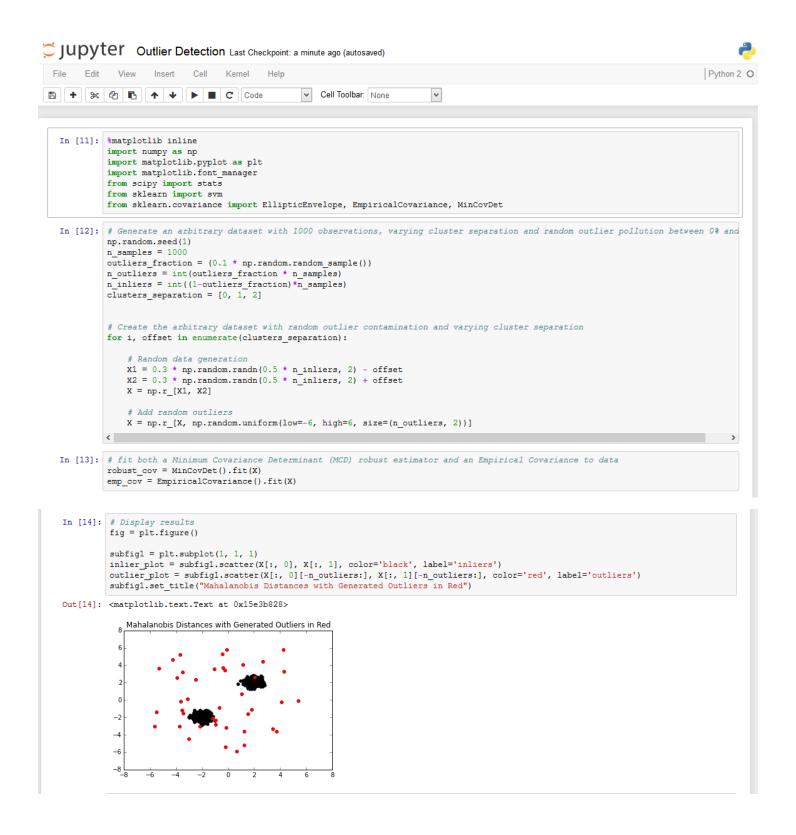
## Outlier Detection Using Anaconda Python in Project Jupyter Notebook



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In [15]: # The above chart shows a number of red dots equal to the randomly generated n_outliers variable.
         # However, I declared an assumption in the main document that I would not know the outlier contamination
         # Therefore, I can not use the "n_outliers" variable for the second stage of this analysis.
         \# Instead, I must ignore that I have that number stored from the task of generating the data and
         # estimate it from the Mahalanobis Distances as if this data was something that I was completely naive to.
         # Also, to graph shows red dots near or invading my black clusters (depending on random seed), so I disagree
         # with some of the assignments and would have assigned fewer outliers by hand.
         emp_mahal = emp_cov.mahalanobis(X - np.mean(X, 0)) ** .333
         robust_mahal = robust_cov.mahalanobis(X - robust_cov.location_) ** .333
         ave_mahal = (emp_mahal + robust_mahal)/2
         mean = np.mean(ave mahal)
         sd = np.std(ave_mahal)
         limit = mean + (3*sd)
         contamination = np.float(np.sum(ave_mahal > limit)) / np.float(np.sum(ave_mahal > -100))
In [16]: contamination
Out[16]: 0.025025025025025027
In [17]: # Just for laughs, compare this new estimate to the outlier contamination rate imposed at random during data generation,
          # and there are indeed fewer assignments than before, exactly as desired:
         (np.float(n_outliers))/(np.float(n_samples))
Out[17]: 0.041
In [22]: # Moving on, I take that new estimate of outlier contamination, and make it an assumption as I compare
          # two outlier detection tools, a single class support vector machine tuned for outlier detection (better
         # with separated clusters or strongly non-Gaussian distributions, and a Robust Covariance Estimator (a.k.a
         # elliptic envelope...better for Gaussian distributions, but fails otherwise)
         classifiers = {
             "One-Class SVM": svm.OneClassSVM(nu=0.95 * outliers_fraction + 0.05,
                                               kernel="rbf", gamma=0.1),
             "Robust Covariance Estimator": EllipticEnvelope(contamination=.1)}
         # Compare classifiers
         xx, yy = np.meshgrid(np.linspace(-10, 10, 500), np.linspace(-10, 10, 500))
         contaminants = np.sum(ave_mahal > limit)
```

```
In [23]: # Fit and Plot
         plt.figure(figsize=(10, 5))
         for i, (clf name, clf) in enumerate(classifiers.items()):
             # fit the data and tag outliers
             clf.fit(X)
             y_pred = clf.decision_function(X).ravel()
             threshold = stats.scoreatpercentile(y_pred,100 * outliers_fraction)
             y_pred = y_pred > threshold
             n errors = np.sum((y pred != ground proj))
             # plot the levels lines and the points
             Z = clf.decision_function(np.c_[xx.ravel(), yy.ravel()])
             Z = Z.reshape(xx.shape)
            subplot = plt.subplot(1, 2, i + 1)
             subplot.set title("Outlier Detection")
            subplot.contourf(xx, yy, Z, levels=np.linspace(Z.min(), threshold, 7),cmap=plt.cm.Blues r)
             a = subplot.contour(xx, yy, Z, levels=[threshold],linewidths=2, colors='red')
             subplot.contourf(xx, yy, Z, levels=[threshold, Z.max()],colors='yellow')
             b = subplot.scatter(X[:-contaminants, 0], X[:-contaminants, 1], c='white')
             c = subplot.scatter(X[-contaminants:, 0], X[-contaminants:, 1], c='black')
             subplot.axis('tight')
             subplot.legend(
                  [a.collections[0], b, c],
                  ['Learned Function Boundary', 'Projected Inliers', 'Projected Outliers'],
                  prop=matplotlib.font_manager.FontProperties(size=10))
             subplot.set xlabel("%d. %s" % (i + 1, clf name))
             subplot.set_xlim((-10, 10))
             subplot.set ylim((-10, 10))
         plt.subplots_adjust(0.04, 0.1, 0.96, 0.94, 0.1, 0.26)
         plt.show()
```

ground\_proj = np.ones(n\_samples, dtype=int)

ground\_proj[-contaminants:] = 0

