Instructions

This notebook goes from micro-variable climate data (same data as used in our UAI 2016 paper "Unsupervised Discovery of El Nino Using Causal Feature Learning on Micro-Level Climate Data") to macro-level states.

I simplified our code a little bit, but you should at least get a "Westerly Winds" and "El Nino" clusters at the end of this notebook, and see that they correlate strongly.

To use your own data, you only need to modify the "The Dataset" cell -- substitute your own data for our X and Y arrays.

If you like, you can also experiment with neural net training in the "Learning $P(Y \mid X)$ " cell. Make sure the validation error converges. Feel free to experiment with other neural net architectures if you're comfortable with Lasagne! If your PC has a GPU, this code will automatically use it. If you have no GPU and the training is slow, you can stop training at any time by pressing ctrl+c (on Linux) and the script will load the best network learned during training on the next step (during training, the current-best net is saved to a file).

You need to write your own code to visualize your results, the last cell here is tailored to display climate maps.

Good Luck!

Krzysztof Chalupka, Caltech, July 2016.

```
In [6]: """ This cell sets up the Python environment. Make sure you have the follow
        ing packages installed:
        numpy
        matplotlib
        theano
        lasagne
        sklearn
        All of them are available through $ pip install [package_name].
        This is only tested on Ubuntu Linux, I've heard Theano is a pain to get wor
        king on Windows.
        import sys
        import numpy as np
        np.random.seed(1423)
        import matplotlib
        %matplotlib inline
        from matplotlib import pyplot as plt
        from sklearn.cluster import KMeans
        from sklearn.preprocessing import StandardScaler
        import theano
        import lasagne
        sys.setrecursionlimit(1000000) # Otherwise, joblib won't save deep nets.
        imshape = (55, 9) # This is the grid shape of our images, stored here for p
        lotting reference.
```

The Dataset

The following code loads our data into the correct format:

- X -- numpy array of size (n datapoints, n input dim). Each row corresponds to one input value.
- Y -- numpy array of size (n datapoints, n output dim). Each row is one output value.

In our case, each row of X is a (flattened) map of Pacific zonal wind strength, and each row of Y a (flattened) map of Pacific water temperature over the same region. In our case, n input dim == n output dim, but this need not be the case at all.

```
In [8]: import joblib
        # Load the data.
        ## PLUG YOUR OWN DATA HERE. 'coords' is only needed to display climate maps
         . Your data
        ## can only contain X and Y arrays.
        X, Y, coords = joblib.load('elnino_data.pkl')
        # Create a randomized, normalized training and validation set.
        shuffled ids = np.random.permutation(X.shape[0])
        X \text{ tr} = X[\text{shuffled ids}[:int(X.shape[0]*.9)]]
        Y tr = Y[shuffled ids[:int(Y.shape[0]*.9)]]
        X ts = X[shuffled ids[int(X.shape[0]*.9):]]
        Y ts = Y[shuffled ids[int(Y.shape[0]*.9):]]
        x_scaler = StandardScaler().fit(X_tr)
        y_scaler = StandardScaler().fit(Y_tr)
        X_{tr} = x_{scaler.transform}(X_{tr})
        X_{ts} = x_{scaler.transform}(X_{ts})
        Y_tr = y_scaler.transform(Y_tr)
        Y_ts = y_scaler.transform(Y_ts)
```

Learning P(Y | X)

The first step of Causal Feature Learning (CFL) is to cluster x's according to $P(Y \mid x)$ conditional densities. In this demonstration, for simplicity, we will approximate $P(Y \mid x)$ with its expected value $E[P(Y \mid x)]$. This means we assume that if two distributions have equal means, they themselves are equal. It is possible to efficiently relax this assumption by using Mixture Density Networks (Bishop 1995) to approximate all moments of a distribution.

Learning E[P(Y | x)] amounts to regressing y on x. We do this using a neural network and the Lasagne package.

```
In [9]: | # Define the network architecture.
        dropout = lasagne.layers.dropout
        batch_norm = lasagne.layers.batch_norm
        net = lasagne.layers.InputLayer(shape=(None, X[0].size))
        net = dropout(net, p=.2)
        net = lasagne.layers.DenseLayer(net, num_units=1024, W=lasagne.init.HeNorma
        l(gain=1))
        net = dropout(net, p=.5)
        net = lasagne.layers.DenseLayer(net, num units=1024, W=lasagne.init.HeNorma
        l(gain=1))
        net = dropout(net, p=.5)
        net = lasagne.layers.DenseLayer(net, num units=Y[0].size,
            nonlinearity=lasagne.nonlinearities.identity)
        # Define the cost function in Theano.
        input var = lasagne.layers.get all layers(net)[0].input var
        target var = theano.tensor.matrix('target')
        train_cost = lasagne.objectives.squared_error(lasagne.layers.get_output(net
        ), target var).mean() +\
            0.0001 * lasagne.regularization.regularize_network_params(net, lasagne.
        regularization.12)
        valid_cost = lasagne.objectives.squared_error(
            lasagne.layers.get_output(net, deterministic=True), target_var).mean()
        # Compile the training algorithm in Lasagne.
        parameters = lasagne.layers.get_all_params(net, trainable=True)
        updates = lasagne.updates.adam(train_cost, parameters, learning_rate=le-4)
        train = theano.function([input_var, target_var],
            train_cost, updates=updates, allow_input_downcast=True)
        valid = theano.function([input_var, target_var], valid_cost, allow_input_do
        wncast=True)
        # Train.
        BATCHSIZE = 128
        val loss = valid(X ts, Y ts)
        print('Validation mse before training: {:.2g}.'.format(float(val loss)))
        best loss = np.inf
        for epoch id in range(1000):
            tr loss = 0
            for batch_id in range(int(np.floor(X_tr.shape[1]/BATCHSIZE))):
                tr_loss += train(X_tr[batch_id*BATCHSIZE:(batch_id+1)*BATCHSIZE],
                                  Y_tr[batch_id*BATCHSIZE:(batch_id+1)*BATCHSIZE])
            val_loss = valid(X_ts, Y_ts)
            if np.isfinite(val_loss) and val_loss < best_loss:</pre>
                joblib.dump(net, 'nn_params/nn.pkl')
                best_loss = val_loss
            sys.stdout.write('\rEpoch {}. Valid loss {:.4g} [best {:.4g}]. Train lo
        ss {:.4g}.'.format(
                     epoch_id, float(val_loss), float(best_loss), tr_loss/float(np.f
        loor(X_tr.shape[1]/BATCHSIZE))))
            sys.stdout.flush()
```

Validation mse before training: 1.3. Epoch 99. Valid loss 0.5163 [best 0.5163]. Train loss 0.8422.

Finding the Observational Partition of X

Finding the observational partition of X amounts to clustering the X data according to $P(Y \mid X)$. That is, we put two x's in the same bucket if the neural net we trained maps them to the same values.

```
In [10]: # Load the best network found and compile it.
    net = joblib.load('nn_params/nn.pkl')
    input_var = lasagne.layers.get_all_layers(net)[0].input_var
    p_y_x = theano.function([input_var], lasagne.layers.get_output(net))

# Cluster Xs.
N_CLASSES = 4
x_lbls = KMeans(n_clusters=N_CLASSES, n_init=10, n_jobs=-1).fit_predict(p_y_x(np.vstack([X_tr, X_ts])))
```

Finding the Observational Partition of Y

To find the partition of Y, we want to put together all y1 and y2 if $P(y1 \mid x) == P(y2 \mid x)$ for each x. This procedure is a little bit more interesting than clustering the x's, and is described in our UAI 2016 paper. We describe it briefly here.

We've already clustered x's in a way that that guarantees that if x1 and x2 belong to the same x_lbls class, then $P(y \mid x1) == P(y \mid x2)$. Thus, the requirement $P(y1 \mid x) == P(y2 \mid x)$ can be substituted by $P(y1 \mid x_lbls==0) == P(y2 \mid x_lbls==0)$ for any observational X-class O. Since x_lbls is discrete, we should have plenty of data per each x_lbls class. We will approximate $P(y \mid x_lbls==0)$ using the distance of y to the closest (except for itself) Y-point whose corresponding x belongs to x_lbls==O.

```
In [11]: y ftrs = np.zeros((Y.shape[0], np.unique(x lbls).size))
         # Loop, not vectorized, to save memory. Can take a while.
         for y_id, y in enumerate(np.vstack([Y_tr, Y_ts])):
             if y id % 100==0:
                 sys.stdout.write('\rComputing P(y | x lbls) features, iter \{\}/\{\}...
          '.format(y id, Y.shape[0]))
                 sys.stdout.flush()
             for x lbl id, x lbl in enumerate(np.unique(x lbls)):
                 # Find ids of xs in this x lbls class.
                 x lbl ids = np.where(x lbls==x lbl)[0]
                 # Compute distances of y to all y's in this x_lbls class and sort t
         hem.
                 sorted_dists = np.sort(np.sum((y-np.vstack([Y_tr, Y_ts])[x_lbl_ids]
         )**2, axis=1))
                 # Find the mean distance to the 4 closest points (exclude the actua
         lly closest point though).
                 y_ftrs[y_id][x_lbl_id] = sorted_dists[1:5].mean()
         print('Done. Clustering P(y | x_lbls).')
         y_lbls = KMeans(n_clusters=N_CLASSES, n_init=10, n_jobs=-1).fit_predict(y_f
```

Computing P(y \mid x_lbls) features, iter 13100/13140...Done. Clustering P(y \mid x_lbls).

Understanding the Results

Visualizing the observational partition is data-specific. In our case, since both X and Y are images, we can visualize the means of each observational cluster to gain some insight into what it contains.

But, one of the virtues of the method is that it is interpretation-agnostic. The observational partition can be used as a causal hypothesis to drive experimentation. This can be done whether the inputs and outputs are easily interpretable or not at all.

```
In [13]: fig = plt.figure(figsize=(15,10), facecolor='white')
         X_raw = x_scaler.inverse_transform(np.vstack([X_tr, X_ts]))
         Y_raw = y_scaler.inverse_transform(np.vstack([Y_tr, Y_ts]))
         levels = np.linspace(-4,4,30)
         for x_cluster_id in range(4):
             ax = plt.subplot2grid((4,2), (x_cluster_id, 0))
             # Plot the cluster's mean difference from all frames' mean.
             cluster_mean = (X_raw[x_lbls==x_cluster_id].mean(axis=0)-X_raw.mean(axi
         s=0)).reshape(imshape).T
             im=ax.contourf(coords['x'].ravel(), coords['y'].ravel(), cluster mean,
         levels=levels, cmap='BrBG r')
             ax.set_xticks([]); ax.set_yticks([])
         levels=np.linspace(-3,5.5,30)
         for y cluster id in range(4):
             ax = plt.subplot2grid((4,2), (y cluster id, 1))
             # Plot the cluster's mean difference from all frames' mean.
             cluster_mean = (Y_raw[y_lbls==y_cluster_id].mean(axis=0)-Y_raw.mean(axi
         s=0)).reshape(imshape).T
             im=ax.contourf(coords['x'].ravel(), coords['y'].ravel(), cluster_mean,
         levels=levels, cmap='coolwarm')
             ax.set_xticks([]); ax.set_yticks([])
         # Compute and print P(y_lbl | x_lbl)
         P_CE = np.array([np.bincount(y_lbls.astype(int)[x_lbls==x_lbl],
             minlength=y_lbls.max()+1).astype(float) for x_lbl in np.sort(np.unique(
         x_lbls))])
         P_CE = P_CE/P_CE.sum()
         P_E_given_C = P_CE/P_CE.sum(axis=1, keepdims=True)
         print('P(TempCluster | WindCluster):')
         print(P_E_given_C)
```

