#### Chapter 5 - Support Vector Machines

This notebook contains all the sample code and solutions to the exercises in chapter 5.



Run in Google Colab (https://colab.research.google.com/github/ageron/handson-ml2/blob/master/05\_support\_vector\_machines.ipynb)

#### Setup

First, let's import a few common modules, ensure MatplotLib plots figures inline and prepare a function to save the figures. We also check that Python 3.5 or later is installed (although Python 2.x may work, it is deprecated so we strongly recommend you use Python 3 instead), as well as Scikit-Learn ≥0.20.

```
In [1]: # Python ≥3.5 is required
        import sys
        assert sys.version_info >= (3, 5)
        # Scikit-Learn ≥0.20 is required
        import sklearn
        assert sklearn.__version__ >= "0.20"
        # Common imports
        import numpy as np
        import os
        # to make this notebook's output stable across runs
        np.random.seed(42)
        # To plot pretty figures
        %matplotlib inline
        import matplotlib as mpl
        import matplotlib.pyplot as plt
        mpl.rc('axes', labelsize=14)
mpl.rc('xtick', labelsize=12)
        mpl.rc('ytick', labelsize=12)
        # Where to save the figures
        PROJECT ROOT DIR = "."
        CHAPTER ID = "svm"
        IMAGES_PATH = os.path.join(PROJECT_ROOT_DIR, "images", CHAPTER_ID)
        os.makedirs(IMAGES PATH, exist ok=True)
        def save_fig(fig_id, tight_layout=True, fig_extension="png", resolution=300):
            path = os.path.join(IMAGES_PATH, fig_id + "." + fig_extension)
             print("Saving figure", fig id)
             if tight layout:
                 plt.tight_layout()
             plt.savefig(path, format=fig extension, dpi=resolution)
```

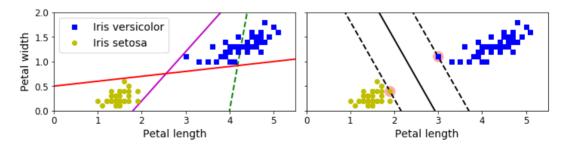
# Large margin classification

The next few code cells generate the first figures in chapter 5. The first actual code sample comes after:

```
In [2]: from sklearn.svm import SVC from sklearn import datasets
```

```
In [3]: # Bad models
           x0 = np.linspace(0, 5.5, 200)
           pred 1 = 5*x0 - 20
          pred^{-}2 = x0 - 1.8
          pred_3 = 0.1 * x0 + 0.5
           def plot_svc_decision_boundary(svm_clf, xmin, xmax):
                w = svm clf.coef [0]
                b = svm clf.intercept [0]
                # At the decision boundary, w0*x0 + w1*x1 + b = 0
                \# => x1 = -w0/w1 * x0 - b/w1
                x0 = np.linspace(xmin, xmax, 200)
                decision boundary = -w[0]/w[1] * x0 - b/w[1]
                margin = 1/w[1]
                gutter up = decision boundary + margin
                gutter down = decision boundary - margin
               svs = svm_clf.support_vectors_
plt.scatter(svs[:, 0], svs[:, 1], s=180, facecolors='#FFAAAA')
               plt.plot(x0, decision_boundary, "k-", linewidth=2)
plt.plot(x0, gutter_up, "k--", linewidth=2)
                plt.plot(x0, gutter_down, "k--", linewidth=2)
           fig, axes = plt.subplots(ncols=2, figsize=(10,2.7), sharey=True)
           plt.sca(axes[0])
          plt.sed(dxes[0])
plt.plot(x0, pred_1, "g--", linewidth=2)
plt.plot(x0, pred_2, "m-", linewidth=2)
plt.plot(x0, pred_3, "r-", linewidth=2)
          plt.plot(X[:, 0][y==1], X[:, 1][y==1], "bs", label="Iris versicolor")
plt.plot(X[:, 0][y==0], X[:, 1][y==0], "yo", label="Iris setosa")
          plt.xlabel("Petal length", fontsize=14)
plt.ylabel("Petal width", fontsize=14)
           plt.legend(loc="upper left", fontsize=14)
           plt.axis([0, 5.5, 0, 2])
           plt.sca(axes[1])
           plot svc decision boundary(svm clf, 0, 5.5)
          plt.plot(X[:, 0][y==1], X[:, 1][y==1], "bs")
plt.plot(X[:, 0][y==0], X[:, 1][y==0], "yo")
           plt.xlabel("Petal length", fontsize=14)
           plt.axis([0, 5.5, 0, 2])
           save_fig("large_margin_classification_plot")
           plt.show()
```

Saving figure large\_margin\_classification\_plot

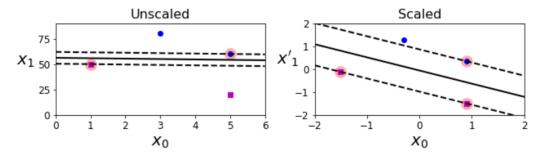


## Sensitivity to feature scales

```
In [4]: Xs = np.array([[1, 50], [5, 20], [3, 80], [5, 60]]).astype(np.float64)
    ys = np.array([0, 0, 1, 1])
    svm_clf = SVC(kernel="linear", C=100)
```

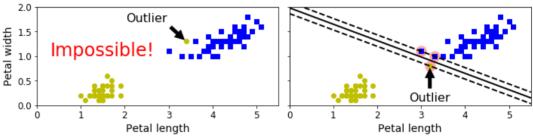
```
svm_clf.fit(Xs, ys)
plt.figure(figsize=(9,2.7))
plt.subplot(121)
plt.plot(Xs[:, 0][ys==1], Xs[:, 1][ys==1], "bo")
plt.plot(Xs[:, 0][ys==0], Xs[:, 1][ys==0], "ms")
plot_svc_decision_boundary(svm_clf, 0, 6)
plt.xlabel("$x_0$", fontsize=20)
plt.ylabel("$x_1$
                         ", fontsize=20, rotation=0)
plt.title("Unscaled", fontsize=16)
plt.axis([0, 6, 0, 90])
from sklearn.preprocessing import StandardScaler
scaler = StandardScaler()
X scaled = scaler.fit transform(Xs)
svm clf.fit(X scaled, ys)
plt.subplot(122)
plt.plot(X_scaled[:, 0][ys==1], X_scaled[:, 1][ys==1], "bo")
plt.plot(X_scaled[:, 0][ys==0], X_scaled[:, 1][ys==0], "ms")
plot svc decision boundary(svm_clf, -2, 2)
plt.xlabel("$x_0$", fontsize=20)
plt.ylabel("$x'_1$ ", fontsize=20, rotation=0)
plt.title("Scaled", fontsize=16)
plt.axis([-2, 2, -2, 2])
save_fig("sensitivity_to_feature_scales_plot")
```

Saving figure sensitivity\_to\_feature\_scales\_plot



## Sensitivity to outliers

```
arrowprops=dict(facecolor='black', shrink=0.1),
             fontsize=16,
plt.axis([0, 5.5, 0, 2])
plt.sca(axes[1])
plt.plot(Xo2[:, 0][yo2==1], Xo2[:, 1][yo2==1], "bs")
plt.plot(Xo2[:, 0][yo2==0], Xo2[:, 1][yo2==0], "yo")
plot_svc_decision_boundary(svm_clf2, 0, 5.5)
plt.xlabel("Petal length", fontsize=14)
plt.annotate("Outlier"
             xy=(X outliers[1][0], X outliers[1][1]),
             xytext=(3.2, 0.08),
             ha="center",
             arrowprops=dict(facecolor='black', shrink=0.1),
             fontsize=16,
plt.axis([0, 5.5, 0, 2])
save_fig("sensitivity_to_outliers_plot")
plt.show()
Saving figure sensitivity_to_outliers_plot
```



#### Large margin vs margin violations

This is the first code example in chapter 5:

```
In [6]: import numpy as np
        from sklearn import datasets
        from sklearn.pipeline import Pipeline
        from sklearn.preprocessing import StandardScaler
        from sklearn.svm import LinearSVC
        iris = datasets.load iris()
        X = iris["data"][:, (2, 3)] # petal length, petal width
        y = (iris["target"] == 2).astype(np.float64) # Iris virginica
        svm clf = Pipeline([
                ("scaler", StandardScaler()),
                ("linear svc", LinearSVC(C=1, loss="hinge", random state=42)),
            ])
        svm clf.fit(X, y)
Out[6]: Pipeline(memory=None,
             steps=[('scaler', StandardScaler(copy=True, with_mean=True, with_std=Tru
        e)), ('linear svc', LinearSVC(C=1, class weight=None, dual=True, fit intercept
             intercept_scaling=1, loss='hinge', max_iter=1000, multi_class='ovr',
             penalty='l2', random_state=42, tol=0.0001, verbose=0))])
```

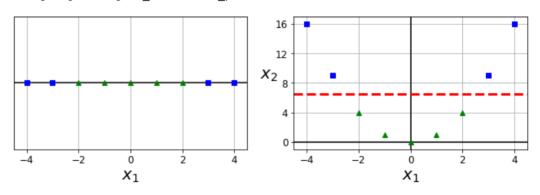
```
In [7]: svm clf.predict([[5.5, 1.7]])
 Out[7]: array([1.])
          Now let's generate the graph comparing different regularization settings:
          scaler = StandardScaler()
          svm clf1 = LinearSVC(C=1, loss="hinge", random state=42)
          svm clf2 = LinearSVC(C=100, loss="hinge", random_state=42)
          scaled_svm_clf1 = Pipeline([
                    ("scaler", scaler),
                    ("linear_svc", svm_clf1),
          scaled svm clf2 = Pipeline([
                    ("scaler", scaler),
                    ("linear_svc", svm_clf2),
               ])
          scaled svm clf1.fit(X, y)
          scaled svm clf2.fit(X, y)
          /Users/ageron/miniconda3/envs/tf2b/lib/python3.7/site-packages/sklearn/svm/bas
          e.py:931: ConvergenceWarning: Liblinear failed to converge, increase the numbe
          r of iterations.
             "the number of iterations.", ConvergenceWarning)
Out[8]: Pipeline(memory=None,
                steps=[('scaler', StandardScaler(copy=True, with_mean=True, with_std=Tru
          e)), ('linear_svc', LinearSVC(C=100, class_weight=None, dual=True, fit_interce
          pt=True.
                intercept scaling=1, loss='hinge', max iter=1000, multi class='ovr',
                penalty='l2', random_state=42, tol=0.0001, verbose=0))])
In [9]: # Convert to unscaled parameters
          b1 = svm_clf1.decision_function([-scaler.mean_ / scaler.scale_])
          b2 = svm_clf2.decision_function([-scaler.mean_ / scaler.scale_])
          w1 = svm_clf1.coef_[0] / scaler.scale_
          w2 = svm_clf2.coef_[0] / scaler.scale_
          svm_clf1.intercept_ = np.array([b1])
svm_clf2.intercept_ = np.array([b2])
          svm_clf1.coef_ = np.array([w1])
svm_clf2.coef_ = np.array([w2])
          # Find support vectors (LinearSVC does not do this automatically)
          t = y * 2 - 1
          support_vectors_idx1 = (t * (X.dot(w1) + b1) < 1).ravel()
          support_vectors_idx2 = (t * (X.dot(w2) + b2) < 1).ravel()
          svm_clf1.support_vectors = X[support_vectors_idx1]
svm_clf2.support_vectors = X[support_vectors_idx2]
In [10]: fig, axes = plt.subplots(ncols=2, figsize=(10,2.7), sharey=True)
          plt.sca(axes[0])
          plt.plot(X[:, 0][y==1], X[:, 1][y==1], "g^", label="Iris virginica")
plt.plot(X[:, 0][y==0], X[:, 1][y==0], "bs", label="Iris versicolor")
plot_svc_decision_boundary(svm_clf1, 4, 5.9)
          plt.xlabel("Petal length", fontsize=14)
plt.ylabel("Petal width", fontsize=14)
          plt.legend(loc="upper left", fontsize=14)
          plt.title("$C = {}$".format(svm_clf1.C), fontsize=16)
          plt.axis([4, 5.9, 0.8, 2.8])
          plt.sca(axes[1])
          plt.plot(X[:, \ 0][y==1], \ X[:, \ 1][y==1], \ "g^")
          plt.plot(X[:, 0][y==0], X[:, 1][y==0], "bs")
          plot_svc_decision_boundary(svm_clf2, 4, 5.99)
```

```
plt.xlabel("Petal length", fontsize=14)
plt.title("$C = {}$".format(svm_clf2.C), fontsize=16)
plt.axis([4, 5.9, 0.8, 2.8])
save fig("regularization plot")
Saving figure regularization_plot
                                                                   C = 100
   2.5
             Iris virginica
Petal width
             Iris versicolor
   2.0
   1.0
                                5.25
                                                4.00
                                                                      5.00
                                                                            5.25
     4.00
                          5.00
                                     5.50
                                           5.75
                                                            4.50
                                                                                 5.50
                                                                                       5.75
               450
                     4 75
                                                                 4 75
                     Petal length
                                                                 Petal length
```

## **Non-linear classification**

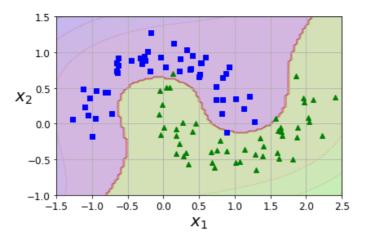
```
In [11]: |X1D = np.linspace(-4, 4, 9).reshape(-1, 1)
          X2D = np.c_{[X1D, X1D**2]}
          y = np.array([0, 0, 1, 1, 1, 1, 1, 0, 0])
          plt.figure(figsize=(10, 3))
          plt.subplot(121)
          plt.grid(True, which='both')
          plt.axhline(y=0, color='k')
          plt.plot(X1D[:, 0][y==0], np.zeros(4), "bs")
          plt.plot(X1D[:, 0][y==1], np.zeros(5), "g^")
          plt.gca().get_yaxis().set_ticks([])
          plt.xlabel(r"$x_1$", fontsize=20)
          plt.axis([-4.5, 4.5, -0.2, 0.2])
          plt.subplot(122)
          plt.grid(True, which='both')
          plt.axhline(y=0, color='k')
          plt.axvline(x=0, color='k')
          plt.plot(X2D[:, 0][y==0], X2D[:, 1][y==0], "bs")
plt.plot(X2D[:, 0][y==1], X2D[:, 1][y==1], "g^")
          plt.xlabel(r"$x_1$", fontsize=20)
plt.ylabel(r"$x_2$ ", fontsize=20, rotation=0)
          plt.gca().get_yaxis().set_ticks([0, 4, 8, 12, 16])
          plt.plot([-4.5, 4.5], [6.5, 6.5], "r--", linewidth=3)
          plt.axis([-4.5, 4.5, -1, 17])
          plt.subplots adjust(right=1)
          save_fig("higher_dimensions_plot", tight_layout=False)
          plt.show()
```

Saving figure higher dimensions plot



```
In [12]:
          from sklearn.datasets import make moons
          X, y = make_moons(n_samples=100, noise=0.15, random_state=42)
          def plot dataset(X, y, axes):
              plt.plot(X[:, 0][y==0], X[:, 1][y==0], "bs")
              plt.plot(X[:, 0][y==1], X[:, 1][y==1], "g^")
              plt.axis(axes)
              plt.grid(True, which='both')
              plt.xlabel(r"$x_1$", fontsize=20)
plt.ylabel(r"$x_2$", fontsize=20, rotation=0)
          plot dataset(X, y, [-1.5, 2.5, -1, 1.5])
          plt.show()
              1.5
              1.0
              0.5
          X_2
              0.0
             -0.5
            -1.0
                -1.5
                    -1.0
                          -0.5
                                0.0
                                     0.5
                                          1.0
                                                1.5
                                                     2.0
                                                          2.5
                                     x_1
In [13]: from sklearn.datasets import make moons
          from sklearn.pipeline import Pipeline
          from sklearn.preprocessing import PolynomialFeatures
          polynomial svm clf = Pipeline([
                  ("poly_features", PolynomialFeatures(degree=3)),
                  ("scaler", StandardScaler()),
("svm_clf", LinearSVC(C=10, loss="hinge", random_state=42))
              1)
          polynomial svm clf.fit(X, y)
Out[13]: Pipeline(memory=None,
               steps=[('poly features', PolynomialFeatures(degree=3, include bias=True,
          interaction_only=False)), ('scaler', StandardScaler(copy=True, with_mean=True,
          with_std=True)), ('svm_clf', LinearSVC(C=10, class_weight=None, dual=True, fit
          intercept=True,
               intercept_scaling=1, loss='hinge', max_iter=1000, multi_class='ovr',
               penalty='\bar{l}2', random_state=42, tol=0.0001, verbose=0))])
In [14]: def plot_predictions(clf, axes):
              x0s = np.linspace(axes[0], axes[1], 100)
              x1s = np.linspace(axes[2], axes[3], 100)
              x0, x1 = np.meshgrid(x0s, x1s)
              X = np.c_[x0.ravel(), x1.ravel()]
              y_pred = clf.predict(X).reshape(x0.shape)
              y decision = clf.decision function(X).reshape(x0.shape)
              plt.contourf(x0, x1, y_pred, cmap=plt.cm.brg, alpha=0.2)
              plt.contourf(x0, x1, y_decision, cmap=plt.cm.brg, alpha=0.1)
          plot_predictions(polynomial_svm_clf, [-1.5, 2.5, -1, 1.5])
          plot_dataset(X, y, [-1.5, 2.5, -1, 1.5])
          save fig("moons polynomial svc plot")
          plt.show()
```

Saving figure moons\_polynomial\_svc\_plot



```
In [15]: from sklearn.svm import SVC
         poly_kernel_svm_clf = Pipeline([
                  ("scaler", StandardScaler()),
("svm_clf", SVC(kernel="poly", degree=3, coef0=1, C=5))
         poly kernel svm clf.fit(X, y)
Out[15]: Pipeline(memory=None,
               steps=[('scaler', StandardScaler(copy=True, with_mean=True, with_std=Tru
          e)), ('svm_clf', SVC(C=5, cache_size=200, class_weight=None, coef0=1,
            decision_function_shape='ovr', degree=3, gamma='auto_deprecated',
            kernel='poly', max_iter=-1, probability=False, random_state=None,
            shrinking=True, tol=0.001, verbose=False))])
In [16]: poly100_kernel_svm_clf = Pipeline([
                  ("scaler", StandardScaler()),
("svm_clf", SVC(kernel="poly", degree=10, coef0=100, C=5))
         poly100 kernel svm clf.fit(X, y)
Out[16]: Pipeline(memory=None,
               steps=[('scaler', StandardScaler(copy=True, with_mean=True, with_std=Tru
          e)), ('svm_clf', SVC(C=5, cache_size=200, class_weight=None, coef0=100,
            decision_function_shape='ovr', degree=10, gamma='auto_deprecated',
            kernel='poly', max_iter=-1, probability=False, random_state=None,
            shrinking=True, tol=0.001, verbose=False))])
```

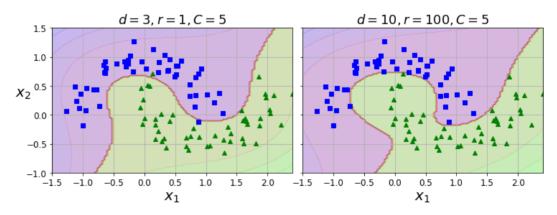
```
In [17]: fig, axes = plt.subplots(ncols=2, figsize=(10.5, 4), sharey=True)

plt.sca(axes[0])
plot_predictions(poly_kernel_svm_clf, [-1.5, 2.45, -1, 1.5])
plot_dataset(X, y, [-1.5, 2.4, -1, 1.5])
plt.title(r"$d=3, r=1, C=5$", fontsize=18)

plt.sca(axes[1])
plot_predictions(poly100_kernel_svm_clf, [-1.5, 2.45, -1, 1.5])
plot_dataset(X, y, [-1.5, 2.4, -1, 1.5])
plt.title(r"$d=10, r=100, C=5$", fontsize=18)
plt.ylabel("")

save_fig("moons_kernelized_polynomial_svc_plot")
plt.show()
```

Saving figure moons\_kernelized\_polynomial\_svc\_plot



```
In [18]: def gaussian_rbf(x, landmark, gamma):
              return np.exp(-qamma * np.linalq.norm(x - landmark, axis=1)**2)
         qamma = 0.3
         x1s = np.linspace(-4.5, 4.5, 200).reshape(-1, 1)
         x2s = gaussian_rbf(x1s, -2, gamma)
         x3s = gaussian_rbf(x1s, 1, gamma)
         XK = np.c_[gaussian_rbf(X1D, -2, gamma), gaussian_rbf(X1D, 1, gamma)]
         yk = np.array([0, 0, 1, 1, 1, 1, 1, 0, 0])
         plt.figure(figsize=(10.5, 4))
         plt.subplot(121)
         plt.grid(True, which='both')
         plt.axhline(y=0, color='k')
         plt.scatter(x=[-2, 1], y=[0, 0], s=150, alpha=0.5, c="red")
         plt.plot(X1D[:, 0][yk==0], np.zeros(4), "bs")
         plt.plot(X1D[:, 0][yk==1], np.zeros(5), "g^")
         plt.plot(x1s, x2s, "g--")
plt.plot(x1s, x3s, "b:")
         plt.gca().get_yaxis().set_ticks([0, 0.25, 0.5, 0.75, 1])
         plt.xlabel(r"$x_1$", fontsize=20)
         plt.ylabel(r"Similarity", fontsize=14)
         plt.annotate(r'$\mathbf{x}$'
                       xy=(X1D[3, 0], 0),
                       xytext=(-0.5, 0.20),
                       ha="center"
                       arrowprops=dict(facecolor='black', shrink=0.1),
                       fontsize=18,
         plt.text(-2, 0.9, "$x_2$", ha="center", fontsize=20)
         plt.text(1, 0.9, "$x_3$", ha="center", fontsize=20)
```

```
plt.axis([-4.5, 4.5, -0.1, 1.1])
          plt.subplot(122)
          plt.grid(True, which='both')
          plt.axhline(y=0, color='k')
          plt.axvline(x=0, color='k')
          plt.plot(XK[:, \ 0][yk==0], \ XK[:, \ 1][yk==0], \ "bs")
          plt.plot(XK[:, 0][yk==1], XK[:, 1][yk==1], "q^")
          plt.xlabel(r"$x_2$", fontsize=20)
plt.ylabel(r"$x_3$ ", fontsize=20, rotation=0)
          plt.annotate(r'$\phi\left(\mathbf{x}\right)$',
                        xy=(XK[3, 0], XK[3, 1]),
                        xytext=(0.65, 0.50),
                        ha="center".
                        arrowprops=dict(facecolor='black', shrink=0.1),
                        fontsize=18,
          plt.subplots adjust(right=1)
          save fig("kernel method plot")
          plt.show()
Saving figure kernel method plot
             1.00
                                                        1.0
                                     X<sub>3</sub>
                                                        0.8
             0.75
                                                                      •
                                                     X3<sup>0.6</sup>
          Similarity
                                                                               \phi(\mathbf{x})
             0.50
                                                        0.4
             0.25
                                                        0.2
             0.00
                                                        0.0
                                  Ó
                                                             0.0
                                                                  0.2
                                                                        0.4
                                                                              0.6
                                                                                    0.8
                                                                                          1.0
                                 x_1
                                                                           x_2
In [19]: x1_{example} = X1D[3, 0]
          for landmark in (-2, 1):
              k = gaussian_rbf(np.array([[x1_example]]), np.array([[landmark]]), gamma)
              print("Phi({}, {}) = {}".format(x1 example, landmark, k))
          Phi(-1.0, -2) = [0.74081822]
Phi(-1.0, 1) = [0.30119421]
In [20]: rbf_kernel_svm_clf = Pipeline([
                   ("scaler", StandardScaler()),
                   ("svm clf", SVC(kernel="rbf", gamma=5, C=0.001))
          rbf kernel svm clf.fit(X, y)
Out[20]: Pipeline(memory=None,
               steps=[('scaler', StandardScaler(copy=True, with mean=True, with std=Tru
          e)), ('svm_clf', SVC(C=0.001, cache_size=200, class_weight=None, coef0=0.0,
            decision_function_shape='ovr', degree=3, gamma=5, kernel='rbf'
            max_iter=-1, probability=False, random_state=None, shrinking=True,
            tol=0.001, verbose=False))])
In [21]: from sklearn.svm import SVC
          gamma1, gamma2 = 0.1, 5
          C1, C2 = 0.001, 1000
          hyperparams = (gamma1, C1), (gamma1, C2), (gamma2, C1), (gamma2, C2)
```

```
svm clfs = []
for gamma, C in hyperparams:
    rbf_kernel_svm_clf = Pipeline([
             ("scaler", StandardScaler()),
("svm_clf", SVC(kernel="rbf", gamma=gamma, C=C))
    rbf_kernel_svm_clf.fit(X, y)
    svm_clfs.append(rbf_kernel_svm_clf)
fig, axes = plt.subplots(nrows=2, ncols=2, figsize=(10.5, 7), sharex=True, shar
for i, svm clf in enumerate(svm clfs):
    plt.sca(axes[i // 2, i % 2])
    plot_predictions(svm_clf, [-1.5, 2.45, -1, 1.5])
    plot_dataset(X, y, [-1.5, 2.45, -1, 1.5])
    gamma, C = hyperparams[i]
    plt.title(r"\$\gamma = {}, C = {}\$".format(gamma, C), fontsize=16)
    if i in (0, 1):
        plt.xlabel("")
    if i in (1, 3):
        plt.ylabel("")
save fig("moons rbf svc plot")
plt.show()
Saving figure moons_rbf_svc_plot
```

