

# Hey, I'm Stefanie

- Open Source Developer at probabl since 2024
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#### Connecting

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# Scikit-learn's new Metadata Routing API

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#### What is metadata routing?

Introducing scikit-learn's metadata routing API

#### Metadata

- Data to apply on top of tabular data
- Data a function can handle besides X and y

#### Use

- Fairness related, business logic, custom behaviour, etc.
- Benefit won't be a better score, but a more realistic model

#### **Routing**

• Passing metadata between several components of a data science pipeline

## What is metadata routing?

Introducing scikit-learn's metadata routing API

#### **Prior to metadata routing API**

- Restricted use of sample\_weight
- No metadata passing in nested structures

#### With metadata routing API

- Nested sample\_weight and groups passing
- Using metadata from other libraries' objects with scikit-learn estimators
- Custom definition of new metadata

In depth into the "groups" metadata in scikit-learn

#### Data

	Sex	Age	Severity	Medication
a <b>s</b>	Male	88	6	1
1 {	Female	100	4	0
3 <b>5</b>	Female	55	10	1
1 {	Female	70	7	1
3 C	Male	65	5	0
	Male	68	0	1

#### **Target**

Recovery Time
33
9
10
25
70
29

In depth into the "groups" metadata in scikit-learn

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Sex	Age	Severity	Medication
Male	88	6	1
Female	100	4	0
Female	55	10	1
Female	70	7	1
Male	65	5	0
Male	68	0	1

#### groups

#### Target

Hospital	Recovery Time
Blue	33
Blue	9
Black	10
Black	25
Orange	70
Orange	29

Groups handling during cross-validation



In depth into the "groups" metadata in scikit-learn

```
from sklearn.model selection import GroupKFold, cross validate
from sklearn.linear model import Ridge
ridge = Ridge()
# GroupKFold considers groups while splitting:
cv = GroupKFold(n splits=2)
# providing `groups` for GroupKFold splitter:
cross validate(ridge, X, y, cv=cv, groups=groups)
{'fit_time': array([0.00135875, 0.00036836]),
 'score time': array([0.00036907, 0.00023532]),
 'test score': array([-0.2923336 , -0.12131859]}
```

Using the "sample\_weight" metadata in scikit-learn

sample\_weight draws a model's attention to a sub-group of samples, in order to adjust their distribution (e.g. IPTW in medical settings)

• in distinction to a randomised trial, real-world observations often come with imbalanced and biased data

Sex	Age	Severity	Medication
Male	88	6	1
Female	100	4	0
Female	55	10	1

Recovery Time		
33		
9		
10		

Using the "sample\_weight" metadata in scikit-learn

```
from sklearn.model_selection import cross_validate
from sklearn.linear_model import Ridge

# Ridge().fit() can consume `sample_weight`
ridge = Ridge()

cross_validate(ridge, X, y, cv=cv, sample_weight=sample_weight)

TypeError: got an unexpected keyword argument 'sample weight'
```

```
scoring = get scorer("neg mean squared error") # can consume sample weight
search = GridSearchCV(
  ridge, param grid={"alpha": [0.1, 1, 10]}, cv=cv, scoring=scoring
cross validate(
   search, X, y, cv=cv, scoring=scoring,
   sample weight=sample_weight, groups=groups,
```

ridge = Ridge() # can consume sample weight

cv=GroupKFold(n splits=2) # consumes groups

```
    Cross-validation on a

        GridSearchCV that

        searches Ridge

        hyperparameters
```

Passing sample\_weight and groups into it

```
TypeError: got an unexpected
keyword argument 'sample_weight'
```

# Using the metadata routing API

```
ridge = Ridge()
cv=GroupKFold(n splits=2)
scoring = get_scorer("neg_mean_squared_error")
search = GridSearchCV(
 ridge, param grid={"alpha": [0.1, 1, 10]}, cv=cv, scoring=scoring
cross_validate(
   search, X, y, cv=cv, scoring=scoring,
   params={"sample weight": sample weight, "groups": groups},
```

# Using the metadata routing API

```
import sklearn
sklearn.set_config(enable_metadata_routing=True)
ridge = Ridge()
cv=GroupKFold(n splits=2)
scoring = get scorer("neg mean squared error")
search = GridSearchCV(
 ridge, param grid={"alpha": [0.1, 1, 10]}, cv=cv, scoring=scoring
cross validate(
   search, X, y, cv=cv, scoring=scoring,
   params={"sample weight": sample weight, "groups": groups},
```

1. Enable metadata routing

# Using the metadata routing API

```
import sklearn
```

```
sklearn.set_config(enable_metadata_routing=True)
ridge = Ridge()
```

```
cv=GroupKFold(n_splits=2)
scoring = get scorer("neg mean squared error")
```

search, X, y, cv=cv, scoring=scoring,

```
search = GridSearchCV(
    ridge, param_grid={"alpha": [0.1, 1, 10]}, cv=cv, scoring=scoring
)
cross validate(
```

params={"sample weight": sample weight, "groups": groups},

```
    Enable metadata routing
```

```
Pass metadata to top level tool
```

```
Using the metadata routing API
import sklearn
sklearn.set config(enable metadata routing=True)
ridge = Ridge().set_fit_request(sample_weight=True)
```

```
cv=GroupKFold(n splits=2)
scoring = get scorer("neg mean squared error")
scoring.set_score_request(sample_weight=True)
search = GridSearchCV(
  ridge, param grid={"alpha": [0.1, 1, 10]}, cv=cv, scoring=scoring
```

cross validate(

search, X, y, cv=cv, scoring=scoring,

```
Pass metadata to top level
tool
```

Enable metadata routing

```
Set requests where
metadata is consumed
```

```
params={"sample weight": sample weight, "groups": groups},
```

Voilà: Routing Success!

## Metadata routing API

Ongoing work

- Expect compatibility for all scikit-learn estimators in 1.7 release
- Defining default settings to relieve users from setting all the requests
- Keep flexibility for custom scenarios

#### Metadata routing API

#### References and Links

#### **Metadata routing**

- User Guide on Metadata Routing
- Developer Guide on Metadata Routing
- Adrin Jalali's talk on the internal logic of metadata routing

#### Use cases using `sample\_weight`

- :probabl. Whiteboard Series by Vincent Warmerdam: Improving models via subsets
- Blogpost by Florian Wilhelm on Inverse Probability of Treatment Weighting

#### Metadata routing API

Thank you for your attention!

# Time for your questions

#### Connecting

https://www.linkedin.com/in/StefanieSenger https://github.com/StefanieSenger