Linear Models

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```
.. currentmodule:: sklearn.linear_model
```

The following are a set of methods intended for regression in which the target value is expected to be a linear combination of the features. In mathematical notation, if $\hat{y}$ is the predicted value.

$$\hat{y}(w, x) = w_0 + w_1 x_1 + ... + w_p x_p$$

Across the module, we designate the vector $w = (w_1, ..., w_p)$ as `coef_` and w_0 as `intercept_`.

To perform classification with generalized linear models, see :ref:`Logistic_regression`.

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Ordinary Least Squares

:class:`LinearRegression` fits a linear model with coefficients $w = (w_1, ..., w_p)$ to minimize the residual sum of squares between the observed targets in the dataset, and the targets predicted by the linear approximation. Mathematically it solves a problem of the form:

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$$\min_w ||Xw - y||_2^2$$

:class:`LinearRegression` will take in its `fit` method arrays X, y and will store the coefficients w of the linear model in its `coef_` member:

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```
>>> from sklearn import linear_model
>>> reg = linear_model.LinearRegression()
>>> reg.fit([[0, 0], [1, 1], [2, 2]], [0, 1, 2])
LinearRegression()
>>> reg.coef_
array([0.5, 0.5])
```

The coefficient estimates for Ordinary Least Squares rely on the independence of the features. When features are correlated and the columns of the design matrix X have an approximately linear dependence, the design matrix becomes close to singular and as a result, the least-squares estimate becomes highly sensitive to random errors in the observed target, producing a large variance. This situation of *multicollinearity* can arise, for example, when data are collected without an experimental design.

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_ols.py`

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Non-Negative Least Squares

It is possible to constrain all the coefficients to be non-negative, which may be useful when they represent some physical or naturally non-negative quantities (e.g., frequency counts or prices of goods). :class:`LinearRegression` accepts a boolean `positive` parameter: when set to *True* Non-Negative Least Squares are then applied.

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Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_nnls.py`

> **System Message: ERROR/3 (D:\onboarding-resources\sample-onboarding-resources\scikit-learn-main\doc\modules\(scikit-learn-main)(doc)(modules)linear_model.rst, line 76); *backlink***
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Ordinary Least Squares Complexity

The least squares solution is computed using the singular value decomposition of X. If X is a matrix of shape *(n_samples, n_features)* this method has a cost of $O(n_{\text{samples}} n^2_{\text{features}})$, assuming that $n_{\text{samples}} \geq n_{\text{features}}$.

Ridge regression and classification

Regression

:class:`Ridge` regression addresses some of the problems of :ref:`ordinary_least_squares` by imposing a penalty on the size of the coefficients. The ridge coefficients minimize a penalized residual sum of squares:

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$$\min_w ||Xw - y||_2^2 + \alpha ||w||_2^2$$

The complexity parameter $\alpha \geq 0$ controls the amount of shrinkage: the larger the value of α, the greater the amount of shrinkage and thus the coefficients become more robust to collinearity.

As with other linear models, :class:`Ridge` will take in its `fit` method arrays X, y and will store the coefficients w of the linear model in its `coef_` member:

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```
>>> from sklearn import linear_model
>>> reg = linear_model.Ridge(alpha=.5)
>>> reg.fit([[0, 0], [0, 0], [1, 1]], [0, .1, 1])
Ridge(alpha=0.5)
>>> reg.coef_
array([0.34545455, 0.34545455])
>>> reg.intercept_
0.13636...
```

Classification

The :class:`Ridge` regressor has a classifier variant: :class:`RidgeClassifier`. This classifier first converts binary targets to `{-1, 1}` and then treats the problem as a regression task, optimizing the same objective as above. The predicted class corresponds to the sign of the regressor's prediction. For multiclass classification, the problem is treated as multi-output regression, and the predicted class corresponds to the output with the highest value.

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It might seem questionable to use a (penalized) Least Squares loss to fit a classification model instead of the more traditional logistic or hinge losses. However, in practice, all those models can lead to similar cross-validation scores in terms of accuracy or precision/recall, while the penalized least squares loss used by the :class:`RidgeClassifier` allows for a very different choice of the numerical solvers with distinct computational performance profiles.

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The :class:`RidgeClassifier` can be significantly faster than e.g. :class:`LogisticRegression` with a high number of classes because it can compute the projection matrix $(X^T X)^{-1} X^T$ only once.

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This classifier is sometimes referred to as a Least Squares Support Vector Machines with a linear kernel.

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_ridge_path.py`

  > **System Message: ERROR/3** (D:\onboarding-resources\sample-onboarding-resources\scikit-learn-main\doc\modules\(scikit-learn-main)(doc)(modules)linear_model.rst, **line 160**); *backlink*
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- :ref:`sphx_glr_auto_examples_text_plot_document_classification_20newsgroups.py`

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- :ref:`sphx_glr_auto_examples_inspection_plot_linear_model_coefficient_interpretation.py`

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Ridge Complexity

This method has the same order of complexity as :ref:`ordinary_least_squares`.

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Setting the regularization parameter: leave-one-out Cross-Validation

:class:`RidgeCV` implements ridge regression with built-in cross-validation of the alpha parameter. The object works in the same way as GridSearchCV except that it defaults to Leave-One-Out Cross-Validation:

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```
>>> import numpy as np
>>> from sklearn import linear_model
>>> reg = linear_model.RidgeCV(alphas=np.logspace(-6, 6, 13))
>>> reg.fit([[0, 0], [0, 0], [1, 1]], [0, .1, 1])
RidgeCV(alphas=array([1.e-06, 1.e-05, 1.e-04, 1.e-03, 1.e-02, 1.e-01, 1.e+00, 1.e+01,
      1.e+02, 1.e+03, 1.e+04, 1.e+05, 1.e+06]))
>>> reg.alpha_
0.01
```

Specifying the value of the :term:`cv` attribute will trigger the use of cross-validation with :class:`~sklearn.model_selection.GridSearchCV`, for example *cv=10* for 10-fold cross-validation, rather than Leave-One-Out Cross-Validation.

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References

- "Notes on Regularized Least Squares", Rifkin & Lippert (technical report, course slides).

Lasso

The :class:`Lasso` is a linear model that estimates sparse coefficients. It is useful in some contexts due to its tendency to prefer solutions with fewer non-zero coefficients, effectively reducing the number of features upon which the given solution is dependent. For this reason, Lasso and its variants are fundamental to the field of compressed sensing. Under certain conditions, it can recover the exact set of non-zero coefficients (see :ref:`sphx_glr_auto_examples_applications_plot_tomography_l1_reconstruction.py`).

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Mathematically, it consists of a linear model with an added regularization term. The objective function to minimize is:

$$\min_w \frac{1}{2n_{\text{samples}}} \|Xw - y\|_2^2 + \alpha \|w\|_1$$

The lasso estimate thus solves the minimization of the least-squares penalty with $\alpha\|w\|_1$ added, where α is a constant and $\|w\|_1$ is the ℓ_1-norm of the coefficient vector.

The implementation in the class :class:`Lasso` uses coordinate descent as the algorithm to fit the coefficients. See :ref:`least_angle_regression` for another implementation:

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```
>>> from sklearn import linear_model
>>> reg = linear_model.Lasso(alpha=0.1)
>>> reg.fit([[0, 0], [1, 1]], [0, 1])
Lasso(alpha=0.1)
>>> reg.predict([[1, 1]])
array([0.8])
```

The function :func:`lasso_path` is useful for lower-level tasks, as it computes the coefficients along the full path of possible values.

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Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_lasso_and_elasticnet.py`

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- :ref:`sphx_glr_auto_examples_applications_plot_tomography_l1_reconstruction.py`

  System Message: ERROR/3 (D:\onboarding-resources\sample-onboarding-resources\scikit-learn-main\doc\modules\(scikit-learn-main)(doc)(modules)linear_model.rst, line 246); *backlink*

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- :ref:`sphx_glr_auto_examples_inspection_plot_linear_model_coefficient_interpretation.py`

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Note

Feature selection with Lasso

As the Lasso regression yields sparse models, it can thus be used to perform feature selection, as detailed in :ref:`l1_feature_selection`.

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The following two references explain the iterations used in the coordinate descent solver of scikit-learn, as well as the duality gap computation used for convergence control.

References

- "Regularization Path For Generalized linear Models by Coordinate Descent", Friedman, Hastie & Tibshirani, J Stat Softw, 2010 (Paper).
- "An Interior-Point Method for Large-Scale L1-Regularized Least Squares," S. J. Kim, K. Koh, M. Lustig, S. Boyd and D. Gorinevsky, in IEEE Journal of Selected Topics in Signal Processing, 2007 (Paper)

Setting regularization parameter

The `alpha` parameter controls the degree of sparsity of the estimated coefficients.

Using cross-validation

scikit-learn exposes objects that set the Lasso `alpha` parameter by cross-validation: :class:`LassoCV` and :class:`LassoLarsCV`. :class:`LassoLarsCV` is based on the :ref:`least_angle_regression` algorithm explained below.

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For high-dimensional datasets with many collinear features, :class:`LassoCV` is most often preferable. However, :class:`LassoLarsCV` has the advantage of exploring more relevant values of *alpha* parameter, and if the number of samples is very small compared to the number of features, it is often faster than :class:`LassoCV`.

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```
.. centered:: |lasso_cv_1| |lasso_cv_2|
```

Information-criteria based model selection

Alternatively, the estimator :class:`LassoLarsIC` proposes to use the Akaike information criterion (AIC) and the Bayes Information criterion (BIC). It is a computationally cheaper alternative to find the optimal value of alpha as the regularization path is computed only once instead of k+1 times when using k-fold cross-validation.

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Indeed, these criteria are computed on the in-sample training set. In short, they penalize the over-optimistic scores of the different Lasso models by their flexibility (cf. to "Mathematical details" section below).

However, such criteria need a proper estimation of the degrees of freedom of the solution, are derived for large samples (asymptotic results) and assume the correct model is candidates under investigation. They also tend to break when the problem is badly conditioned (e.g. more features than samples).

Mathematical details

The definition of AIC (and thus BIC) might differ in the literature. In this section, we give more information regarding the criterion computed in scikit-learn. The AIC criterion is defined as:

$$AIC = -2\log(\hat{L}) + 2d$$

where $\hat{L}$ is the maximum likelihood of the model and d is the number of parameters (as well referred to as degrees of freedom in the previous section).

The definition of BIC replace the constant 2 by $\log(N)$:

$$BIC = -2\log(\hat{L}) + \log(N)d$$

where N is the number of samples.

For a linear Gaussian model, the maximum log-likelihood is defined as:

$$\log(\hat{L}) = -\frac{n}{2}\log(2\pi) - \frac{n}{2}\ln(\sigma^2) - \frac{\sum_{i=1}^{n}(y_i - \hat{y}_i)^2}{2\sigma^2}$$

where σ^2 is an estimate of the noise variance, y_i and $\hat{y}_i$ are respectively the true and predicted targets, and n is the number of samples.

Plugging the maximum log-likelihood in the AIC formula yields:

$$AIC = n\log(2\pi\sigma^2) + \frac{\sum_{i=1}^{n}(y_i - \hat{y}_i)^2}{\sigma^2} + 2d$$

The first term of the above expression is sometimes discarded since it is a constant when σ^2 is provided. In addition, it is sometimes stated that the AIC is equivalent to the C_p statistic [12]. In a strict sense, however, it is equivalent only up to some constant and a multiplicative factor.

At last, we mentioned above that σ^2 is an estimate of the noise variance. In :class:`LassoLarsIC` when the parameter *noise_variance* is not provided (default), the noise variance is estimated via the unbiased estimator [13] defined as:

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$$\sigma^2 = \frac{\sum_{i=1}^{n} (y_i - \hat{y}_i)^2}{n - p}$$

where p is the number of features and $\hat{y}_i$ is the predicted target using an ordinary least squares regression. Note, that this formula is valid only when *n_samples* > *n_features*.

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_lasso_model_selection.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_lasso_lars_ic.py`

  > **System Message: ERROR/3 (D:\onboarding-resources\sample-onboarding-resources\scikit-learn-main\doc\modules\(scikit-learn-main)(doc)(modules)linear_model.rst, line 383); *backlink***
  >
  > Unknown interpreted text role "ref".

References

[12] :arxiv:`Zou, Hui, Trevor Hastie, and Robert Tibshirani. "On the degrees of freedom of the lasso." The Annals of Statistics 35.5 (2007): 2173-2192. <0712.0881.pdf>`

> **System Message: ERROR/3 (D:\onboarding-resources\sample-onboarding-resources\scikit-learn-main\doc\modules\(scikit-learn-main)(doc)(modules)linear_model.rst, line 387); *backlink***
>
> Unknown interpreted text role "arxiv".

[13] Cherkassky, Vladimir, and Yunqian Ma. "Comparison of model selection for regression." Neural computation 15.7 (2003): 1691-1714.

Comparison with the regularization parameter of SVM

The equivalence between `alpha` and the regularization parameter of SVM, `C` is given by `alpha = 1 / C` or `alpha = 1 / (n_samples * C)`, depending on the estimator and the exact objective function optimized by the model.

Multi-task Lasso

The :class:`MultiTaskLasso` is a linear model that estimates sparse coefficients for multiple regression problems jointly: `y` is a 2D array, of shape `(n_samples, n_tasks)`. The constraint is that the selected features are the same for all the regression problems, also called tasks.

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The following figure compares the location of the non-zero entries in the coefficient matrix W obtained with a simple Lasso or a MultiTaskLasso. The Lasso estimates yield scattered non-zeros while the non-zeros of the MultiTaskLasso are full columns.

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```
.. centered:: |multi_task_lasso_1| |multi_task_lasso_2|
```

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```
.. centered:: Fitting a time-series model, imposing that any active feature be active at all times
```

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_multi_task_lasso_support.py`

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Mathematically, it consists of a linear model trained with a mixed ℓ_1 ℓ_2-norm for regularization. The objective function to minimize is:

$$\min_W \frac{1}{2n_{\text{samples}}} \|XW - Y\|_{\text{Fro}}^2 + \alpha \|W\|_{21}$$

where Fro indicates the Frobenius norm

$$\|A\|_{\text{Fro}} = \sqrt{\sum_{ij} a_{ij}^2}$$

and ℓ_1 ℓ_2 reads

$$\|A\|_{21} = \sum_i \sqrt{\sum_j a_{ij}^2}.$$

The implementation in the class :class:`MultiTaskLasso` uses coordinate descent as the algorithm to fit the coefficients.

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Elastic-Net

:class:`ElasticNet` is a linear regression model trained with both ℓ_1 and ℓ_2-norm regularization of the coefficients. This combination allows for learning a sparse model where few of the weights are non-zero like :class:`Lasso`, while still maintaining the regularization properties of :class:`Ridge`. We control the convex combination of ℓ_1 and ℓ_2 using the `l1_ratio` parameter.

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Elastic-net is useful when there are multiple features that are correlated with one another. Lasso is likely to pick one of these at random, while elastic-net is likely to pick both.

A practical advantage of trading-off between Lasso and Ridge is that it allows Elastic-Net to inherit some of Ridge's stability under rotation.

The objective function to minimize is in this case

$$\min_w \frac{1}{2n_{\text{samples}}} \|Xw - y\|_2^2 + \alpha\rho\|w\|_1 + \frac{\alpha(1-\rho)}{2}\|w\|_2^2$$

The class :class:`ElasticNetCV` can be used to set the parameters `alpha` (α) and `l1_ratio` (ρ) by cross-validation.

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Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_lasso_and_elasticnet.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_lasso_coordinate_descent_path.py`

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The following two references explain the iterations used in the coordinate descent solver of scikit-learn, as well as the duality gap computation used for convergence control.

References

- "Regularization Path For Generalized linear Models by Coordinate Descent", Friedman, Hastie & Tibshirani, J Stat Softw, 2010 (Paper).
- "An Interior-Point Method for Large-Scale L1-Regularized Least Squares," S. J. Kim, K. Koh, M. Lustig, S. Boyd and D. Gorinevsky, in IEEE Journal of Selected Topics in Signal Processing, 2007 (Paper)

Multi-task Elastic-Net

The :class:`MultiTaskElasticNet` is an elastic-net model that estimates sparse coefficients for multiple regression problems jointly: `Y` is a 2D array of shape `(n_samples, n_tasks)`. The constraint is that the selected features are the same for all the regression problems, also called tasks.

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Mathematically, it consists of a linear model trained with a mixed ℓ_1 ℓ_2-norm and ℓ_2-norm for regularization. The objective function to minimize is:

$$\min_W \frac{1}{2n_{\text{samples}}}\|XW - Y\|^2_{\text{Fro}} + \alpha\rho\|W\|_{21} + \frac{\alpha(1-\rho)}{2}\|W\|^2_{\text{Fro}}$$

The implementation in the class :class:`MultiTaskElasticNet` uses coordinate descent as the algorithm to fit the coefficients.

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The class :class:`MultiTaskElasticNetCV` can be used to set the parameters `alpha` (α) and `l1_ratio` (ρ) by cross-validation.

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Least Angle Regression

Least-angle regression (LARS) is a regression algorithm for high-dimensional data, developed by Bradley Efron, Trevor Hastie, Iain Johnstone and Robert Tibshirani. LARS is similar to forward stepwise regression. At each step, it finds the feature most correlated with the target. When there are multiple features having equal correlation, instead of continuing along the same feature, it proceeds in a direction equiangular between the features.

The advantages of LARS are:

- It is numerically efficient in contexts where the number of features is significantly greater than the number of samples.
- It is computationally just as fast as forward selection and has the same order of complexity as ordinary least squares.
- It produces a full piecewise linear solution path, which is useful in cross-validation or similar attempts to tune the model.
- If two features are almost equally correlated with the target, then their coefficients should increase at approximately the same rate. The algorithm thus behaves as intuition would expect, and also is more stable.
- It is easily modified to produce solutions for other estimators, like the Lasso.

The disadvantages of the LARS method include:

- Because LARS is based upon an iterative refitting of the residuals, it would appear to be especially sensitive to the effects of noise. This problem is discussed in detail by Weisberg in the discussion section of the Efron et al. (2004) Annals of Statistics article.

The LARS model can be used using via the estimator :class:`Lars`, or its low-level implementation :func:`lars_path` or :func:`lars_path_gram`.

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LARS Lasso

:class:`LassoLars` is a lasso model implemented using the LARS algorithm, and unlike the implementation based on coordinate descent, this yields the exact solution, which is piecewise linear as a function of the norm of its coefficients.

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```
>>> from sklearn import linear_model
>>> reg = linear_model.LassoLars(alpha=.1, normalize=False)
>>> reg.fit([[0, 0], [1, 1]], [0, 1])
LassoLars(alpha=0.1, normalize=False)
>>> reg.coef_
array([0.6..., 0.        ])
```

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_lasso_lars.py`

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The Lars algorithm provides the full path of the coefficients along the regularization parameter almost for free, thus a common operation is to retrieve the path with one of the functions :func:`lars_path` or :func:`lars_path_gram`.

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Mathematical formulation

The algorithm is similar to forward stepwise regression, but instead of including features at each step, the estimated coefficients are increased in a direction equiangular to each one's correlations with the residual.

Instead of giving a vector result, the LARS solution consists of a curve denoting the solution for each value of the ℓ_1 norm of the parameter vector. The full coefficients path is stored in the array `coef_path_` of shape *(n_features, max_features + 1)*. The first column is always zero.

References:

- Original Algorithm is detailed in the paper Least Angle Regression by Hastie et al.

Orthogonal Matching Pursuit (OMP)

:class:`OrthogonalMatchingPursuit` and :func:`orthogonal_mp` implements the OMP algorithm for approximating the fit of a linear model with constraints imposed on the number of non-zero coefficients (ie. the ℓ_0 pseudo-norm).

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Being a forward feature selection method like :ref:`least_angle_regression`, orthogonal matching pursuit can approximate the optimum solution vector with a fixed number of non-zero elements:

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$$\underset{w}{\operatorname{arg\,min}} \|y - Xw\|_2^2 \text{ subject to } \|w\|_0 \leq n_{\text{nonzero\_coefs}}$$

Alternatively, orthogonal matching pursuit can target a specific error instead of a specific number of non-zero coefficients. This can be expressed as:

$$\underset{w}{\operatorname{arg\,min}} \|w\|_0 \text{ subject to } \|y - Xw\|_2^2 \leq \text{tol}$$

OMP is based on a greedy algorithm that includes at each step the atom most highly correlated with the current residual. It is similar to the simpler matching pursuit (MP) method, but better in that at each iteration, the residual is recomputed using an orthogonal projection on the space of the previously chosen dictionary elements.

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_omp.py`

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References:

- https://www.cs.technion.ac.il/~ronrubin/Publications/KSVD-OMP-v2.pdf
- Matching pursuits with time-frequency dictionaries, S. G. Mallat, Z. Zhang,

Bayesian Regression

Bayesian regression techniques can be used to include regularization parameters in the estimation procedure: the regularization parameter is not set in a hard sense but tuned to the data at hand.

This can be done by introducing uninformative priors over the hyper parameters of the model. The ℓ_2 regularization used in :ref:`ridge_regression` is equivalent to finding a maximum a posteriori estimation under a Gaussian prior over the coefficients w with precision λ^{-1}. Instead of setting *lambda* manually, it is possible to treat it as a random variable to be estimated from the data.

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To obtain a fully probabilistic model, the output y is assumed to be Gaussian distributed around Xw:

$$p(y|X, w, \alpha) = \mathcal{N}(y|Xw, \alpha)$$

where α is again treated as a random variable that is to be estimated from the data.

The advantages of Bayesian Regression are:

- It adapts to the data at hand.
- It can be used to include regularization parameters in the estimation procedure.

The disadvantages of Bayesian regression include:

- Inference of the model can be time consuming.

References

- A good introduction to Bayesian methods is given in C. Bishop: Pattern Recognition and Machine learning
- Original Algorithm is detailed in the book *Bayesian learning for neural networks* by Radford M. Neal

Bayesian Ridge Regression

:class:`BayesianRidge` estimates a probabilistic model of the regression problem as described above. The prior for the coefficient w is given by a spherical Gaussian:

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$$p(w|\lambda) = \mathcal{N}(w|0, \lambda^{-1}\mathbf{I}_p)$$

The priors over α and λ are chosen to be gamma distributions, the conjugate prior for the precision of the Gaussian. The resulting model is called *Bayesian Ridge Regression*, and is similar to the classical :class:`Ridge`.

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The parameters w, α and λ are estimated jointly during the fit of the model, the regularization parameters α and λ being estimated by maximizing the *log marginal likelihood*. The scikit-learn implementation is based on the algorithm described in Appendix A of (Tipping, 2001) where the update of the parameters α and λ is done as suggested in (MacKay, 1992). The initial value of the maximization procedure can be set with the hyperparameters `alpha_init` and `lambda_init`.

There are four more hyperparameters, α_1, α_2, λ_1 and λ_2 of the gamma prior distributions over α and λ. These are usually chosen to be *non-informative*. By default $\alpha_1 = \alpha_2 = \lambda_1 = \lambda_2 = 10^{-6}$.

Bayesian Ridge Regression is used for regression:

```
>>> from sklearn import linear_model
>>> X = [[0., 0.], [1., 1.], [2., 2.], [3., 3.]]
>>> Y = [0., 1., 2., 3.]
>>> reg = linear_model.BayesianRidge()
>>> reg.fit(X, Y)
BayesianRidge()
```

After being fitted, the model can then be used to predict new values:

```
>>> reg.predict([[1, 0.]])
array([0.50000013])
```

The coefficients w of the model can be accessed:

```
>>> reg.coef_
array([0.49999993, 0.49999993])
```

Due to the Bayesian framework, the weights found are slightly different to the ones found by :ref:`ordinary_least_squares`. However, Bayesian Ridge Regression is more robust to ill-posed problems.

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Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_bayesian_ridge.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_bayesian_ridge_curvefit.py`

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References:

- Section 3.3 in Christopher M. Bishop: Pattern Recognition and Machine Learning, 2006
- David J. C. MacKay, Bayesian Interpolation, 1992.
- Michael E. Tipping, Sparse Bayesian Learning and the Relevance Vector Machine, 2001.

Automatic Relevance Determination - ARD

:class:`ARDRegression` is very similar to Bayesian Ridge Regression, but can lead to sparser coefficients w [1] [2]. :class:`ARDRegression` poses a different prior over w, by dropping the assumption of the Gaussian being spherical.

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Instead, the distribution over w is assumed to be an axis-parallel, elliptical Gaussian distribution.

This means each coefficient w_i is drawn from a Gaussian distribution, centered on zero and with a precision λ_i:

$$p(w|\lambda) = \mathcal{N}(w|0, A^{-1})$$

with $\text{diag}(A) = \lambda = \{\lambda_1, ..., \lambda_p\}$.

In contrast to Bayesian Ridge Regression, each coordinate of w_i has its own standard deviation λ_i. The prior over all λ_i is chosen to be the same gamma distribution given by hyperparameters λ_1 and λ_2.

ARD is also known in the literature as *Sparse Bayesian Learning* and *Relevance Vector Machine* [3] [4].

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_ard.py`

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References:

[1] Christopher M. Bishop: Pattern Recognition and Machine Learning, Chapter 7.2.1

[2] David Wipf and Srikantan Nagarajan: A new view of automatic relevance determination

[3] Michael E. Tipping: Sparse Bayesian Learning and the Relevance Vector Machine

[4] Tristan Fletcher: Relevance Vector Machines explained

Logistic regression

Logistic regression, despite its name, is a linear model for classification rather than regression. Logistic regression is also known in the literature as logit regression, maximum-entropy classification (MaxEnt) or the log-linear classifier. In this model, the probabilities describing the possible outcomes of a single trial are modeled using a logistic function.

Logistic regression is implemented in :class:`LogisticRegression`. This implementation can fit binary, One-vs-Rest, or multinomial logistic regression with optional ℓ_1, ℓ_2 or Elastic-Net regularization.

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> **Note**
>
> Regularization is applied by default, which is common in machine learning but not in statistics. Another advantage of regularization is that it improves numerical stability. No regularization amounts to setting C to a very high value.

As an optimization problem, binary class ℓ_2 penalized logistic regression minimizes the following cost function:

$$\min_{w, c} \frac{1}{2} w^T w + C \sum_{i=1}^{n} \log(\exp(-y_i(X_i^T w + c)) + 1).$$

Similarly, ℓ_1 regularized logistic regression solves the following optimization problem:

$$\min_{w, c} \|w\|_1 + C \sum_{i=1}^{n} \log(\exp(-y_i(X_i^T w + c)) + 1).$$

Elastic-Net regularization is a combination of ℓ_1 and ℓ_2, and minimizes the following cost function:

$$\min_{w, c} \frac{1 - \rho}{2} w^T w + \rho \|w\|_1 + C \sum_{i=1}^{n} \log(\exp(-y_i(X_i^T w + c)) + 1),$$

where ρ controls the strength of ℓ_1 regularization vs. ℓ_2 regularization (it corresponds to the *l1_ratio* parameter).

Note that, in this notation, it's assumed that the target y_i takes values in the set $-1, 1$ at trial i. We can also see that Elastic-Net is equivalent to ℓ_1 when $\rho = 1$ and equivalent to ℓ_2 when $\rho = 0$.

The solvers implemented in the class :class:`LogisticRegression` are "liblinear", "newton-cg", "lbfgs", "sag" and "saga":

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The solver "liblinear" uses a coordinate descent (CD) algorithm, and relies on the excellent C++ LIBLINEAR library, which is shipped with scikit-learn. However, the CD algorithm implemented in liblinear cannot learn a true multinomial (multiclass) model; instead, the optimization problem is decomposed in a "one-vs-rest" fashion so separate binary classifiers are trained for all classes. This happens under the hood, so :class:`LogisticRegression` instances using this solver behave as multiclass classifiers. For ℓ_1 regularization :func:`sklearn.svm.l1_min_c` allows to calculate the lower bound for C in order to get a non "null" (all feature weights to zero) model.

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The "lbfgs", "sag" and "newton-cg" solvers only support ℓ_2 regularization or no regularization, and are found to converge faster for some high-dimensional data. Setting *multi_class* to "multinomial" with these solvers learns a true multinomial logistic regression model [5], which means that its probability estimates should be better calibrated than the default "one-vs-rest" setting.

The "sag" solver uses Stochastic Average Gradient descent [6]. It is faster than other solvers for large datasets, when both the number of samples and the number of features are large.

The "saga" solver [7] is a variant of "sag" that also supports the non-smooth *penalty="l1"*. This is therefore the solver of choice for sparse multinomial logistic regression. It is also the only solver that supports *penalty="elasticnet"*.

The "lbfgs" is an optimization algorithm that approximates the Broydenâ€“Fletcherâ€“Goldfarbâ€“Shanno algorithm [8], which belongs to quasi-Newton methods. The "lbfgs" solver is recommended for use for small data-sets but for larger datasets its performance suffers. [9]

The following table summarizes the penalties supported by each solver:

	Solvers				
Penalties	**'liblinear'**	**'lbfgs'**	**'newton-cg'**	**'sag'**	**'saga'**
Multinomial + L2 penalty	no	yes	yes	yes	yes
OVR + L2 penalty	yes	yes	yes	yes	yes
Multinomial + L1 penalty	no	no	no	no	yes
OVR + L1 penalty	yes	no	no	no	yes
Elastic-Net	no	no	no	no	yes
No penalty ('none')	no	yes	yes	yes	yes
Behaviors					
Penalize the intercept (bad)	yes	no	no	no	no
Faster for large datasets	no	no	no	yes	yes
Robust to unscaled datasets	yes	yes	yes	no	no

The "lbfgs" solver is used by default for its robustness. For large datasets the "saga" solver is usually faster. For large dataset, you may also consider using :class:`SGDClassifier` with 'log' loss, which might be even faster but requires more tuning.

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Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_logistic_l1_l2_sparsity.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_logistic_path.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_logistic_multinomial.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_sparse_logistic_regression_20newsgroups.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_sparse_logistic_regression_mnist.py`

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Differences from liblinear:

There might be a difference in the scores obtained between :class:`LogisticRegression` with `solver=liblinear` or :class:`LinearSVC` and the external liblinear library directly, when `fit_intercept=False` and the fit `coef_` (or) the data to be predicted are zeroes. This is because for the sample(s) with `decision_function` zero, :class:`LogisticRegression` and :class:`LinearSVC` predict the negative class, while liblinear predicts the positive class. Note that a model with `fit_intercept=False` and having many samples with `decision_function` zero, is likely to be a underfit, bad model and you are advised to set `fit_intercept=True` and increase the intercept_scaling.

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>
> Unknown interpreted text role "class".

> **Note**
>
> **Feature selection with sparse logistic regression**
>
> A logistic regression with ℓ_1 penalty yields sparse models, and can thus be used to perform feature selection, as detailed in :ref:`l1_feature_selection`.
>
> > **System Message: ERROR/3 (`D:\onboarding-resources\sample-onboarding-resources\scikit-learn-main\doc\modules\(scikit-learn-main)(doc)(modules)linear_model.rst`, line 984); *backlink***
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> **Note**
>
> **P-value estimation**
>
> It is possible to obtain the p-values and confidence intervals for coefficients in cases of regression without penalization. The *statsmodels package <https://pypi.org/project/statsmodels/>* natively supports this. Within sklearn, one could use bootstrapping instead as well.

:class:`LogisticRegressionCV` implements Logistic Regression with built-in cross-validation support, to find the optimal *C* and *l1_ratio* parameters according to the `scoring` attribute. The "newton-cg", "sag", "saga" and "lbfgs" solvers are found to be faster for high-dimensional dense data, due to warm-starting (see :term:`Glossary <warm_start>`).

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References:

[5] Christopher M. Bishop: Pattern Recognition and Machine Learning, Chapter 4.3.4

[6] Mark Schmidt, Nicolas Le Roux, and Francis Bach: Minimizing Finite Sums with the Stochastic Average Gradient.

[7] Aaron Defazio, Francis Bach, Simon Lacoste-Julien: :arxiv:`SAGA: A Fast Incremental Gradient Method With Support for Non-Strongly Convex Composite Objectives. <1407.0202>`

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[8] https://en.wikipedia.org/wiki/Broyden%E2%80%93Fletcher%E2%80%93Goldfarb%E2%80%93Shanno_algorithm

[9] "Performance Evaluation of Lbfgs vs other solvers"

Generalized Linear Regression

Generalized Linear Models (GLM) extend linear models in two ways [10]. First, the predicted values $\hat{y}$ are linked to a linear combination of the input variables X via an inverse link function h as

$$\hat{y}(w, X) = h(Xw).$$

Secondly, the squared loss function is replaced by the unit deviance d of a distribution in the exponential family (or more precisely, a reproductive exponential dispersion model (EDM) [11]).

The minimization problem becomes:

$$\min_w \frac{1}{2n_{\text{samples}}} \sum_i d(y_i, \hat{y}_i) + \frac{\alpha}{2}\|w\|_2^2,$$

where α is the L2 regularization penalty. When sample weights are provided, the average becomes a weighted average.

The following table lists some specific EDMs and their unit deviance (all of these are instances of the Tweedie family):

Distribution	Target Domain	Unit Deviance $d(y, \hat{y})$
Normal	$y \in (-\infty, \infty)$	$(y-\hat{y})^2$
Poisson	$y \in [0, \infty)$	$2(y\log\frac{y}{\hat{y}} - y + \hat{y})$
Gamma	$y \in (0, \infty)$	$2(\log\frac{\hat{y}}{y} + \frac{y}{\hat{y}} - 1)$
Inverse Gaussian	$y \in (0, \infty)$	$\frac{(y-\hat{y})^2}{y\hat{y}^2}$

The Probability Density Functions (PDF) of these distributions are illustrated in the following figure,

PDF of a random variable Y following Poisson, Tweedie (power=1.5) and Gamma distributions with different mean values (μ). Observe the point mass at $Y = 0$ for the Poisson distribution and the Tweedie (power=1.5) distribution, but not for the Gamma distribution which has a strictly positive target domain.

The choice of the distribution depends on the problem at hand:

- If the target values y are counts (non-negative integer valued) or relative frequencies (non-negative), you might use a Poisson

deviance with log-link.
- If the target values are positive valued and skewed, you might try a Gamma deviance with log-link.
- If the target values seem to be heavier tailed than a Gamma distribution, you might try an Inverse Gaussian deviance (or even higher variance powers of the Tweedie family).

Examples of use cases include:

- Agriculture / weather modeling: number of rain events per year (Poisson), amount of rainfall per event (Gamma), total rainfall per year (Tweedie / Compound Poisson Gamma).
- Risk modeling / insurance policy pricing: number of claim events / policyholder per year (Poisson), cost per event (Gamma), total cost per policyholder per year (Tweedie / Compound Poisson Gamma).
- Predictive maintenance: number of production interruption events per year (Poisson), duration of interruption (Gamma), total interruption time per year (Tweedie / Compound Poisson Gamma).

References:

[10] McCullagh, Peter; Nelder, John (1989). Generalized Linear Models, Second Edition. Boca Raton: Chapman and Hall/CRC. ISBN 0-412-31760-5.

[11] JÃ¸rgensen, B. (1992). The theory of exponential dispersion models and analysis of deviance. Monografias de matemÃ¡tica, no. 51. See also Exponential dispersion model.

Usage

:class:`TweedieRegressor` implements a generalized linear model for the Tweedie distribution, that allows to model any of the above mentioned distributions using the appropriate `power` parameter. In particular:

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- `power = 0`: Normal distribution. Specific estimators such as :class:`Ridge`, :class:`ElasticNet` are generally more appropriate in this case.

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- `power = 1`: Poisson distribution. :class:`PoissonRegressor` is exposed for convenience. However, it is strictly equivalent to *TweedieRegressor(power=1, link='log')*.

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- `power = 2`: Gamma distribution. :class:`GammaRegressor` is exposed for convenience. However, it is strictly equivalent to *TweedieRegressor(power=2, link='log')*.

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- `power = 3`: Inverse Gaussian distribution.

The link function is determined by the *link* parameter.

Usage example:

```
>>> from sklearn.linear_model import TweedieRegressor
>>> reg = TweedieRegressor(power=1, alpha=0.5, link='log')
>>> reg.fit([[0, 0], [0, 1], [2, 2]], [0, 1, 2])
TweedieRegressor(alpha=0.5, link='log', power=1)
>>> reg.coef_
array([0.2463..., 0.4337...])
>>> reg.intercept_
-0.7638...
```

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_poisson_regression_non_normal_loss.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_tweedie_regression_insurance_claims.py`

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Practical considerations

The feature matrix X should be standardized before fitting. This ensures that the penalty treats features equally.

Since the linear predictor Xw can be negative and Poisson, Gamma and Inverse Gaussian distributions don't support negative values, it is necessary to apply an inverse link function that guarantees the non-negativeness. For example with *link='log'*, the inverse link function becomes $h(Xw) = \exp(Xw)$.

If you want to model a relative frequency, i.e. counts per exposure (time, volume, ...) you can do so by using a Poisson distribution and passing $y = \frac{\text{counts}}{\text{exposure}}$ as target values together with exposure as sample weights. For a concrete example see e.g. :ref:`sphx_glr_auto_examples_linear_model_plot_tweedie_regression_insurance_claims.py`.

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When performing cross-validation for the *power* parameter of *TweedieRegressor*, it is advisable to specify an explicit *scoring* function, because the default scorer :meth:`TweedieRegressor.score` is a function of *power* itself.

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Stochastic Gradient Descent - SGD

Stochastic gradient descent is a simple yet very efficient approach to fit linear models. It is particularly useful when the number of samples (and the number of features) is very large. The `partial_fit` method allows online/out-of-core learning.

The classes :class:`SGDClassifier` and :class:`SGDRegressor` provide functionality to fit linear models for classification and regression using different (convex) loss functions and different penalties. E.g., with `loss="log"`, :class:`SGDClassifier` fits a logistic regression model, while with `loss="hinge"` it fits a linear support vector machine (SVM).

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References

- :ref:`sgd`

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Perceptron

The :class:`Perceptron` is another simple classification algorithm suitable for large scale learning. By default:

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- It does not require a learning rate.
- It is not regularized (penalized).
- It updates its model only on mistakes.

The last characteristic implies that the Perceptron is slightly faster to train than SGD with the hinge loss and that the resulting models are sparser.

Passive Aggressive Algorithms

The passive-aggressive algorithms are a family of algorithms for large-scale learning. They are similar to the Perceptron in that they do not require a learning rate. However, contrary to the Perceptron, they include a regularization parameter `C`.

For classification, :class:`PassiveAggressiveClassifier` can be used with `loss='hinge'` (PA-I) or `loss='squared_hinge'` (PA-II). For regression, :class:`PassiveAggressiveRegressor` can be used with `loss='epsilon_insensitive'` (PA-I) or `loss='squared_epsilon_insensitive'` (PA-II).

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References:

- "Online Passive-Aggressive Algorithms" K. Crammer, O. Dekel, J. Keshat, S. Shalev-Shwartz, Y. Singer - JMLR 7 (2006)

Robustness regression: outliers and modeling errors

Robust regression aims to fit a regression model in the presence of corrupt data: either outliers, or error in the model.

Different scenario and useful concepts

There are different things to keep in mind when dealing with data corrupted by outliers:

- **Outliers in X or in y?**

Outliers in the y direction	Outliers in the X direction

- **Fraction of outliers versus amplitude of error**

  The number of outlying points matters, but also how much they are outliers.

Small outliers	Large outliers

An important notion of robust fitting is that of breakdown point: the fraction of data that can be outlying for the fit to start missing the inlying data.

Note that in general, robust fitting in high-dimensional setting (large *n_features*) is very hard. The robust models here will probably not work in these settings.

Trade-offs: which estimator?

Scikit-learn provides 3 robust regression estimators: :ref:`RANSAC <ransac_regression>`, :ref:`Theil Sen <theil_sen_regression>` and :ref:`HuberRegressor <huber_regression>`.

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- :ref:`HuberRegressor <huber_regression>` should be faster than :ref:`RANSAC <ransac_regression>` and :ref:`Theil Sen <theil_sen_regression>` unless the number of samples are very large, i.e `n_samples` >> `n_features`. This is because :ref:`RANSAC <ransac_regression>` and :ref:`Theil Sen <theil_sen_regression>` fit on smaller subsets of the data. However, both :ref:`Theil Sen <theil_sen_regression>` and :ref:`RANSAC <ransac_regression>` are unlikely to be as robust as :ref:`HuberRegressor <huber_regression>` for the default parameters.

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- :ref:`RANSAC <ransac_regression>` is faster than :ref:`Theil Sen <theil_sen_regression>` and scales much better with the number of samples.

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- :ref:`RANSAC <ransac_regression>` will deal better with large outliers in the y direction (most common situation).

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- :ref:`Theil Sen <theil_sen_regression>` will cope better with medium-size outliers in the X direction, but this property will disappear in high-dimensional settings.

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When in doubt, use :ref:`RANSAC <ransac_regression>`.

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RANSAC: RANdom SAmple Consensus

RANSAC (RANdom SAmple Consensus) fits a model from random subsets of inliers from the complete data set.

RANSAC is a non-deterministic algorithm producing only a reasonable result with a certain probability, which is dependent on the number of iterations (see *max_trials* parameter). It is typically used for linear and non-linear regression problems and is especially popular in the field of photogrammetric computer vision.

The algorithm splits the complete input sample data into a set of inliers, which may be subject to noise, and outliers, which are e.g. caused by erroneous measurements or invalid hypotheses about the data. The resulting model is then estimated only from the determined inliers.

Details of the algorithm

Each iteration performs the following steps:

1. Select `min_samples` random samples from the original data and check whether the set of data is valid (see `is_data_valid`).
2. Fit a model to the random subset (`base_estimator.fit`) and check whether the estimated model is valid (see `is_model_valid`).
3. Classify all data as inliers or outliers by calculating the residuals to the estimated model (`base_estimator.predict(X) - y`) - all data samples with absolute residuals smaller than or equal to the `residual_threshold` are considered as inliers.
4. Save fitted model as best model if number of inlier samples is maximal. In case the current estimated model has the same number of inliers, it is only considered as the best model if it has better score.

These steps are performed either a maximum number of times (`max_trials`) or until one of the special stop criteria are met (see `stop_n_inliers` and `stop_score`). The final model is estimated using all inlier samples (consensus set) of the previously determined best model.

The `is_data_valid` and `is_model_valid` functions allow to identify and reject degenerate combinations of random sub-samples. If the estimated model is not needed for identifying degenerate cases, `is_data_valid` should be used as it is called prior to fitting the model and thus leading to better computational performance.

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_ransac.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_robust_fit.py`

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  Unknown interpreted text role "ref".

References:

- https://en.wikipedia.org/wiki/RANSAC
- "Random Sample Consensus: A Paradigm for Model Fitting with Applications to Image Analysis and Automated Cartography" Martin A. Fischler and Robert C. Bolles - SRI International (1981)
- "Performance Evaluation of RANSAC Family" Sunglok Choi, Taemin Kim and Wonpil Yu - BMVC (2009)

Theil-Sen estimator: generalized-median-based estimator

The :class:`TheilSenRegressor` estimator uses a generalization of the median in multiple dimensions. It is thus robust to multivariate outliers. Note however that the robustness of the estimator decreases quickly with the dimensionality of the problem. It loses its robustness properties and becomes no better than an ordinary least squares in high dimension.

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Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_theilsen.py`

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- :ref:`sphx_glr_auto_examples_linear_model_plot_robust_fit.py`

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References:

- https://en.wikipedia.org/wiki/Theil%E2%80%93Sen_estimator

Theoretical considerations

:class:`TheilSenRegressor` is comparable to the :ref:`Ordinary Least Squares (OLS) <ordinary_least_squares>` in terms of asymptotic efficiency and as an unbiased estimator. In contrast to OLS, Theil-Sen is a non-parametric method which means it makes no assumption about the underlying distribution of the data. Since Theil-Sen is a median-based estimator, it is more robust against corrupted data aka outliers. In univariate setting, Theil-Sen has a breakdown point of about 29.3% in case of a simple linear regression which means that it can tolerate arbitrary corrupted data of up to 29.3%.

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The implementation of :class:`TheilSenRegressor` in scikit-learn follows a generalization to a multivariate linear regression model [14] using the spatial median which is a generalization of the median to multiple dimensions [15].

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In terms of time and space complexity, Theil-Sen scales according to

$$\binom{n_{\text{samples}}}{n_{\text{subsamples}}}$$

which makes it infeasible to be applied exhaustively to problems with a large number of samples and features. Therefore, the magnitude of a subpopulation can be chosen to limit the time and space complexity by considering only a random subset of all possible combinations.

Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_theilsen.py`

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References:

[14] Xin Dang, Hanxiang Peng, Xueqin Wang and Heping Zhang: Theil-Sen Estimators in a Multiple Linear Regression Model.

[15] T. KÃ¤rkkÃ¤inen and S. Ã„yrÃ¤mÃ¶: On Computation of Spatial Median for Robust Data Mining.

Huber Regression

The :class:`HuberRegressor` is different to :class:`Ridge` because it applies a linear loss to samples that are classified as outliers. A sample is classified as an inlier if the absolute error of that sample is lesser than a certain threshold. It differs from :class:`TheilSenRegressor` and :class:`RANSACRegressor` because it does not ignore the effect of the outliers but gives a lesser weight to them.

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The loss function that :class:`HuberRegressor` minimizes is given by

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$$\min_{w, \sigma} {\sum_{i=1}^n\left(\sigma + H_{\epsilon}\left(\frac{X_{i}w - y_{i}}{\sigma}\right)\sigma\right) + \alpha {||w||_2}^2}$$

where

$$H_{\epsilon}(z) = \begin{cases} z^2, & \text {if } |z| < \epsilon, \\ 2\epsilon|z| - \epsilon^2, & \text{otherwise} \end{cases}$$

It is advised to set the parameter `epsilon` to 1.35 to achieve 95% statistical efficiency.

Notes

The :class:`HuberRegressor` differs from using :class:`SGDRegressor` with loss set to *huber* in the following ways.

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- :class:`HuberRegressor` is scaling invariant. Once `epsilon` is set, scaling `X` and `y` down or up by different values would produce the same robustness to outliers as before. as compared to :class:`SGDRegressor` where `epsilon` has to be set again when `X` and `y` are scaled.

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- :class:`HuberRegressor` should be more efficient to use on data with small number of samples while :class:`SGDRegressor` needs a number of passes on the training data to produce the same robustness.

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Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_huber_vs_ridge.py`

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References:

- Peter J. Huber, Elvezio M. Ronchetti: Robust Statistics, Concomitant scale estimates, pg 172

Note that this estimator is different from the R implementation of Robust Regression (http://www.ats.ucla.edu/stat/r/dae/rreg.htm) because the R implementation does a weighted least squares implementation with weights given to each sample on the basis of how much the residual is greater than a certain threshold.

Quantile Regression

Quantile regression estimates the median or other quantiles of y conditional on X, while ordinary least squares (OLS) estimates the conditional mean.

As a linear model, the :class:`QuantileRegressor` gives linear predictions $\hat{y}(w, X) = Xw$ for the q-th quantile, $q \in (0, 1)$. The weights or coefficients w are then found by the following minimization problem:

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$$\min_w \frac{1}{n_{\text{samples}}} \sum_i PB_q(y_i - X_i w) + \alpha \|w\|_1.$$

This consists of the pinball loss (also known as linear loss), see also :class:`~sklearn.metrics.mean_pinball_loss`,

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$$PB_q(t) = q\max(t, 0) + (1 - q)\max(-t, 0) = \begin{cases} qt, & t > 0, \\ 0, & t = 0, \\ (1 - q)t, & t < 0 \end{cases}$$

and the L1 penalty controlled by parameter `alpha`, similar to :class:`Lasso`.

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As the pinball loss is only linear in the residuals, quantile regression is much more robust to outliers than squared error based estimation of the mean. Somewhat in between is the :class:`HuberRegressor`.

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Quantile regression may be useful if one is interested in predicting an interval instead of point prediction. Sometimes, prediction intervals are calculated based on the assumption that prediction error is distributed normally with zero mean and constant variance. Quantile regression provides sensible prediction intervals even for errors with non-constant (but predictable) variance or non-normal distribution.

Based on minimizing the pinball loss, conditional quantiles can also be estimated by models other than linear models. For example, :class:`~sklearn.ensemble.GradientBoostingRegressor` can predict conditional quantiles if its parameter `loss` is set to `"quantile"` and parameter `alpha` is set to the quantile that should be predicted. See the example in :ref:`sphx_glr_auto_examples_ensemble_plot_gradient_boosting_quantile.py`.

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Most implementations of quantile regression are based on linear programming problem. The current implementation is based on :func:`scipy.optimize.linprog`.

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Examples:

- :ref:`sphx_glr_auto_examples_linear_model_plot_quantile_regression.py`

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References:

- Koenker, R., & Bassett Jr, G. (1978). Regression quantiles. Econometrica: journal of the Econometric Society, 33-50.
- Portnoy, S., & Koenker, R. (1997). :doi:`The Gaussian hare and the Laplacian tortoise: computability of squared-error versus absolute-error estimators. Statistical Science, 12, 279-300 <10.1214/ss/1030037960>`.

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- Koenker, R. (2005). :doi:`Quantile Regression <10.1017/CBO9780511754098>`. Cambridge University Press.

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Polynomial regression: extending linear models with basis functions

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> ```
> .. currentmodule:: sklearn.preprocessing
> ```

One common pattern within machine learning is to use linear models trained on nonlinear functions of the data. This approach maintains the generally fast performance of linear methods, while allowing them to fit a much wider range of data.

For example, a simple linear regression can be extended by constructing **polynomial features** from the coefficients. In the standard linear regression case, you might have a model that looks like this for two-dimensional data:

$$\hat{y}(w, x) = w_0 + w_1 x_1 + w_2 x_2$$

If we want to fit a paraboloid to the data instead of a plane, we can combine the features in second-order polynomials, so that the model looks like this:

$$\hat{y}(w, x) = w_0 + w_1 x_1 + w_2 x_2 + w_3 x_1 x_2 + w_4 x_1^2 + w_5 x_2^2$$

The (sometimes surprising) observation is that this is *still a linear model*: to see this, imagine creating a new set of features

$$z = [x_1, x_2, x_1 x_2, x_1^2, x_2^2]$$

With this re-labeling of the data, our problem can be written

$$\hat{y}(w, z) = w_0 + w_1 z_1 + w_2 z_2 + w_3 z_3 + w_4 z_4 + w_5 z_5$$

We see that the resulting *polynomial regression* is in the same class of linear models we considered above (i.e. the model is linear in w) and can be solved by the same techniques. By considering linear fits within a higher-dimensional space built with these basis functions, the model has the flexibility to fit a much broader range of data.

Here is an example of applying this idea to one-dimensional data, using polynomial features of varying degrees:

This figure is created using the :class:`PolynomialFeatures` transformer, which transforms an input data matrix into a new data matrix of a given degree. It can be used as follows:

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```
>>> from sklearn.preprocessing import PolynomialFeatures
>>> import numpy as np
>>> X = np.arange(6).reshape(3, 2)
>>> X
array([[0, 1],
       [2, 3],
       [4, 5]])
>>> poly = PolynomialFeatures(degree=2)
>>> poly.fit_transform(X)
array([[ 1.,  0.,  1.,  0.,  0.,  1.],
       [ 1.,  2.,  3.,  4.,  6.,  9.],
       [ 1.,  4.,  5., 16., 20., 25.]])
```

The features of X have been transformed from $[x_1, x_2]$ to $[1, x_1, x_2, x_1^2, x_1 x_2, x_2^2]$, and can now be used within any linear model.

This sort of preprocessing can be streamlined with the :ref:`Pipeline <pipeline>` tools. A single object representing a simple polynomial regression can be created and used as follows:

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```
>>> from sklearn.preprocessing import PolynomialFeatures
>>> from sklearn.linear_model import LinearRegression
>>> from sklearn.pipeline import Pipeline
>>> import numpy as np
>>> model = Pipeline([('poly', PolynomialFeatures(degree=3)),
...                   ('linear', LinearRegression(fit_intercept=False))])
>>> # fit to an order-3 polynomial data
>>> x = np.arange(5)
>>> y = 3 - 2 * x + x ** 2 - x ** 3
>>> model = model.fit(x[:, np.newaxis], y)
>>> model.named_steps['linear'].coef_
array([ 3., -2.,  1., -1.])
```

The linear model trained on polynomial features is able to exactly recover the input polynomial coefficients.

In some cases it's not necessary to include higher powers of any single feature, but only the so-called *interaction features* that multiply together at most d distinct features. These can be gotten from :class:`PolynomialFeatures` with the setting `interaction_only=True`.

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For example, when dealing with boolean features, $x_i^n = x_i$ for all n and is therefore useless; but $x_i x_j$ represents the conjunction of two booleans. This way, we can solve the XOR problem with a linear classifier:

```
>>> from sklearn.linear_model import Perceptron
>>> from sklearn.preprocessing import PolynomialFeatures
>>> import numpy as np
>>> X = np.array([[0, 0], [0, 1], [1, 0], [1, 1]])
>>> y = X[:, 0] ^ X[:, 1]
>>> y
array([0, 1, 1, 0])
>>> X = PolynomialFeatures(interaction_only=True).fit_transform(X).astype(int)
>>> X
array([[1, 0, 0, 0],
       [1, 0, 1, 0],
       [1, 1, 0, 0],
       [1, 1, 1, 1]])
>>> clf = Perceptron(fit_intercept=False, max_iter=10, tol=None,
...                  shuffle=False).fit(X, y)
```

And the classifier "predictions" are perfect:

```
>>> clf.predict(X)
array([0, 1, 1, 0])
>>> clf.score(X, y)
1.0
```