SVMs

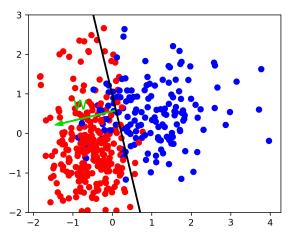
Presented by Yasin Ceran

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Linear Models for Binary Classification

2 Kernel SVMs

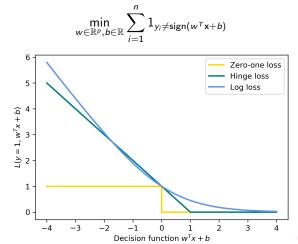
Linear Binary Classification



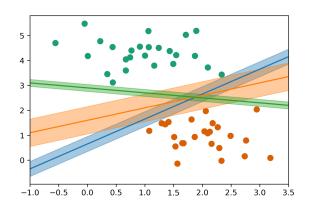
$$\hat{y} = \operatorname{sign}(w^T \mathbf{x} + b) = \operatorname{sign}\left(\sum_i w_i x_i + b\right)$$

Picking A Loss

$$\hat{y} = \operatorname{sign}(w^T \mathbf{x} + b)$$



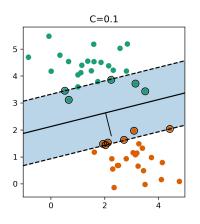
Max-Margin and Support Vectors (1)

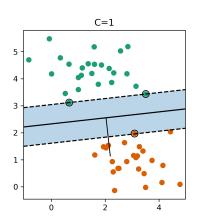


- $\min_{w \in \mathbb{R}^p, b \in \mathbb{R}} C \sum_{i=1}^n \max(0, 1 y_i(w^T \mathbf{x} + b)) + ||w||_2^2$
- Within margin $\Leftrightarrow y_i(w^Tx + b) < 1$
- Smaller $w \Rightarrow$ larger margin

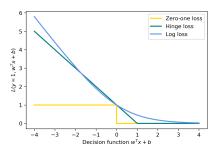


Max-Margin and Support Vectors (2)





Logistic Regression vs SVM



- $\min_{w \in P, b \in \mathbb{R}} C \sum_{i=1}^{n} \log(\exp(-y_i(w^T \mathbf{x}_i + b)) + 1) + ||w||_2^2$
- $\min_{w \in P, b \in \mathbb{R}} C \sum_{i=1}^{n} \max(0, 1 y_i(w^T \mathbf{x}_i + b)) + ||w||_2^2$
- Do you need probability estimates?
 - If yes, use Logistic Regression
 - If it doesn't matter, try either/both
- Need compact model or believe solution is sparse, use L_1 .

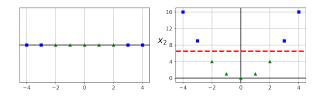
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1 Linear Models for Binary Classification

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Motivation

- Go from linear models to more powerful nonlinear ones.
- Keep convexity (ease of optimization).
- Generalize the concept of feature engineering.



Reminder on Linear SVM

$$\min_{w \in \mathbb{R}^p, b \in \mathbf{R}} C \sum_{i=1}^n \max(0, 1 - y_i(w^T \mathbf{x} + b)) + ||w||_2^2$$

$$\hat{y} = \operatorname{sign}(w^T \mathbf{x} + b)$$

```
#####POLYNOMIAL FEATURES#####
    from sklearn.datasets import

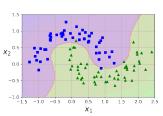
    make moons

 3 from sklearn.pipeline import
    → Pipeline
 4 from sklearn.preprocessing import

→ PolynomialFeatures.StandardScaler

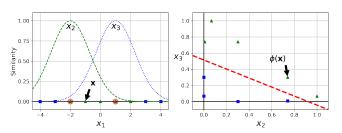
    from sklearn.svm import LinearSVC
 7
     polynomial_svm_clf = Pipeline([
            ("poly_features",
           → PolynomialFeatures(degree=3)).
            ("scaler",
 9

→ StandardScaler()),
10
            ("svm clf", LinearSVC(C=10.
           → loss="hinge",
           11
        1)
12
13
     polynomial sym clf.fit(X, v)
```



Similarity Features

- You can use a similarity function to measure how much each instance resembles a particular landmark
- Gaussian Radial Basis Function (RBF): $\phi_{\gamma}(\mathbf{x}, I) = \exp(\gamma ||\mathbf{x} I||^2)$



Reformulate Linear Models

Optimization Theory

$$w = \sum_{i=1}^{n} \alpha_i \mathbf{x}_i$$

(alpha are dual coefficients. Non-zero for support vectors only)

$$\hat{y} = \operatorname{sign}(w^T \mathbf{x}) \Longrightarrow \hat{y} = \operatorname{sign}\left(\sum_{i=1}^{n} \alpha_i(\mathbf{x}_i^T \mathbf{x})\right)$$

$$\alpha_i <= C$$

Introducing Kernels

$$\hat{y} = \operatorname{sign}\left(\sum_{i}^{n} \alpha_{i}(\mathbf{x}_{i}^{T}\mathbf{x})\right) \longrightarrow \hat{y} = \operatorname{sign}\left(\sum_{i}^{n} \alpha_{i}(\phi(\mathbf{x}_{i})^{T}\phi(\mathbf{x}))\right)$$
$$\phi(\mathbf{x}_{i})^{T}\phi(\mathbf{x}_{j}) \longrightarrow k(\mathbf{x}_{i}, \mathbf{x}_{j})$$

Examples of Kernels

$$k_{\mathsf{linear}}(\mathbf{x}, \mathbf{x}') = \mathbf{x}^T \mathbf{x}'$$

$$k_{\mathsf{poly}}(\mathbf{x},\mathbf{x}') = (\mathbf{x}^T\mathbf{x}' + c)^d$$

$$k_{\mathsf{rbf}}(\mathbf{x}, \mathbf{x}') = \exp(-\gamma ||\mathbf{x} - \mathbf{x}'||^2)$$

$$k_{\text{sigmoid}}(\mathbf{x}, \mathbf{x}') = \tanh\left(\gamma \mathbf{x}^T \mathbf{x}' + r\right)$$

$$k_{\cap}(\mathbf{x},\mathbf{x}') = \sum_{i=1}^{p} \min(x_i,x_i')$$

- If k and k' are kernels, so are k + k', kk', ck', ...

Polynomial Kernel vs Features

$$k_{\mathsf{poly}}(\mathbf{x}, \mathbf{x}') = (\mathbf{x}^T \mathbf{x}' + c)^d$$

Complexity:

Explicit polynomials $\rightarrow n$ samples *n features^d

 $\mathsf{Kernel\ trick} \to n_\mathit{samples} * n_\mathit{samples} * n_\mathit{features}$

For a single feature:

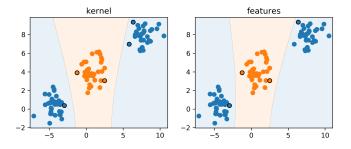
$$(x^2, \sqrt{2}x, 1)^T(x'^2, \sqrt{2}x', 1) = x^2x'^2 + 2xx' + 1 = (xx' + 1)^2$$

Poly kernels vs explicit features

```
poly = PolynomialFeatures(include_bias=False)

X_poly = poly.fit_transform(X)
print(X.shape, X_poly.shape)
print(poly.get_feature_names())

((100, 2), (100, 5))
['x0', 'x1', 'x0'2', 'x0 x1', 'x1'2']
```



Understanding Dual Coefficients

$$y = sign(0.139x_0 + 0.06x_1 - 0.201x_0^2 + 0.048x_0x_1 + 0.019x_1^2)$$

```
1 linear_svm.coef_
2 array([[0.139, 0.06, -0.201, 0.048, 0.019]])
```

$$y = \operatorname{sign}(-0.03\phi(\mathbf{x}_1)^T \phi(x) - 0.003\phi(\mathbf{x}_{26})^T \phi(\mathbf{x}) + 0.003\phi(\mathbf{x}_{42})^T \phi(\mathbf{x}) + 0.03\phi(\mathbf{x}_{62})^T \phi(\mathbf{x}))$$

```
1 linear_svm.dual_coef_
2
3 array([[-0.03, -0.003, 0.003]])
4
5 linear_svm.support_
6
7 array([1,26,42,62], dtype=int32)
```

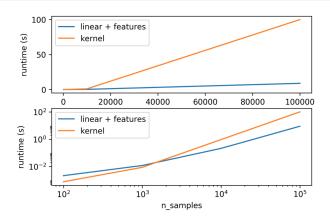
With Kernel

$$y = \operatorname{sign}\left(\sum_{i}^{n} \alpha_{i} k(\mathbf{x}_{i}, \mathbf{x})\right)$$

```
1 poly_svm.dual_coef_
2 array([[-0.057, -0., -0.012, 0.008, 0.062]])
4 poly_svm.support_
6 array([1,26,41,42,62], dtype=int32)
```

$$y = \operatorname{sign}(-0.057(\mathbf{x}_1^T \mathbf{x} + 1)^2 - 0.012(\mathbf{x}_{41}^T \mathbf{x} + 1)^2 + 0.008(\mathbf{x}_{42}^T \mathbf{x} + 1)^2 + 0.062(\mathbf{x}_{62}, \mathbf{x} + 1)^2)$$

Runtime Considerations



Kernels in Practice

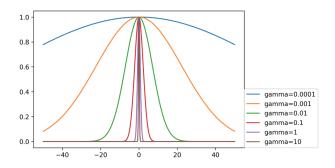
- Dual coefficients less interpretable
- Long runtime for "large" datasets (100k samples)
- As a rule of thumb, try linear kernel first (remember 'LinearSVC' is much faster than 'SVC(kernel=linear)'
- If the training data is not so large, try Gaussian RBF kernel

Preprocessing

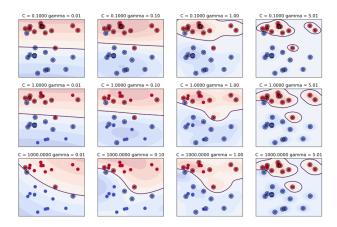
- Kernel use inner products or distances.
- StandardScaler or MinMaxScaler
- Gamma parameter in RBF directly relates to scaling of data and n_features - the default is 1/(X.var() * n_features)

Parameters for RBF Kernels

- Regularization parameter C is limit on alphas (for any kernel)
- Gamma is bandwidth: $k_{\rm rbf}(\mathbf{x}, \mathbf{x}') = \exp(-\gamma ||\mathbf{x} \mathbf{x}'||^2)$

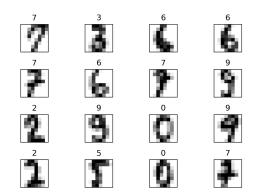


Parameters for RBF Kernels



MNIST Example

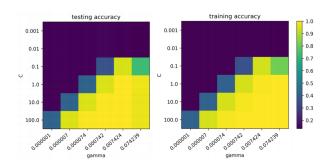
- 1 from sklearn.datasets import load_digits
- 3 digits = load_digits()



Scaling and Default Params

```
# gamma : f'scale', 'auto'} or float, optional (default='scale')
 2 # Kernel coefficient for 'rbf', 'poly' and 'sigmoid'.
    # if gamma='scale' (default) is passed then it uses 1 / (n_features * X.var())
    # as value of gamma
    # if 'auto', uses 1 / n_features.
    print('auto', np.mean(cross_val_score(SVC(gamma='auto'), X_train, v_train, cv=10)))
    print('scale', np.mean(cross_val_score(SVC(gamma='scale'), X_train, y_train, cv=10)))
    scaled_svc = make_pipeline(StandardScaler(), SVC())
10
     print('pipe', np.mean(cross_val_score(scaled_svc, X_train, y_train, cv=10)))
11
12
13
    auto 0.563
14
    scale 0.987
15
    pipe 0.977
16
17
    gamma = (1. / (X_train.shape[1] * X_train.var()))
     print(np.mean(cross_val_score(SVC(gamma=gamma), X_train, y_train, cv=10)))
18
19
20
    0.987
```

Grid-Searching Parameters



Summary

 Logistic Regression and Linear SVM differ from each other by their loss functions