Feature Selection - Feature Extraction

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Why?

Given some data, we often wish to perform preprocessing in order to:

- transform it to a format that our algorithms can take as input
- reduce time complexity: Less computation
- reduce space complexity: Less parameters
- make it have properties (0-mean, unit variance, sparseness,...)
- More interpretable: simpler explanation
- Data visualization (structure, groups, outliers, etc)

Curse of dimensionality

Assume n data lives in $[0, 1]^d$.

To capture a cube neighborhood representing a fraction s of $[0,1]^d$, need the edge length to be $l=s^{1/d}.$

- d=10, s=0.1 \Rightarrow l=0.63
- d=10, s=0.01 \Rightarrow l=0.8

 \Rightarrow Neighborhoods are no longer local

The volume V of an hypercube with an edge length l= 0.1 is V = 0.1^d

$$\Rightarrow d
earrow$$
 , $V \searrow$ 0

 \Rightarrow Probability to capture points becomes pprox 0



Curse of dimensionality

Points in high dimensional spaces are isolated Immustration:

- Uniform sample n point in $[0,1]^d$
- Compute the volume occupied by the k nearest neighbors of a point: $l^d pprox k/n$



Feature selection / extraction

- Feature selection
 - \circ Choosing k < d important features, ignoring the remaining d - k \Rightarrow Subset selection algorithmss
- Feature extraction
 - Project the original d dimensions to **new** k < d dimensions \Rightarrow Discover low dimensional representations (smooth manifold) for data in high dimension. **Manifold Learning**



Implementation issues - scikit-learn

Install User Guide API Examples Community 🗗 More 🔻

Q D 1.5.1 (stable) -

scikit-learn

Machine Learning in Python

Getting Started Release Highlights for 1.5

Classification

learn

Identifying which category an object belongs to.

Applications: Spam detection, image recognition. Algorithms: Gradient boosting, nearest neighbors, random forest, logistic regression, and more...



Examples

- Simple and efficient tools for predictive data analysis
- Accessible to everybody, and reusable in various contexts
- Built on NumPy, SciPy, and matplotlib
- Open source, commercially usable BSD license

Regression

Predicting a continuous-valued attribute associated with an object.

Applications: Drug response, stock prices. Algorithms: Gradient boosting, nearest neighbors, random forest, ridge, and more...



Examples

Clustering

Automatic grouping of similar objects into sets.

Applications: Customer segmentation, grouping experiment outcomes. Algorithms: <u>k-Means</u>, <u>HDBSCAN</u>, <u>hierarchical</u> clustering, and more...



Examples

Feature selection - Filter

• Variance threshold

-Compute the variance of each feature

- Assume that features with a higher variance may contain more useful information
- Select the subset of features based on a user-specified threshold

늘 Good: fast!

Does not take the relationship among features into account

from sklearn.feature_selection import VarianceThreshold

Feature selection - Filter

- Information gain
 - $\,\circ\,$ Measures the amount of information that feature f provides on dataset Z
 - Features with high information gain are more important
 - Based on the entropy

IG(Z, f) = H(Z) - H(Z|f)

H(Z): entropy of the class distribution in Z H(Z|f): conditional entropy of the class distribution in Z given f



Feature selection - Wrappers

Recursive Feature Elimination (RFE)

- Fit model to the training set (the model has to provide information about feature importance)
- Eliminate feature with the smallest coefficient
- Repeat until k features are reached

```
from sklearn.feature_selection import RFE
from sklearn.datasets import load_iris
from sklearn.tree import DecisionTreeClassifier
X, y = load_iris(return_X_y=True)
estimator = DecisionTreeClassifier()
s = RFE(estimator, n_features_to_select=2, step=1)
s.fit(X, y)
print("Taille des données avant sélection",X.shape)
print("Variables sélectionnées : ", s.get_support())
print("Classement des variables : ",s.ranking_)
Taille des données avant sélection (150, 4)
Variables sélectionnées : [False False True True]
Classement des variables : [2 3 1 1]
```

Feature selection - Wrappers

Permutation Importance

For each feature, do n times

- Shuffle feature column
- Assess performance w.r.t. original

```
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.inspection import permutation_importance
from sklearn.cluster import KMeans
data = load_iris()
X = data.data
y = data.target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=0)
model = KMeans(n_clusters=2).fit(X_train, y_train)
r = permutation_importance(model, X_test, y_test,n_repeats=30,random_state=0)
for i in r.importances_mean.argsort()[::-1]:
         print(f"{data.feature_names[i]:<8}"</pre>
               f"{r.importances_mean[i]:.3f}"
               f" +/- {r.importances_std[i]:.3f}")
petal length (cm)77.803 +/- 12.963
petal width (cm)39.936 +/- 6.399
sepal length (cm)30.218 +/- 6.479
sepal width (cm)5.483 +/- 1.083
```

Feature selection - Wrappers

Shapley values

Algorithm 6 (Estimation de la valeur de Shapley du descripteur i)

Entrée : Nombre d'itérations M, exemple \mathbf{x} , ensemble des exemples \mathbf{X} , i, modèle f

Sortie : Estimation de la valeur de Shapley du descripteur *i*

1. Pour tout $j \in \llbracket 1, M
rbracket$

1. Tirer un exemple ${f z}$ dans ${f X}$

2. Tirer une permutation aléatoire σ de l'ensemble $\{1 \cdots d\}$

3.
$$\mathbf{x}_{\sigma}=(x_{\sigma(1)}\cdots x_{\sigma(d)})$$
 et $\mathbf{z}_{\sigma}=(z_{\sigma(1)}\cdots z_{\sigma(d)})$

4. Créer deux nouveaux exemples :

1. $\mathbf{x}_{+i} = (x_{\sigma(1)} \cdots x_{\sigma(i-1)}, x_{\sigma(i)}; z_{\sigma(i+1)} \cdots z_{\sigma(d)})$ 2. $\mathbf{x}_{-i} = (x_{\sigma(1)} \cdots x_{\sigma(i-1)}, z_{\sigma(i)}; z_{\sigma(i+1)} \cdots z_{\sigma(d)})$ 5. Calculer la contribution marginale du descripteur $i : \phi_i^j = f(\mathbf{x}_{+i}) - f(\mathbf{x}_{-i})$ 2. Calculer un estimateur de la valeur de Shapley du descripteur $i : \phi_i(\mathbf{x}) = \frac{1}{M} \sum_{i=1}^M \phi_i^j$

Feature selection- Wrappers

Shapley values



Feature selection - Embedded

LASSO regularization

Least Absolute Shrinkage and Selection Operator

 $Minf_w(x)+\lambda\|w\|_1$

from sklearn.linear_model import LogisticRegression
model = LogisticRegression(penalty='l1')

Feature selection: Filters/Embedded methods

- Filters measure the relevance of features by their correlation with the dependent variable, while embedded methods measure the usefulness of a subset of features by training a model on them.
- Filters are much faster than enveloping methods
- Filters use statistical methods to evaluate a subset of features, while embedded methods use cross-validation.
- Filters may fail to find the best subset of features on many occasions, but enveloping methods can always provide the best subset of features.

Feature extraction

- Linear methods: PCA, MDS, LDA
- Non linear methods: ISOMAP, LLE, t-SNE

One standard method for decoupling and dimensionality reduction of continuous data is Principal Component Analysis (PCA)

- Find a low-dimensional space such that when x is projected there, information loss is minimized.
- The projection of x on the direction of w is: $z = w^T x$
- Find w such that $\mathbb{V}(z)$ is maximized

$$egin{aligned} V(z) &= \mathbb{V}(w^Tx) = \mathbb{E}\left((w^Tx-w^T\mu)^2
ight) \ &= \mathbb{E}\left((w^Tx-w^T\mu)(w^Tx-w^T\mu)
ight) \ &= \mathbb{E}\left(w^T(x-\mu)(x-\mu)^Tw
ight) \ &= w^T\mathbb{E}\left((x-\mu)(x-\mu)^T
ight) w \ &= w^T\Sigma w \end{aligned}$$

• First PC*

Maximize $\mathbb{V}(z)$ subject to $||w|| = 1 \Rightarrow \max_{u} u^T \Sigma u - \alpha(u^T u - 1)$ $\Sigma u = \alpha u \Rightarrow u$ eigenvector of Σ Choose u with the largest eigenvalue for $\mathbb{V}(z)$ to be maximized

Second PC*

Deflation or $\mathbb{V}(z_2)$ subject to $\|w\|=1$ and orthogonal to u

$$\max_{w} w^T \Sigma w - lpha (w^T w - 1) - eta (w^T u)$$

 $\Sigma w = lpha w \Rightarrow w$ eigenvector of Σ

• continue k times*

What PCA does

$$z = W^T(x - m)$$

where $W_{.,j}$ is the j^{th} eigenvector of Σ , and m is the sample mean. 22 x_2 Z_1 Z x_1 Z_1

POV: proportion of variance.



- Typically, k/ POV>threshold
- Scree graph plot of POV, stop at elbow



```
from sklearn.decomposition import PCA
X = ...
pca = PCA(n_components=2)
pca.fit(X)
PCA(n_components=2)
print(pca.explained_variance_ratio_)
```

Given pairwise distances between N points, d_{ij} , $i, j \in \{1 \cdots N\}$, place on a low dimensional map such as distances are preserved.

Sammon stress

$$z = g(x| heta)$$

Find θ minimizing

$$\mathbb{E}(heta|X) = \sum_{r,s} rac{\left(\|z^r-z^s\|-\|x^r-x^s\|
ight)^2}{\|x^r-x^s\|^2}$$

Feature extraction - MDS

from sklearn.manifold import MDS
X = ..
embedding = MDS(n_components=2)
X2 = embedding.fit_transform(X)





Example: 3D points of swiss roll given by their distance matrix.

Linear (or Fisher) Discriminant Analysis Find a low-dimensional space such that when xis projected, classes are well-separated.





Data Scattering

$$\max_{w} rac{m_1 - m_2}{s_1^2 + s_2^2}$$

Inter class

$$(m_1-m_2)^2 = ig(w^T\mu_1-w^T\mu_2ig)^2 \ = w^T(\mu_1-\mu_2)(\mu_1-\mu_2)^Tw = w^TS_Bw$$
 where $S_B = (\mu_1-\mu_2)(\mu_1-\mu_2)^T$

Data Scattering

$$\max_{w} rac{m_1 - m_2}{s_1^2 + s_2^2}$$

Intra class

$$s_1^2 = \sum_t r_t ig(w^T x_t - m_1 ig)^2 \ = \sum_t r_t w^T (x_t - m_1) (x_t - m_1)^T w = w^T S_1 w$$
 where $S_1 = \sum_t r_t (x_t - m_1) (x_t - m_1)^T$ $s_1^2 + s_2^2 = w^T S_W w$, $S_W = S_1 + S_2$

Fisher's discriminant

Find w maximizing

 $rac{w^T S_B w}{w^T S_W w}$

Solutions

• LDA:
$$w=c.\,S_W^{-1}(m_1-m_2)$$

• Parametric: $w = \Sigma^{-1}(\mu_1 - \mu_2)$, when $p(x|C_i) pprox \mathcal{N}(\mu_i, \Sigma)$

from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
X,y = ..
lda = LinearDiscriminantAnalysis(n_components=2)
X2 = lda.fit(X, y).transform(X)

Fisher's discriminant

What about the multiple class case (C>2) ?

• Inter class:
$$S_B = \sum_{i=1}^C N_i (\mu_i - \mu) (\mu_i - \mu)^T$$
 $m = \frac{1}{C} \sum_{i=1}^C \mu_i$
• Intra class: $S_W = \sum_{i=1}^C S_i = \sum_{i=1}^C r_{t,i} (x_t - m_i) (x_t - m_i)^T$

$$\max_w rac{W^T S_B W}{W^T S_W W}$$

- ightarrow The largest eigenvectors of $S_W^{-1}S_B$
- ightarrow Maximum rank of C-1

Non Linear Feature extraction - Introduction

Deficiencies of Linear Methods

Data may not be best summarized by linear combination of features.

Example: PCA cannot discover 1D structure of a helix



Seminal Paper: J. B. Tenenbaum, V. de Silva and J. C. Langford, A Global Geometric Framework for Nonlinear Dimensionality Reduction, Science 290 (5500) 2000

Algorithm

- 1. Constructing neighbourhood graph ${\cal G}$
- 2. \forall pair of points in *G*: shortest path distances \approx geodesic distances.
- 3. Use MDS with geodesic distances.

- Construction of the neighbourhood graph G (K-nearest neighborhood (K=7)). D_G : 1000×1000 (Euclidean) distance matrix (fig A)
- Shortest paths in $G: D_G: 1000 \times 1000$ geodesic distance matrix of two arbitrary points along the manifold (fig B)
- Embedding G in \mathbb{R}^d using MDS: Find a d-D Euclidean space preserving pairwise distances (fig C)



Example: unfolding the swiss roll

Example: Embedding MNIST data (\mathbb{R}^{784}) into \mathbb{R}^2





Advantages

- Nonlinear
- Globally optimal low-dimensional Euclidean representation even though input space is highly folded, twisted, or curved.
- Guarantee asymptotically to recover the true dimensionality.

Disadvantages

- May not be stable, depends on the topology of the manifold
- asymptotically recover geometric structure of nonlinear manifolds
 - $\circ~N$ high: pairwise distances pprox geodesics, but costly
 - $\circ~N$ small: geodesic distances very inaccurate
- Distance matrix is dense \Rightarrow does not scale to large datasets \rightarrow Landmark Isomap

from sklearn.manifold import Isomap
X = ...
iso = Isomap(n_components=2)
X2= iso.fit_transform(X)

Seminal Paper} Sam T. Roweis and Lawrence K. Saul, Nonlinear Dimensionality Reduction by Locally Linear Embedding, Science 22:Vol. 290, 2000

ISOMAP vs. LLE

• Local Linear Embedding \Rightarrow local approach

 \Rightarrow The resulting matrix is sparse...Apply efficient sparse matrix solvers

Characterictics of a Manifold

- Locally \boldsymbol{M} is a linear patch
- How to combine all local patches together?



Algorithm

Assumption: manifold ${\cal M}$ is roughly linear when viewed locally

Approximation error can be made small:

$$Min_W \|x_i - \sum_{j=1}^k w_{ij} x_j\|^2$$
 .

- 1. W: a linear representation of every data point by its neighbors. This is an intrinsic geometrical property of the manifold
- 2. A good projection should preserve this local geometric property as much as possible

Algorithm

- We expect each data point and its neighbors to lie on or close to a locally linear patch of \mathcal{M} .
- Each point can be written as a linear combination of its neighbors. The weights chosen to minimize the reconstruction error.



Optimal weights: weights that minimize the reconstruction errors are invariant to rotation, rescaling and translation of the data points.

- Invariance to translation is enforced by adding the constraint that the weights sum to one.
- The weights characterize the intrinsic geometric properties of each neighborhood.

Local geometry is preserved: the same weights that reconstruct the data points in D dimensions should reconstruct it in the manifold in d dimensions.

Algorithm

Low-dimensional embedding $Y\in \mathcal{M}_{d,N}(\mathbb{R})$

$$\min_{Y} \sum_{i=1}^{N} \|Y_{.,i} - YW_{i,.}\|^2$$

Use the same weights from the original space



LLE - Constrained LS Problem

Optimization: Compute the optimal weight for each point individually:

$$\|x_i - \sum_{j=1}^k w_{ij} x_j \|^2 = \|\sum_{j=1}^k w_{ij} (x_i - x_j) \|^2 = \sum_{j=1}^k \sum_k w_{ij} w_{ik} C_{jk}$$

where $C_{jk} = (x_i - x_j)^T (x_i - x_k)$ \ Can be minimized using a Lagrange multiplier for $\sum_j w_{ij} = 1$

Solution:

$$w_{ij} = rac{\displaystyle \sum_k C_{jk}^{-1}}{\displaystyle \sum_{lm} C_{lm}^{-1}}$$

LLE space

- $Y_{.,i} \in \mathbb{R}^k$: projected vector for X_i
- The geometrical property is best preserved if $E(Y) = \sum_i \|Y_{.,i} \sum_j w_{ij}Y_{.,j}\|^2$ is small
- Y: eigenvectors of the lowest d non-zero eigenvalues of $M = (I-W)^T (I-W)$
- ightarrow Eigenvalue problem}: $E(Y) = Tr(YMY^T)$

 $U = (U_1 \cdots U_d)$: bottom eigenvectors of M. Then

$$Y = U^T ext{ and } M_{ij} = \delta_{ij} - w_{ij} - w_{ji} + \sum_k w_{ki} w_{kj}$$

from sklearn.manifold import LocallyLinearEmbedding
X = ...
lle = LocallyLinearEmbedding(n_components=2)
X2 = lle.fit_transform(X)

Example: 3D points of swiss roll given by their distance matrix.

Example: Embedding MNIST data (\mathbb{R}^{784}) into \mathbb{R}^2





Limitations

- Require dense data points on the manifold for good estimation
- Need for a good neighborhood \Rightarrow How to choose k?
 - $\circ~\mbox{small} \rightarrow \mbox{rank}$ deficient tangent space and lead to over-fitting
 - $\circ~$ large \rightarrow Tangent space will not match local geometry well



t-Distributed Stochastic Neighbor Embedding: a 4 step algorithm

1. Transform the distance information between data points x_i into conditional probabilities, expressing a similarity relationship. The similarity between x_i and x_j is the conditional probability $P_{j|i}$, which expresses the fact that x_i will consider x_j as its neighbor according to a Gaussian probability density centered at x_i and of variance σ_i .

$$P_{j|i} = rac{e^{-rac{\|x_i-x_j\|^2}{2\sigma_i^2}}}{\displaystyle{\sum_{k
eq i}e^{-rac{\|x_i-x_k\|^2}{2\sigma_i^2}}}}$$

t-Distributed Stochastic Neighbor Embedding: a 4 step algorithm

2. In the same way, we calculate on ${\cal M}$

$$Q_{j|i} = rac{e^{-rac{\|y_i-y_j\|^2}{2\sigma_i^2}}}{\displaystyle{\sum_{k
eq i}e^{-rac{\|y_i-y_k\|^2}{2\sigma_i^2}}}}$$

By minimizing the difference between $P_{j|i}$ and $Q_{j|i}$, the Stochastic Neighbor Embedding (SNE) algorithm attempts to preserve similarities. SNE minimizes the sum of Kullback-Leibler divergences by gradient descent:

$$D_{KL}(P,Q) = \sum_{i
eq j} P_{j|i} log\left(rac{P_{j|i}}{Q_{j|i}}
ight)$$

t-Distributed Stochastic Neighbor Embedding: a 4 step algorithm

3. tSNE uses other definitions of conditional probabilities in the original space and \mathcal{M} .

$$P_{ij}=rac{P_{j|i}+P_{i|j}}{2n}$$

with $P_{ii} = 0$ for all i, and

$$Q_{ij} = rac{\left(1 + \|y_i - y_j\|^2
ight)^{-1}}{\displaystyle\sum_{k
eq l} \left(1 + \|y_l - y_k\|^2
ight)^{-1}}$$

minimization being performed on the Kullback Leibler divergence. As this cost function is non-symmetrical, optimization favors modeling high values of P_{ij} by high values of Q_{ij} .

$$D_{KL}(P,Q) = \sum_{i
eq j} P_{ij} log\left(rac{P_{ij}}{Q_{ij}}
ight)$$

t-Distributed Stochastic Neighbor Embedding: a 4 step algorithm

4. the variance depends on the observation point. Each value of σ_i induces a probability distribution P_i over the set of x_i points, whose entropy $H(P_i)$ increases with σ_i .

tSNE performs a binary search for a value of σ_i allowing P_i to reach a fixed perplexity $Perp(P_i) = 2^{H(P_i)}$, with $H(P_i) = -\sum_i P_{ij} log_2 P_{ij}$.

Perplexity can be interpreted as a measure of the average number of effective neighbors.

Example: Embedding MNIST data (\mathbb{R}^{784}) into \mathbb{R}^2