can also be explored as part of the downstream path of the table used to train the model.

4.2 Model trained in non-snowflake ecosystem

)]

For the workflows such as: A model trained using snowpark.ml but not a Snowpark DataFrame (like pandas). A model trained without using snowpark.ml or a Snowpark DataFrame. A model trained outside of Snowflake.

You can still associate the lineage between the source data object and the trained model by passing the snowpark dataframe backed by the source data object to model registry's log_model API as sample_input_data.

```
rf = RandomForestRegressor(max_depth=3, n_estimators=20, random_state=42)
    rf.fit(X_train, y_train)
    predictions = rf.predict(X_test)
    mse = mean_squared_error(y_test, predictions)
    accuracy = 100 - mean_absolute_percentage_error(y_test, predictions)
    print(f"MSE: {mse}, Accuracy: {accuracy}")
    return rf
feature_columns = list(set(training_data_df.columns) - set(label_cols) -_u
  →set(join_keys) - set([timestamp_col]))
print(f"feature cols: {feature_columns}")
sklearn_trained_model = train_model_using_sklearn(training_data_df.to_pandas(),_
 →feature_columns)
training_data_df = training_data_df.select(feature_columns)
model_version = registry.log_model(
    model_name="MODEL_TRAINED_ON_PANDAS",
    version_name="v1",
    model=sklearn_trained_model,
    comment="Model trained with pandas dataframe",
    # Passing the snowpark dataframe as sample input data
    sample input data = training data df
)
feature cols: ['MY_NEW_FEATURE', 'PH', 'TOTAL_SULFUR_DIOXIDE', 'CITRIC_ACID',
'CHLORIDES', 'SULPHATES', 'FIXED_ACIDITY', 'ALCOHOL']
/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/site-
packages/sklearn/base.py:1152: DataConversionWarning: A column-vector y was
passed when a 1d array was expected. Please change the shape of y to
(n_samples,), for example using ravel().
 return fit_method(estimator, *args, **kwargs)
/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/contextlib.py:113:
UserWarning: `relax_version` is not set and therefore defaulted to True.
Dependency version constraints relaxed from ==x.y.z to >=x.y, <(x+1). To use
specific dependency versions for compatibility, reproducibility, etc., set
`options={'relax_version': False}` when logging the model.
 return next(self.gen)
MSE: 0.4896289668292035, Accuracy: 99.89502779572355
/opt/homebrew/anaconda3/envs/py38_env/lib/python3.8/site-
packages/snowflake/ml/model/model_signature.py:69: UserWarning: The sample input
has 512 rows, thus a truncation happened before inferring signature. This might
```

```
cause inaccurate signature inference. If that happens, consider specifying
signature manually.
  warnings.warn(
```

Query lineage Query the upstream of the model we just trained.

```
[]: ds = model_version.lineage(direction="upstream")
ds
```

The model can also be explored as part of the downstream path of the dataset used to train the model.

```
[]: print(ds[0].lineage(direction="downstream"))
```

```
[ModelVersion(
  name='MODEL_TRAINED_ON_DATASET',
  version='V1',
), ModelVersion(
  name='MODEL_TRAINED_ON_PANDAS',
  version='V1',
)]
```

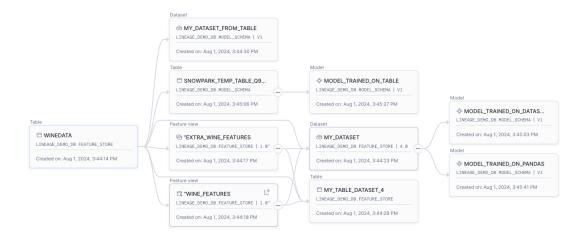
0.1 5. Visualization of lineage

The below image shows the screenshot of complete visualization of lineages of all the objects we created in the notebook from Snowsight UI.

```
[]: from IPython.display import Image, display

# Path to your image file
image_path = 'lineage-graph.png'

# Display the image
display(Image(filename=image_path,width=1000, height=1000))
```



6. Clean up notebook

This cell will drop the schemas have been created at beginning of this notebook, and also drop all objects live in the schemas including source data tables, feature views, datasets, and models.

```
[]: session.sql(f"DROP SCHEMA IF EXISTS {FS_DEMO_SCHEMA}").collect() session.sql(f"DROP SCHEMA IF EXISTS {MODEL_DEMO_SCHEMA}").collect()
```