

custom transformers + patsy & statsmodels

Lecture 17

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Custom sklearn transformers

FunctionTransformer

The simplest way to create a new transformer is to use `FunctionTransformer()` from the preprocessing submodule which allows for converting a Python function into a transformer.

```
1 from sklearn.preprocessing import FunctionTransformer
2 X = pd.DataFrame({"x1": range(1,6), "x2": range(5, 0, -1)})
```

```

1 log_transform = FunctionTransformer(np.log)
2 lt = log_transform.fit(X)
3 lt

```

FunctionTransformer(func=<ufunc 'log'>)

```
1 lt.transform(X)
```

	x1	x2
0	0.0000	1.6094
1	0.6931	1.3863
2	1.0986	1.0986
3	1.3863	0.6931
4	1.6094	0.0000

```
1 lt.get_params()
```

```
{'accept_sparse': False, 'check_inverse': True,
'feature_names_out': None, 'func': <ufunc
'log'>, 'inv_kw_args': None, 'inverse_func':
None, 'kw_args': None, 'validate': False}
```

```
1 dir(lt)
```

```
['__annotations__', '__class__', '__delattr__',
'__dict__', '__dir__', '__doc__', '__eq__',
'__format__', '__ge__', '__getattr__',
'__getstate__', '__gt__', '__hash__',
'__init__', '__init_subclass__', '__le__',
'__lt__', '__module__', '__ne__', '__new__',
'__reduce__', '__reduce_ex__', '__repr__',
'__setattr__', '__setstate__', '__sizeof__',
'__sklearn_clone__', '__sklearn_is_fitted__',
'__sklearn_tags__', '__str__',
'__subclasshook__', '__weakref__',
'_build_request_for_signature',
'_check_feature_names', '_check_input',
'_check_inverse_transform',
'_check_n_features', '_doc_link_module',
'_doc_link_template',
'_doc_link_url_param_generator',
'_get_default_requests', '_get_doc_link']
```

Input types

```
1 def interact(X, y = None):  
2     return np.c_[X, X[:,0] * X[:,1]]  
3 X = pd.DataFrame({"x1": range(1,6), "x2": range(5, 0, -1)})  
4 Z = np.array(X)
```

```
1 FunctionTransformer(  
2     interact  
3 ).fit_transform(X)
```

pandas.errors.InvalidIndexError: (slice(None, None, None), 0)

```
1 FunctionTransformer(  
2     interact  
3 ).fit_transform(Z)
```

```
array([[1, 5, 5],  
       [2, 4, 8],  
       [3, 3, 9],  
       [4, 2, 8],  
       [5, 1, 5]])
```

```
1 FunctionTransformer(  
2     interact, validate=True  
3 ).fit_transform(X)
```

```
array([[1, 5, 5],  
       [2, 4, 8],  
       [3, 3, 9],  
       [4, 2, 8],  
       [5, 1, 5]])
```

```
1 FunctionTransformer(  
2     interact, validate=True  
3 ).fit_transform(Z)
```

```
array([[1, 5, 5],  
       [2, 4, 8],  
       [3, 3, 9],  
       [4, 2, 8],  
       [5, 1, 5]])
```

Build your own transformer

For a more full featured transformer, it is possible to construct it as a class that inherits from `BaseEstimator` and `TransformerMixin` classes from the `base` submodule.

```
1 from sklearn.base import BaseEstimator, TransformerMixin
2
3 class scaler(BaseEstimator, TransformerMixin):
4     def __init__(self, m = 1, b = 0):
5         self.m = m
6         self.b = b
7
8     def fit(self, X, y=None):
9         return self
10
11    def transform(self, X, y=None):
12        return X*self.m + self.b
```

```

1 X = pd.DataFrame({
2     "x1": range(1,6),
3     "x2": range(5, 0, -1)}
4 ); X

```

	x1	x2
0	1	5
1	2	4
2	3	3
3	4	2
4	5	1

```

1 double = scaler(2)
2 double.fit_transform(X)

```

	x1	x2
0	2	10
1	4	8
2	6	6
3	8	4
4	10	2

```

1 double.get_params()

```

```
{'b': 0, 'm': 2}
```

```

1 double.set_params(b=-3).fit_transform(X)

```

	x1	x2
0	-1	7
1	1	5
2	3	3
3	5	1
4	7	-1

What else do we get?

```
1 print(  
2     np.array(dir(double))  
3 )
```

```
['__class__', '__delattr__', '__dict__', '__dir__', '__doc__', '__eq__', '__format__',  
 '__ge__', '__getattr__', '__getstate__', '__gt__', '__hash__', '__init__',  
 '__init_subclass__', '__le__', '__lt__', '__module__', '__ne__', '__new__',  
 '__reduce__', '__reduce_ex__', '__repr__', '__setattr__', '__setstate__',  
 '__sizeof__', '__sklearn_clone__', '__sklearn_tags__', '__str__',  
 '__subclasshook__', '__weakref__', '_build_request_for_signature',  
 '_check_feature_names', '_check_n_features', '_doc_link_module',  
 '_doc_link_template', '_doc_link_url_param_generator', '_get_default_requests',  
 '_get_doc_link', '_get_metadata_request', '_get_param_names', '_get_tags',  
 '_more_tags', '_repr_html_', '_repr_html_inner', '_repr_mimebundle_',  
 '_sklearn_auto_wrap_output_keys', '_validate_data', '_validate_params', 'b', 'fit',  
 'fit_transform', 'get_metadata_routing', 'get_params', 'm', 'set_output',  
 'set_params', 'transform']
```


Demo - Interaction Transformer

Useful methods

We employed a couple of special methods that are worth mentioning in a little more detail.

- `_validate_data()` & `_check_feature_names()` are methods that are inherited from `BaseEstimator` they are responsible for setting and checking the `n_features_in_` and the `feature_names_in_` attributes respectively.
- In general one or both is run during `fit()` with `reset=True` in which case the respective attribute will be set.
- Later, in `transform()` one or both will again be called with `reset=False` and the properties of `X` will be checked against the values in the attribute.
- These are worth using as they promote an interface consistent with sklearn and also provide convenient error checking with useful warning / error messages.

These methods are part of `BaseEstimator` but are not available as part of the published documentation see the code on

`check_is_fitted()`

This is another useful helper function from `sklearn.utils` - it is fairly simplistic in that it checks for the existence of a specified attribute. If no attribute is given then it checks for any attributes ending in `_` that do not begin with `__`.

Again this is useful for providing a consistent interface and useful error / warning messages.

See also the other `check*()` functions in `sklearn.utils`.

Other custom estimators

If you want to implement your own custom modeling function it is possible, there are different Mixin base classes in `sklearn.base` that provide the common core interface.

Class	Description
<code>base.Biclustermixin</code>	Mixin class for all bicluster estimators
<code>base.ClassifierMixin</code>	Mixin class for all classifiers
<code>base.ClusterMixin</code>	Mixin class for all cluster estimators
<code>base.DensityMixin</code>	Mixin class for all density estimators
<code>base.RegressorMixin</code>	Mixin class for all regression estimators
<code>base.TransformerMixin</code>	Mixin class for all transformers
<code>base.OneToOneFeatureMixin</code>	Provides <code>get_feature_names_out</code> for simple transformers

patsy

paty

`paty` is a Python package for describing statistical models (especially linear models, or models that have a linear component) and building design matrices. It is closely inspired by and compatible with the formula mini-language used in R and S.

...

Patsy's goal is to become the standard high-level interface to describing statistical models in Python, regardless of what particular model or library is being used underneath.

Formulas

```
1 from patsy import ModelDesc
```

```
1 ModelDesc.from_formula("y ~ a + a:b + np.log(x)")
```

```
ModelDesc(lhs_termlist=[Term([EvalFactor('y')])],  
          rhs_termlist=[Term([]),  
                        Term([EvalFactor('a')]),  
                        Term([EvalFactor('a'), EvalFactor('b')]),  
                        Term([EvalFactor('np.log(x)')])])])
```

```
1 ModelDesc.from_formula("y ~ a*b + np.log(x) - 1")
```

```
ModelDesc(lhs_termlist=[Term([EvalFactor('y')])],  
          rhs_termlist=[Term([EvalFactor('a')]),  
                        Term([EvalFactor('b')]),  
                        Term([EvalFactor('a'), EvalFactor('b')]),  
                        Term([EvalFactor('np.log(x)')])])])
```

Model matrix

```
1 from patsy import demo_data, dmatrix, dmatrices
```

```
1 data = demo_data("y", "a", "b", "x1", "x2")
2 data
```

```
{'a': ['a1', 'a1', 'a2', 'a2', 'a1', 'a1',
      'a2', 'a2'], 'b': ['b1', 'b2', 'b1', 'b2',
      'b1', 'b2', 'b1', 'b2'], 'x1': array([ 1.7641,
      0.4002, 0.9787, 2.2409, 1.8676, -0.9773,
      0.9501, -0.1514]), 'x2': array([-0.1032,
      0.4106, 0.144 , 1.4543, 0.761 , 0.1217,
      0.4439, 0.3337]), 'y': array([ 1.4941,
      -0.2052, 0.3131, -0.8541, -2.553 , 0.6536,
      0.8644, -0.7422])}
```

```
1 pd.DataFrame(data)
```

	a	b	x1	x2	y
0	a1	b1	1.7641	-0.1032	1.4941
1	a1	b2	0.4002	0.4106	-0.2052
2	a2	b1	0.9787	0.1440	0.3131
3	a2	b2	2.2409	1.4543	-0.8541
4	a1	b1	1.8676	0.7610	-2.5530
5	a1	b2	-0.9773	0.1217	0.6536
6	a2	b1	0.9501	0.4439	0.8644
7	a2	b2	-0.1514	0.3337	-0.7422

```
1 dmatrix("a + a:b + np.exp(x1)", data)
```

```
DesignMatrix with shape (8, 5)
      Intercept  a[T.a2]  a[a1]:b[T.b2]
a[a2]:b[T.b2]  np.exp(x1)
              1          0              0
0      5.83604
              1          0              1
0      1.49206
              1          1              0
0      2.66110
              1          1              0
1      9.40173
              1          0              0
0      6.47247
              1          0              1
0      0.37633
              1          1              0
0      2.58594
              1          1              0
```


Model matrices

```
1 y, X = dmatrices("y ~ a + a:b + np.exp(x1)", data)
```

```
1 y
```

DesignMatrix with shape (8, 1)

y
1.49408
-0.20516
0.31307
-0.85410
-2.55299
0.65362
0.86444
-0.74217

Terms:

'y' (column 0)

```
1 X
```

DesignMatrix with shape (8, 5)

Intercept	a[T.a2]	a[a1]:b[T.b2]	a[a2]:b[T.b2]	np.exp(x1)
1	0	0	0	5.83604
1	0	1	0	1.49206
1	1	0	0	2.66110
1	1	0	1	9.40173
1	0	0	0	6.47247
1	0	1	0	0.37633
1	1	0	0	2.58594
1	1	0	1	0.85954

Terms:

'Intercept' (column 0)
'a' (column 1)
'a:b' (columns 2:4)
'np.exp(x1)' (column 4)

as DataFrames

```
1 dmatrix("a + a:b + np.exp(x1)", data, return_type='dataframe')
```

	Intercept	a[T.a2]	a[a1]:b[T.b2]	a[a2]:b[T.b2]	np.exp(x1)
0	1.0	0.0	0.0	0.0	5.8360
1	1.0	0.0	1.0	0.0	1.4921
2	1.0	1.0	0.0	0.0	2.6611
3	1.0	1.0	0.0	1.0	9.4017
4	1.0	0.0	0.0	0.0	6.4725
5	1.0	0.0	1.0	0.0	0.3763
6	1.0	1.0	0.0	0.0	2.5859
7	1.0	1.0	0.0	1.0	0.8595

Formula Syntax

Code	Description	Example
<code>+</code>	unions terms on the left and right	<code>a+a ⇒ a</code>
<code>-</code>	removes terms on the right from terms on the left	<code>a+b-a ⇒ b</code>
<code>:</code>	constructs interactions between each term on the left and right	<code>(a+b):c ⇒ a:c + b:c</code>
<code>*</code>	short-hand for terms and their interactions	<code>a*b ⇒ a + b + a:b</code>
<code>/</code>	short-hand for left terms and their interactions with right terms	<code>a/b ⇒ a + a:b</code>
<code>I()</code>	used for calculating arithmetic calculations	<code>I(x1 + x2)</code>
<code>Q()</code>	used to quote column names, e.g. columns with spaces or symbols	<code>Q('bad name!')</code>
<code>C()</code>	used for categorical data coding	<code>C(a, Treatment('a2'))</code>

Examples

```
1 dmatrix("x:y", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 2)

	Intercept	x:y
1	-1.72397	
1	0.38018	
1	-0.14814	
1	-0.23130	
1	0.76682	

Terms:

'Intercept' (column 0)
'x:y' (column 1)

```
1 dmatrix("x*y", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 4)

	Intercept	x	y	x:y
1	1.76405	-0.97728	-1.72397	
1	0.40016	0.95009	0.38018	
1	0.97874	-0.15136	-0.14814	
1	2.24089	-0.10322	-0.23130	
1	1.86756	0.41060	0.76682	

Terms:

'Intercept' (column 0)
'x' (column 1)
'y' (column 2)
'x:y' (column 3)

```
1 dmatrix("x/y", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 3)

	Intercept	x	x:y
1	1.76405	-1.72397	
1	0.40016	0.38018	
1	0.97874	-0.14814	
1	2.24089	-0.23130	
1	1.86756	0.76682	

Terms:

'Intercept' (column 0)
'x' (column 1)
'x:y' (column 2)

```
1 dmatrix("x*(y+z)", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 6)

Intercept	x	y	z	x:y	x:z
1	1.76405	-0.97728	0.14404	-1.72397	0.25410
1	0.40016	0.95009	1.45427	0.38018	0.58194
1	0.97874	-0.15136	0.76104	-0.14814	0.74486
1	2.24089	-0.10322	0.12168	-0.23130	0.27266
1	1.86756	0.41060	0.44386	0.76682	0.82894

Terms:

- 'Intercept' (column 0)
- 'x' (column 1)
- 'y' (column 2)
- 'z' (column 3)
- 'x:y' (column 4)
- 'x:z' (column 5)

Intercept Examples (-1)

```
1 dmatrix("x", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 2)

	Intercept	x
1	1.76405	
1	0.40016	
1	0.97874	
1	2.24089	
1	1.86756	

Terms:

'Intercept' (column 0)
'x' (column 1)

```
1 dmatrix("x-1", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 1)

	x
1	1.76405
1	0.40016
1	0.97874
1	2.24089
1	1.86756

Terms:

'x' (column 0)

```
1 dmatrix("-1 + x", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 1)

	x
1	1.76405
1	0.40016
1	0.97874
1	2.24089
1	1.86756

Terms:

'x' (column 0)

Intercept Examples (0)

```
1 dmatrix("x+0", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 1)

```
      x
1.76405
0.40016
0.97874
2.24089
1.86756
Terms:
  'x' (column 0)
```

```
1 dmatrix("x-0", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 2)

```
Intercept      x
1  1.76405
1  0.40016
1  0.97874
1  2.24089
1  1.86756
Terms:
'Intercept' (column 0)
'x' (column 1)
```

```
1 dmatrix("x - (-0)", demo_data("x","y","z"))
```

DesignMatrix with shape (5, 1)

```
      x
1.76405
0.40016
0.97874
2.24089
1.86756
Terms:
  'x' (column 0)
```

Design Info

One of the keep features of the design matrix object is that it retains all the necessary details (including stateful transforms) that are necessary to apply to new data inputs (e.g. for prediction).

```
1 d = dmatrix("a + a:b + np.exp(x1)", data, return_type='dataframe')
2 d.design_info
```

```
DesignInfo(['Intercept',
            'a[T.a2]',
            'a[a1]:b[T.b2]',
            'a[a2]:b[T.b2]',
            'np.exp(x1)'],
            factor_infos={EvalFactor('a'): FactorInfo(factor=EvalFactor('a'),
                                                         type='categorical',
                                                         state=<factor state>,
                                                         categories=('a1', 'a2')),
                          EvalFactor('b'): FactorInfo(factor=EvalFactor('b'),
                                                         type='categorical',
                                                         state=<factor state>,
                                                         categories=('b1', 'b2')),
                          EvalFactor('np.exp(x1)'): FactorInfo(factor=EvalFactor('np.exp(x1)'),
                                                                type='numerical',
                                                                state=<factor state>,
                                                                num_columns=1)},
            term_codings=OrderedDict([(Term([],
```


Stateful transforms

```
1 data = {"x1": np.random.normal(size=10)}  
2 new_data = {"x1": np.random.normal(size=10)}
```

```
1 d = dmatrix("scale(x1)", data)  
2 d
```

DesignMatrix with shape (10, 2)

Intercept	scale(x1)
1	0.57782
1	1.39708
1	-1.27276
1	0.75630
1	-1.73517
1	-0.59511
1	0.64699
1	0.88133
1	-0.93921
1	0.28274

Terms:

'Intercept' (column 0)
'scale(x1)' (column 1)

```
1 np.mean(d, axis=0)
```

array([1., -0.])

```
1 pred = dmatrix(d.design_info, new_data)  
2 pred
```

DesignMatrix with shape (10, 2)

Intercept	scale(x1)
1	-1.57035
1	1.23389
1	-0.79933
1	-1.65573
1	-2.30259
1	2.68184
1	-1.25777
1	-3.52965
1	-2.89942
1	-0.88729

Terms:

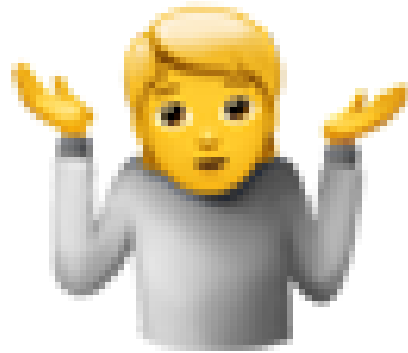
'Intercept' (column 0)
'scale(x1)' (column 1)

```
1 np.mean(pred, axis=0)
```

array([1. , -1.0986])

scikit-learn + Patsy

The state of affairs here is a bit of a mess at the moment - previously the `sklego` package implemented a `PatsyTransformer` class that has since been deprecated in favor of the `FormulaicTransformer` which uses the `formulaic` package for formula handling.



A PatsyTransformer

```
1 from patsy import dmatrix, build_design_matrices
2 from sklearn.utils.validation import check_is_fitted
3 from sklearn.base import BaseEstimator, TransformerMixin
4
5 class PatsyTransformer(TransformerMixin, BaseEstimator):
6     def __init__(self, formula):
7         self.formula = formula
8
9     def fit(self, X, y=None):
10        m = dmatrix(self.formula, X)
11        assert np.array(m).shape[0] == np.array(X).shape[0]
12        self.design_info_ = m.design_info
13        return self
14
15    def transform(self, X):
16        check_is_fitted(self, 'design_info_')
17        return build_design_matrices([self.design_info_], X)[0]
```

```

1 df = pd.DataFrame({
2     "y": [2, 2, 4, 4, 6], "x": [1, 2, 3, 4, 5],
3     "a": ["yes", "yes", "no", "no", "yes"]
4 })
5 X, y = df[["x", "a"]], df[["y"]].values

```

```

1 pt = PatsyTransformer("x*a + np.log(x)")
2 pt.fit_transform(X)

```

DesignMatrix with shape (5, 5)

Intercept	a[T.yes]	x	x:a[T.yes]	np.log(x)
1	1	1	1	0.00000
1	1	2	2	0.69315
1	0	3	0	1.09861
1	0	4	0	1.38629
1	1	5	5	1.60944

Terms:

'Intercept' (column 0)
 'a' (column 1)
 'x' (column 2)
 'x:a' (column 3)
 'np.log(x)' (column 4)

```

1 make_pipeline(
2     PatsyTransformer("x*a + np.log(x)"),
3     StandardScaler()
4 ).fit_transform(X)

```

```

array([[ 0.      ,  0.8165, -1.4142, -0.3235,
        -1.6845],
       [ 0.      ,  0.8165, -0.7071,  0.2157,
        -0.4651],
       [ 0.      , -1.2247,  0.      , -0.8627,
         0.2483],
       [ 0.      , -1.2247,  0.7071, -0.8627,
         0.7544],
       [ 0.      ,  0.8165,  1.4142,  1.8332,
         1.1469]])

```

statsmodels

statsmodels

statsmodels is a Python module that provides classes and functions for the estimation of many different statistical models, as well as for conducting statistical tests, and statistical data exploration. An extensive list of result statistics are available for each estimator. The results are tested against existing statistical packages to ensure that they are correct.

```
1 import statsmodels.api as sm
2 import statsmodels.formula.api as smf
3 import statsmodels.tsa.api as tsa
```

statsmodels uses slightly different terminology for referring to **y** (dependent / response) and **x** (independent / explanatory) variables. Specifically it uses **endog** to refer to the **y** and **exog** to refer to the **x** variable(s).

This is particularly important when using the main API, less so when using the formula API.

OpenIntro Loans data

This data set represents thousands of loans made through the Lending Club platform, which is a platform that allows individuals to lend to other individuals. Of course, not all loans are created equal. Someone who is a essentially a sure bet to pay back a loan will have an easier time getting a loan with a low interest rate than someone who appears to be riskier. And for people who are very risky? They may not even get a loan offer, or they may not have accepted the loan offer due to a high interest rate. It is important to keep that last part in mind, since this data set only represents loans actually made, i.e. do not mistake this data for loan applications!

For the full data dictionary see [here](#). We have removed some of the columns to make the data set more reasonably sized and also dropped any rows with missing values.

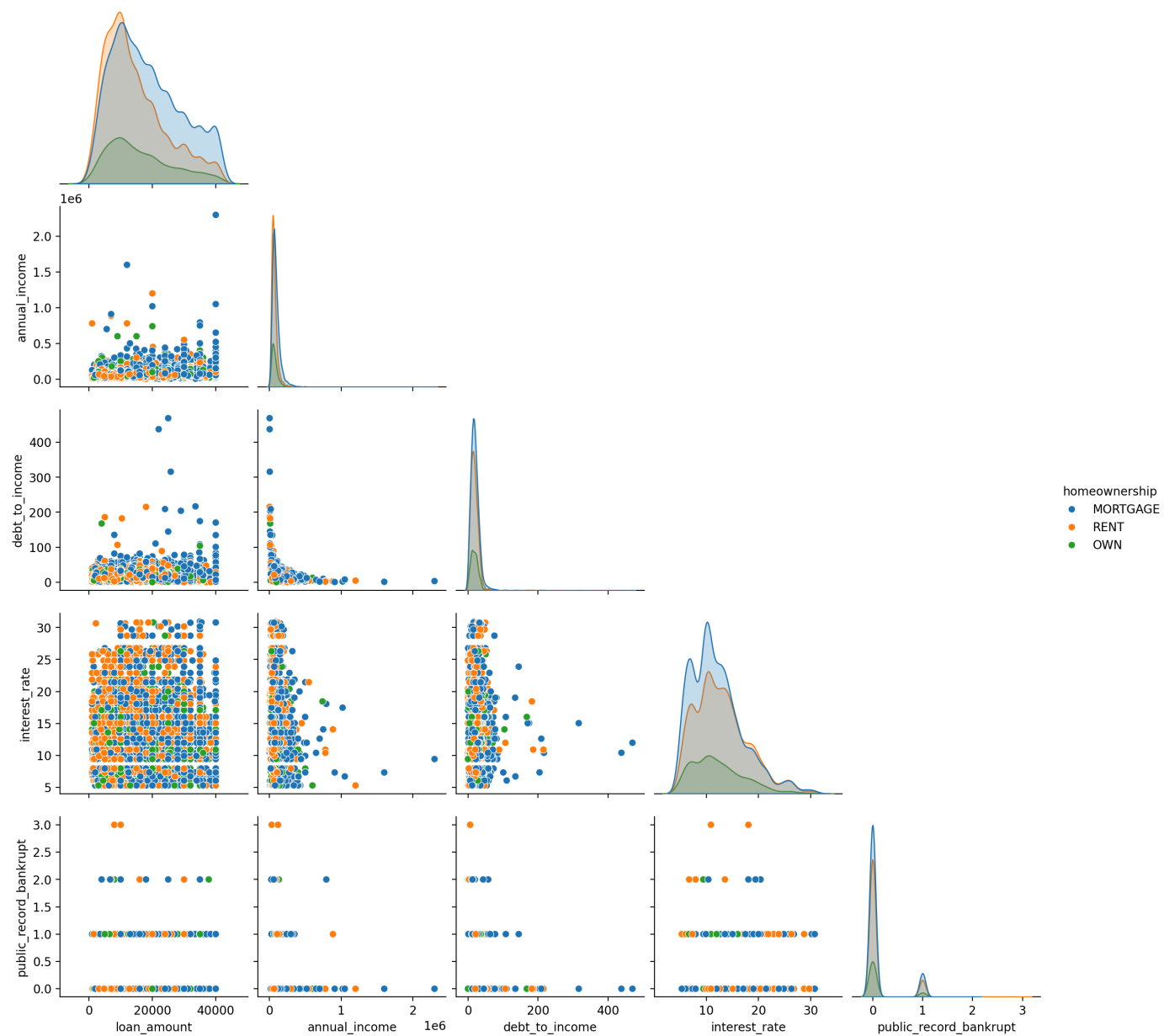
```
1 loans = pd.read_csv("data/openintro_loans.csv")
2 loans
```

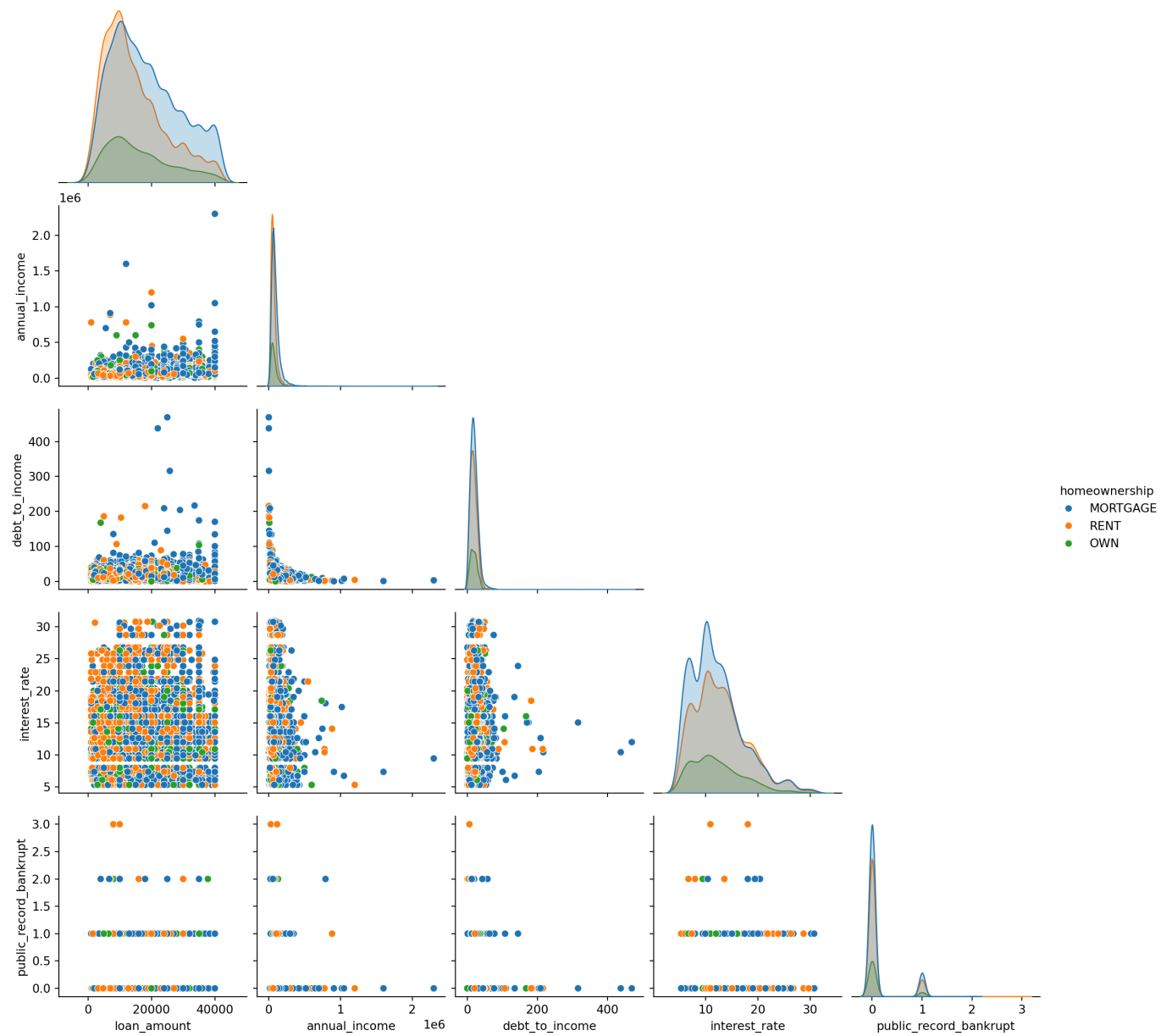
	state	emp_length	term	homeownership	annual_income	...	loan_amount	grade	interest_rate
\									
0	NJ	3	60	MORTGAGE	90000.0	...	28000	C	14.07
1	HI	10	36	RENT	40000.0	...	5000	C	12.61
2	WI	3	36	RENT	40000.0	...	2000	D	17.09
3	PA	1	36	RENT	30000.0	...	21600	A	6.72
4	CA	10	36	RENT	35000.0	...	23000	C	14.07
...
9177	TX	10	36	RENT	108000.0	...	24000	A	7.35
9178	PA	8	36	MORTGAGE	121000.0	...	10000	D	19.03
9179	CT	10	36	MORTGAGE	67000.0	...	30000	E	23.88
9180	WI	1	36	MORTGAGE	80000.0	...	24000	A	5.32
9181	CT	3	36	RENT	66000.0	...	12800	B	10.91

```
public_record_bankrupt  loan_status
0                        0      Current
1                        1      Current
```

```
1 print(loans.columns)
```

```
Index(['state', 'emp_length', 'term', 'homeownership', 'annual_income', 'verified_income',  
      'debt_to_income', 'total_credit_limit', 'total_credit_utilized',  
      'num_cc_carrying_balance',  
      'loan_purpose', 'loan_amount', 'grade', 'interest_rate', 'public_record_bankrupt',  
      'loan_status'],  
      dtype='object')
```



OLS

```
1 y = loans["loan_amount"]
2 X = loans[["homeownership", "annual_income", "debt_to_income", "interest_rate", "p
3
4 model = sm.OLS(endog=y, exog=X)
```

ValueError: Pandas data cast to numpy dtype of object. Check input data with `np.asarray(data)`.

What do you think the issue is here?

The error occurs because `X` contains mixed types - specifically we have categorical data columns which cannot be directly converted to a numeric dtype so we need to take care of the dummy coding for statsmodels (with this interface).

```
1 X_dc = pd.get_dummies(X, dtype='int')
2 model = sm.OLS(endog=y, exog=X_dc)
3 model
```

```
<statsmodels.regression.linear_model.OLS object at 0x1669c8a40>
```

```
1 np.array(dir(model))
```

```
array(['__class__', '__delattr__', '__dict__', '__dir__', '__doc__', '__eq__',
      '__format__', '__ge__', '__getattr__', '__getstate__', '__gt__',
      '__hash__', '__init__', '__init_subclass__', '__le__', '__lt__',
      '__module__', '__ne__', '__new__', '__reduce__', '__reduce_ex__',
      '__repr__', '__setattr__', '__sizeof__', '__str__', '__subclasshook__',
      '__weakref__', '_check_kwargs', '_data_attr', '_df_model', '_df_resid',
      '_fit_collinear', '_fit_ridge', '_fit_zeros', '_formula_max_endog',
      '_get_init_kwds', '_handle_data', '_init_keys', '_kwargs_allowed',
      '_setup_score_hess', '_sqrt_lasso', 'data', 'df_model', 'df_resid',
      'endog', 'endog_names', 'exog', 'exog_names', 'fit', 'fit_regularized',
      'from_formula', 'get_distribution', 'hessian', 'hessian_factor',
      'information', 'initialize', 'k_constant', 'loglike', 'nobs',
      'pinv_wexog', 'predict', 'rank', 'score', 'weights', 'wendog', 'wexog',
      'whiten'], dtype='<U18')
```

Fitting and summary

```
1 res = model.fit()  
2 print(res.summary())
```

OLS Regression Results					
=====					
Dep. Variable:	loan_amount	R-squared:	0.135		
Model:	OLS	Adj. R-squared:	0.135		
Method:	Least Squares	F-statistic:	239.5		
Date:	Thu, 20 Feb 2025	Prob (F-statistic):	2.33e-285		
Time:	11:39:13	Log-Likelihood:	-97245.		
No. Observations:	9182	AIC:	1.945e+05		
Df Residuals:	9175	BIC:	1.946e+05		
Df Model:	6				
Covariance Type:	nonrobust				
=====					
	coef	std err	t	P> t	[0.025
0.975]					
=====					

Formula interface

Most of the modeling interfaces are also provided by `smf` (`statsmodels.formula.api`), in which case `patsy` is used to construct the model matrices.

```
1 model = smf.ols(  
2     "loan_amount ~ homeownership + annual_income + debt_to_income + interest_rate + public_reco  
3     data = loans  
4 )  
5 res = model.fit()  
6 print(res.summary())
```

OLS Regression Results						
=====						
Dep. Variable:	loan_amount	R-squared:	0.135			
Model:	OLS	Adj. R-squared:	0.135			
Method:	Least Squares	F-statistic:	239.5			
Date:	Thu, 20 Feb 2025	Prob (F-statistic):	2.33e-285			
Time:	11:39:13	Log-Likelihood:	-97245.			
No. Observations:	9182	AIC:	1.945e+05			
Df Residuals:	9175	BIC:	1.946e+05			
Df Model:	6					
Covariance Type:	nonrobust					
=====						
	coef	std err	t	P> t	[0.025	0.975]

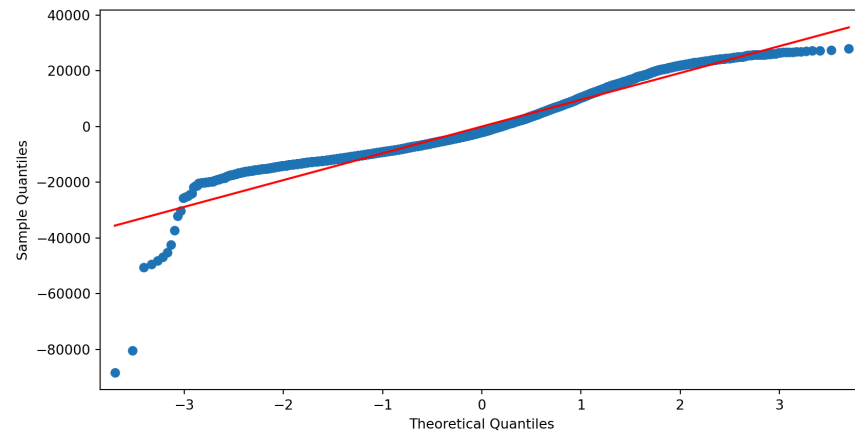
Intercept	1.002e+04	357.245	28.048	0.000	9319.724	1.07e+04
homeownership[T.OWN]	-1139.5893	322.361	-3.535	0.000	-1771.489	-507.690

Result values and model parameters

Diagnostic plots

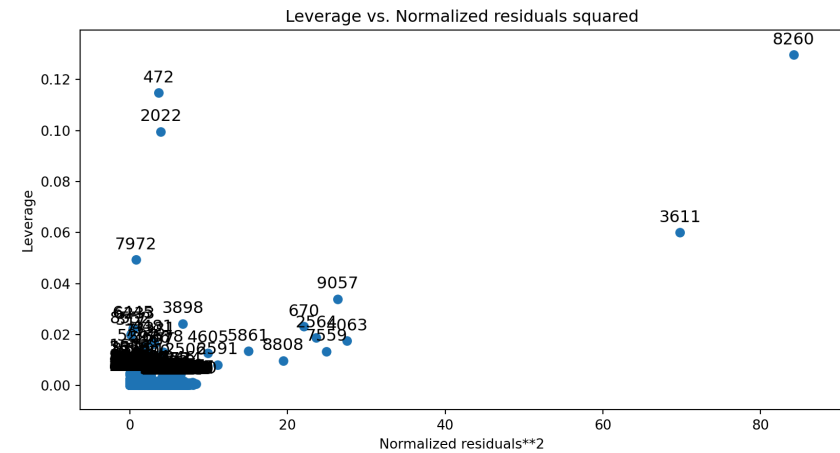
QQ Plot

```
1 plt.figure()
2 sm.graphics.qqplot(res.resid, line="s")
3 plt.show()
```



Leverage plot

```
1 plt.figure()
2 sm.graphics.plot_leverage_resid2(res)
3 plt.show()
```



Alternative model

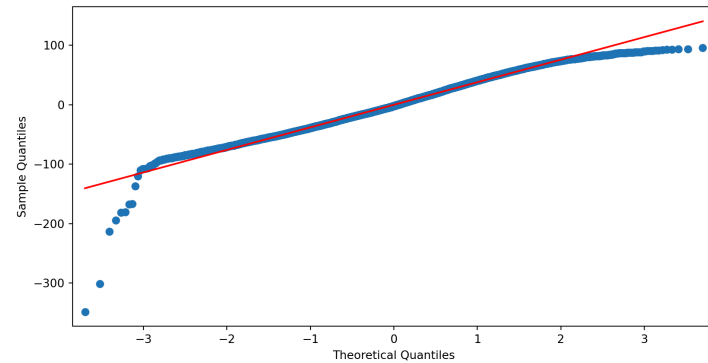
```
1 res = smf.ols(  
2     "np.sqrt(loan_amount) ~ homeownership + annual_income + debt_to_income + interest_rate + pu  
3     data = loans  
4 ).fit()  
5 print(res.summary())
```

OLS Regression Results

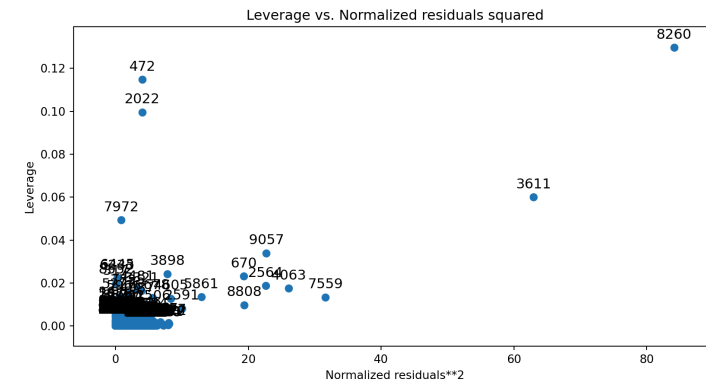
=====						
Dep. Variable:	np.sqrt(loan_amount)	R-squared:	0.132			
Model:	OLS	Adj. R-squared:	0.132			
Method:	Least Squares	F-statistic:	232.7			
Date:	Thu, 20 Feb 2025	Prob (F-statistic):	1.16e-277			
Time:	11:39:14	Log-Likelihood:	-46429.			
No. Observations:	9182	AIC:	9.287e+04			
Df Residuals:	9175	BIC:	9.292e+04			
Df Model:	6					
Covariance Type:	nonrobust					
=====						
	coef	std err	t	P> t	[0.025	0.975]

Intercept	95.4915	1.411	67.687	0.000	92.726	98.257
homeownership[T.OWN]	-4.4495	1.273	-3.495	0.000	-6.945	-1.954
homeownership[T.RENT]	-10.4225	0.873	-11.937	0.000	-12.134	-8.711
annual_income	0.0002	6.24e-06	30.016	0.000	0.000	0.000

```
1 plt.figure()
2 sm.graphics.qqplot(res.resid, line="s")
3 plt.show()
```



```
1 plt.figure()
2 sm.graphics.plot_leverage_resid2(res)
3 plt.show()
```



Bushtail Possums

Data representing possums in Australia and New Guinea. This is a copy of the data set by the same name in the DAAG package, however, the data set included here includes fewer variables.

```
1 possum = pd.read_csv("data/possum.csv")
2 possum
```

	site	pop	sex	age	head_l	skull_w	total_l	tail_l
0	1	Vic	m	8.0	94.1	60.4	89.0	36.0
1	1	Vic	f	6.0	92.5	57.6	91.5	36.5
2	1	Vic	f	6.0	94.0	60.0	95.5	39.0
3	1	Vic	f	6.0	93.2	57.1	92.0	38.0
4	1	Vic	f	2.0	91.5	56.3	85.5	36.0
..
99	7	other	m	1.0	89.5	56.0	81.5	36.5
100	7	other	m	1.0	88.6	54.7	82.5	39.0
101	7	other	f	6.0	92.4	55.0	89.0	38.0
102	7	other	m	4.0	91.5	55.2	82.5	36.5
103	7	other	f	3.0	93.6	59.9	89.0	40.0

[104 rows x 8 columns]



Logistic regression models (GLM)

```
1 y = pd.get_dummies( possum["pop"], drop_first = True, dtype="int")
2 X = pd.get_dummies( possum.drop(["site","pop"], axis=1), dtype="int")
3
4 model = sm.GLM(y, X, family = sm.families.Binomial())
```

`statsmodels.tools.sm_exceptions.MissingDataError: exog contains inf or nans`

What went wrong this time?

Missing values

Missing values can be handled via `missing` argument, possible values are `"none"`, `"drop"`, and `"raise"`.

```
1 model = sm.GLM(y, X, family = sm.families.Binomial(), missing="drop")
2 res = model.fit()
3 print(res.summary())
```

Success vs failure

Note `endog` can be 1d or 2d for binomial models - in the case of the latter each row is interpreted as [success, failure].

```
1 y = pd.get_dummies( possum["pop"], dtype="int")
2 X = pd.get_dummies( possum.drop(["site","pop"], axis=1), dtype="int")
3
4 res = sm.GLM(y, X, family = sm.families.Binomial(), missing="drop").fit()
5 print(res.summary())
```

Generalized Linear Model Regression Results

=====			
Dep. Variable:	['Vic', 'other']	No. Observations:	102
Model:	GLM	Df Residuals:	95
Model Family:	Binomial	Df Model:	6
Link Function:	Logit	Scale:	1.0000
Method:	IRLS	Log-Likelihood:	-31.942
Date:	Thu, 20 Feb 2025	Deviance:	63.885
Time:	11:39:15	Pearson chi2:	154.
No. Iterations:	7	Pseudo R-squ. (CS):	0.5234
Covariance Type:	nonrobust		
=====			

	coef	std err	z	P> z	[0.025	0.975]

age	0.1373	0.183	0.751	0.453	-0.221	0.495
head 1	0.1072	0.150	1.247	0.212	0.507	0.112

Formula interface

```
1 res = smf.glm(  
2   "pop ~ sex + age + head_l + skull_w + total_l + tail_l-1",  
3   data = possum,  
4   family = sm.families.Binomial(),  
5   missing="drop"  
6 ).fit()  
7 print(res.summary())
```

Generalized Linear Model Regression Results

```
=====
Dep. Variable:      ['pop[Vic]', 'pop[other]']    No. Observations:
102
Model:              GLM      Df Residuals:
95
Model Family:      Binomial    Df Model:
6
Link Function:      Logit      Scale:
1.0000
Method:              IRLS      Log-Likelihood:
-31.942
Date:               Thu, 20 Feb 2025    Deviance:
63.885
Time:               11:39:15    Pearson chi2:
154
```


sleepstudy data

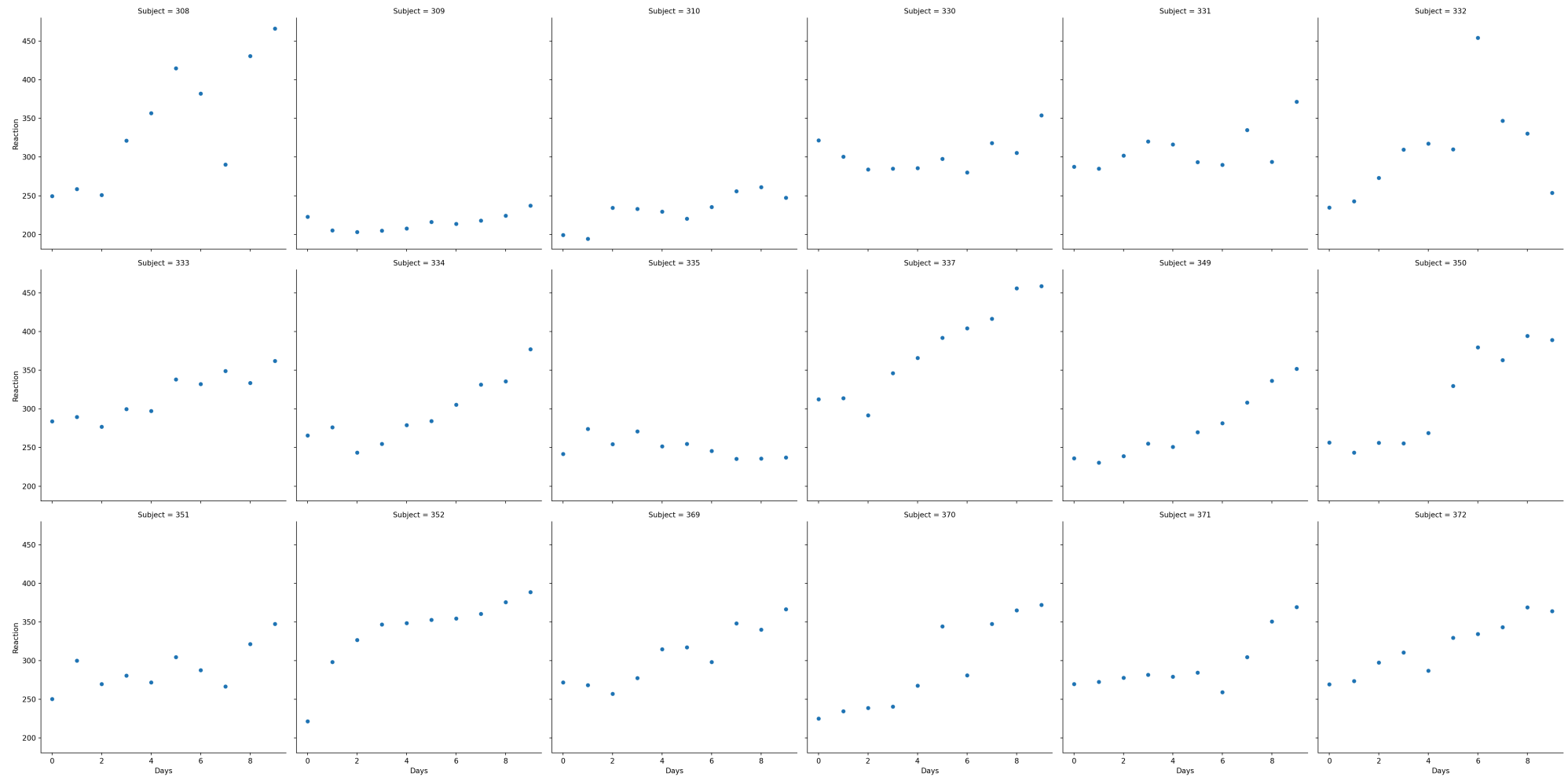
These data are from the study described in Belenky et al. (2003), for the most sleep-deprived group (3 hours time-in-bed) and for the first 10 days of the study, up to the recovery period. The original study analyzed speed ($1/(\text{reaction time})$) and treated day as a categorical rather than a continuous predictor.

```
1 sleep = pd.read_csv("data/sleepstudy.csv")
2 sleep
```

	Reaction	Days	Subject
0	249.5600	0	308
1	258.7047	1	308
2	250.8006	2	308
3	321.4398	3	308
4	356.8519	4	308
..
175	329.6076	5	372
176	334.4818	6	372
177	343.2199	7	372
178	369.1417	8	372
179	364.1236	9	372

[180 rows x 3 columns]

```
1 g = sns.relplot(x="Days", y="Reaction", col="Subject", col_wrap=6, data=sleep)
```



Random intercept model

```
1 me_rand_int = smf.mixedlm(  
2     "Reaction ~ Days", data=sleep, groups=sleep["Subject"],  
3     subset=sleep.Days >= 2  
4 )  
5 res_rand_int = me_rand_int.fit(method=["lbfgs"])  
6 print(res_rand_int.summary())
```

Mixed Linear Model Regression Results

```
=====
Model:                MixedLM Dependent Variable: Reaction
No. Observations:    180      Method:                REML
No. Groups:          18      Scale:                 960.4529
Min. group size:     10      Log-Likelihood:    -893.2325
Max. group size:     10      Converged:         Yes
Mean group size:     10.0

-----
              Coef.   Std.Err.    z      P>|z|   [0.025   0.975]
-----
Intercept    251.405     9.747  25.793  0.000   232.302  270.509
Days         10.467     0.804  13.015  0.000     8.891  12.044
Group Var    1378.232    17.157

=====
```

lme4 version

```
1 summary(  
2   lmer(Reaction ~ Days + (1|Subject), data=sleepstudy)  
3 )
```

Linear mixed model fit by REML ['lmerMod']

Formula: Reaction ~ Days + (1 | Subject)

Data: sleepstudy

REML criterion at convergence: 1786.5

Scaled residuals:

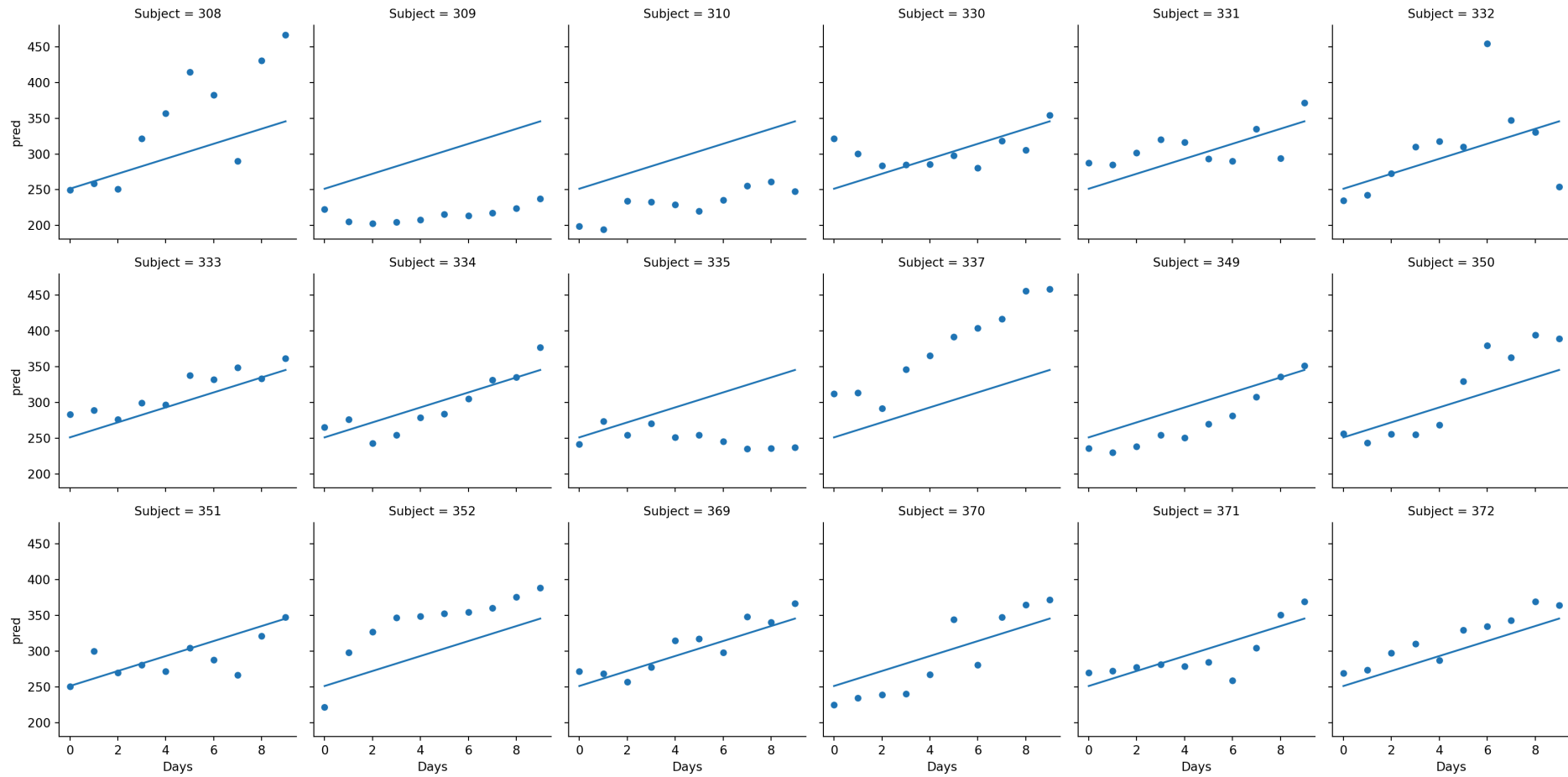
Min	1Q	Median	3Q	Max
-3.2257	-0.5529	0.0109	0.5188	4.2506

Random effects:

Groups	Name	Variance	Std.Dev.
Subject	(Intercept)	1378.2	37.12
Residual		960.5	30.99

Number of obs: 180, groups: Subject, 18

Predictions

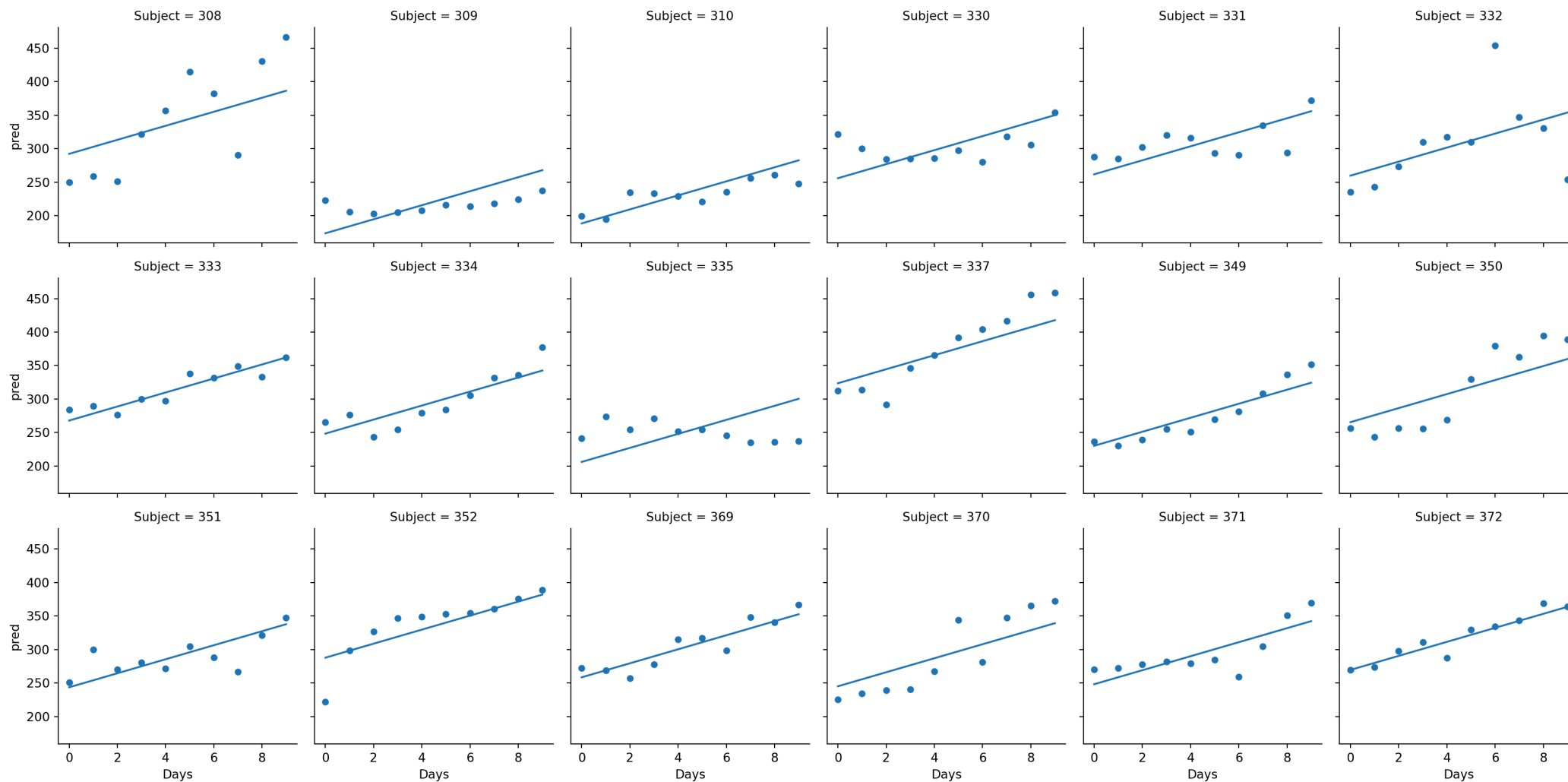


Recovering random effects for prediction

```
1 # Multiply each RE by the random effects design matrix for each group
2 rex = [
3     np.dot(
4         me_rand_int.exog_re_li[j],
5         res_rand_int.random_effects[k]
6     )
7     for (j, k) in enumerate(me_rand_int.group_labels)
8 ]
9 rex[0]
```

```
array([40.7838, 40.7838, 40.7838, 40.7838, 40.7838, 40.7838, 40.7838, 40.7838,
       40.7838, 40.7838])
```

```
1
2 # Add the fixed and random terms to get the overall prediction
3 y_hat = res_rand_int.predict() + np.concatenate(rex)
```



Random intercept and slope model

```
1 me_rand_sl= smf.mixedlm(  
2     "Reaction ~ Days", data=sleep, groups=sleep["Subject"],  
3     subset=sleep.Days >= 2,  
4     re_formula=~Days"  
5 )  
6 res_rand_sl = me_rand_sl.fit(method=["lbfgs"])  
7 print(res_rand_sl.summary())
```

Mixed Linear Model Regression Results

=====						
Model:	MixedLM	Dependent Variable:		Reaction		
No. Observations:	180	Method:		REML		
No. Groups:	18	Scale:		654.9412		
Min. group size:	10	Log-Likelihood:		-871.8141		
Max. group size:	10	Converged:		Yes		
Mean group size:	10.0					

	Coef.	Std.Err.	z	P> z	[0.025	0.975]

Intercept	251.405	6.825	36.838	0.000	238.029	264.781
Days	10.467	1.546	6.771	0.000	7.438	13.497
Group Var	612.089	11.881				
Group x Days Cov	9.605	1.820				
Days Var	35.072	0.610				
=====						

lme4 version

```
1 summary(  
2   lmer(Reaction ~ Days + (Days|Subject), data=sleepstudy)  
3 )
```

Linear mixed model fit by REML ['lmerMod']
Formula: Reaction ~ Days + (Days | Subject)
Data: sleepstudy

REML criterion at convergence: 1743.6

Scaled residuals:

Min	1Q	Median	3Q	Max
-3.9536	-0.4634	0.0231	0.4634	5.1793

Random effects:

Groups	Name	Variance	Std.Dev.	Corr
Subject	(Intercept)	612.10	24.741	
	Days	35.07	5.922	0.07
Residual		654.94	25.592	

Number of obs: 180, groups: Subject, 18

Fixed effects:

Prediction

