## Homework 5

November 30, 2015

## 1 Inference Homework 5

- 1.1 Alex Pine
- $1.2 \quad 2015/11/21$
- 1.3 Question 1

I collaborated with Israel Malkin and Peter Li on this problem.

Let T be a tree MRF, and let T' be a directed tree with the same skeleton as T, and one root node, R, which has all edges are directed away from it.

T' must be a moral graph, where each node has at most one parent. If it were not, that would mean there is some node n with at least two parents,  $p_1$  and  $p_2$ . Then there would be at least two directed paths which end with n, one including  $p_1$ , and another including  $p_2$ . If both of those paths started with R, T would not be a tree, because trees have exactly one path between two nodes by definition. If instead, one of the paths ends in a node that is not R, this node would constitute a second root node, which does not exist by construction. Thus T' must be moral.

Since T' is moral, it encodes the same probability distribution as T, which we'll call  $p_T(\mathbf{x})$ .  $p_T(\mathbf{x})$  can thus be factorized according to T':

$$p_T(\mathbf{x}) = \prod_{x_i \in \mathbf{x}} p(x_i \mid \text{pa}(x))$$

Since all nodes have at most one parent node, we can denote  $pa(x) = x_p$ .

$$p_T(\mathbf{x}) = \prod_{x_i \in \mathbf{x}} p(x_i \mid x_p)$$

Applying Bayes' rule, this becomes

$$p_T(\mathbf{x}) = \prod_{x_i \in \mathbf{x}} \frac{p(x_i, x_p)}{p(x_p)}$$

Multiplying this by  $\prod_{x_i \in \mathbf{x}} \frac{p(x_i)}{p(x_i)} = 1$ , this becomes

$$p_T(\mathbf{x}) = \prod_{x_i \in \mathbf{x}} \frac{p(x_i, x_p)}{p(x_i)p(x_p)} \prod_{x_i \in \mathbf{x}} p(x_i)$$

Since the set of edges T' are between parents and children, and each node is a child of one or zero parents, each i appears once in the edges  $(i,j) \in T'$ . This means we can change the indices of the first product to be the set of edges in T' without changing the value of the expression. If we also denote V to be the set containing the marginal variables of  $\mathbf{x}$ , then this formula can be rewritten as

$$p_T(\mathbf{x}) = \prod_{(i,j) \in T} \frac{p(x_i, x_j)}{p(x_i)p(x_j)} \prod_{x_i \in V} p(x_i)$$

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This probability distribution corresponds to an MRF with the same nodes and edges as T, where each node  $x_i$  has the potential  $p(x_i)$ , and each edge has the potential  $p(x_i, x_j)$ , which is what we were trying to prove.

# 2 Question 2

I collaborated with Maya Rotmensch, Peter Li, and Israel Malkin on this problem.

The output was written to a file called "hw5\_prob2.uai". Some debugging output is shown at the end of the code to illustrate that it works properly.

```
In [3]: import numpy as np
        def read_chowlui_input():
            input_matrix = []
            with open('inference/hw5/data/chowliu-input.txt', 'r') as f:
                for line in f:
                    input_matrix.append([int(token) for token in line.split()])
            return np.array(input_matrix)
        def read_labels_input():
            input_labels = []
            with open('inference/hw5/data/names.txt', 'r') as f:
                for line in f:
                    input_labels.append(line.strip())
            return np.array(input_labels)
        input_matrix = read_chowlui_input()
        labels = read_labels_input()
In [4]: import scipy.sparse.csgraph
        # Given the input matrix, which has one row for each data point, and a column for each item,
        # create two matrices:
        # 1) both_count_matrix:
             When i > j: A[i][j] = the number of times items i and j were both present.
             When i == j: A[i][j] = the number of times item i appears in the data set.
             When i < j: A[i][j] = zero. This portion is not used.
        # 2) single_count_matrix:
             When i != j : B[i][j] = the number of times item i was present and j was not.
             When i == j: B[i][j] = zero. This portion is not used.
        def create_count_matrices(input_matrix):
            both_count_matrix = np.zeros((input_matrix.shape[1], input_matrix.shape[1]), dtype=int)
            single_count_matrix = np.zeros((input_matrix.shape[1], input_matrix.shape[1]), dtype=int)
            for row in input_matrix:
                for i in range(len(row)):
                    for j in range(len(row)):
                        if i > j and row[i] == 1 and row[j] == 1:
                            both_count_matrix[i][j] += 1
                        if row[i] == 1 and row[j] == 0:
                            single_count_matrix[i][j] += 1
                    if row[i] == 1:
                        both_count_matrix[i][i] += 1
            return both_count_matrix, single_count_matrix
        both_count_matrix, single_count_matrix = create_count_matrices(input_matrix)
```

```
In [6]: # Functions needed to create a matrix that has the mutual information between each item,
        # based on their empirical probabilities.
        # I actually take the inverse of the mutual information, so that a minimum spanning tree
        # algorithm can be used to find the maximum spanning tree.
        import itertools
        import math
        import numpy.ma as ma
        # Computes one part of the sum of the "empirical" mutual information between x_i and x_j.
        def mutual_info_part(num_xixj, num_xi, num_xj, total):
            if not num_xixj or not num_xi or not num_xj:
                return 0.0
            p_xixj = 1.0 * num_xixj / total
            p_xi = 1.0 * num_xi / total
            p_xj = 1.0 * num_xj / total
            return p_xixj * math.log(p_xixj / (p_xi * p_xj), 2)
        def mutual_info(i, j, num_data, both_count_matrix, single_count_matrix):
           num_xi_one = both_count_matrix[i][i]
            num_xj_one = both_count_matrix[j][j]
            num_xi_zero = num_data - num_xi_one
            num_xj_zero = num_data - num_xj_one
            info = 0.0
            # 11 case
            info += mutual_info_part(both_count_matrix[i][j], num_xi_one, num_xj_one, num_data)
            # 10 case
            info += mutual_info_part(single_count_matrix[i][j], num_xi_one, num_xj_zero, num_data)
            # 01 case
            info += mutual_info_part(single_count_matrix[j][i], num_xi_zero, num_xj_one, num_data)
            # 00 case
            neither_count = (num_data - both_count_matrix[i][j] - single_count_matrix[i][j]
                             - single_count_matrix[j][i])
            info += mutual_info_part(neither_count, num_xi_zero, num_xj_zero, num_data)
            assert (neither_count + both_count_matrix[i][j] + single_count_matrix[i][j]
                    + single_count_matrix[j][i] == num_data)
            return info
        # Create a matrix where each value is the inverse of the mutual information values
        # between each item.
        def create_inv_mutual_info_matrix(num_data, both_count_matrix, single_count_matrix):
            inv_info_matrix = np.zeros(both_count_matrix.shape)
            # Create a mask for the upper right corner, to mark this matrix as triangular.
            info_mask = np.zeros(both_count_matrix.shape, dtype=int)
            num_items = both_count_matrix.shape[1]
            for i in range(num_items):
                for j in range(i):
                    info = mutual_info(i, j, num_data, both_count_matrix, single_count_matrix)
                    inv_info_matrix[i][j] = 1.0 / info
                for j in range(i, num_items):
                    info_mask[i][j] = 1
            inv_info_matrix = ma.array(inv_info_matrix, mask=info_mask)
            return inv_info_matrix
```

```
num_data = input_matrix.shape[0]
        inv_info_matrix = create_inv_mutual_info_matrix(
            num_data, both_count_matrix, single_count_matrix)
In [7]: import scipy.sparse.csgraph
       mst_rows, mst_columns = scipy.sparse.csgraph.minimum_spanning_tree(inv_info_matrix).nonzero()
       mst_edge_pairs = zip(mst_rows, mst_columns)
In [8]: # Code for calculating the potential functions on nodes and edges, now that we have the
        # structure of the tree.
        def edge_part(num_xixj, num_xi, num_xj, total):
           p_xixj = 1.0 * num_xixj / total
           p_xi = 1.0 * num_xi / total
           p_xj = 1.0 * num_xj / total
           return p_xixj / (p_xi * p_xj)
       def edge_potential(i, j, num_data, both_count_matrix, single_count_matrix):
            num_xi_one = both_count_matrix[i][i]
           num_xj_one = both_count_matrix[j][j]
            num_xi_zero = num_data - num_xi_one
            num_xj_zero = num_data - num_xj_one
            potential = np.zeros((2, 2))
            # 11 case
            potential[1][1] = edge_part(both_count_matrix[i][j], num_xi_one, num_xj_one, num_data)
            # 10 case
            potential[1][0] = edge_part(single_count_matrix[i][j], num_xi_one, num_xj_zero, num_data)
            # 01 case
            potential[0][1] = edge_part(single_count_matrix[j][i], num_xi_zero, num_xj_one, num_data)
            # 00 case
            neither_count = (num_data - both_count_matrix[i][j] - single_count_matrix[i][j]
                             - single_count_matrix[j][i])
            potential[0][0] = edge_part(neither_count, num_xi_zero, num_xj_zero, num_data)
            assert (neither_count + both_count_matrix[i][j] + single_count_matrix[i][j]
                    + single_count_matrix[j][i] == num_data)
            return potential
        def node_potential(i, num_data, both_count_matrix):
            num_xi_one = both_count_matrix[i][i]
            num_xi_zero = num_data - num_xi_one
            return np.array([float(num_xi_one) / num_data, float(num_xi_zero) / num_data])
       def create_graph_potentials(mst_edge_pairs, num_data,
                                    both_count_matrix, single_count_matrix):
            \# Every node i has a 1x2 potential matrix
            node_potentials = np.ndarray((both_count_matrix.shape[0], 2))
            # Every edge (i,j) has a 2x2 potential matrix.
            edge_potentials = np.ndarray((both_count_matrix.shape[0], both_count_matrix.shape[1],
                                          2, 2))
```

```
num_items = both_count_matrix.shape[1]
            for i in range(num_items):
                for j in range(i):
                    edge_potentials[i][j] = edge_potential(i, j, num_data, both_count_matrix,
                                                           single_count_matrix)
                node_potentials[i] = node_potential(i, num_data, both_count_matrix)
            return node_potentials, edge_potentials
       node_potentials, edge_potentials = create_graph_potentials(
            mst_edge_pairs, num_data, both_count_matrix, single_count_matrix)
In [10]: # Creates a string that reprsents the Markov random field in UAI format.
         # http://www.hlt.utdallas.edu/~vgogate/uai14-competition/modelformat.html
         def create_graph_as_uai_file(mst_edge_pairs, node_potentials, edge_potentials):
             num_vars = node_potentials.shape[0]
             network_type = 'MARKOV'
             num_vars_str = str(num_vars)
             var_cardinals = ' '.join(['2']*num_vars)
             num_cliques = str(len(mst_edge_pairs) + num_vars)
             node_cliques = ['1 ' + str(i) for i in range(num_vars)]
             edge_cliques = [' '.join(['2', str(i), str(j)]) for i, j in mst_edge_pairs]
             preamble = ([network_type, num_vars_str, var_cardinals, num_cliques] + node_cliques
                         + edge_cliques)
             function_tables = []
             # node potentials
             for i in range(num_vars):
                 prob_str = ' '.join([' ', str(node_potentials[i][0]), str(node_potentials[i][1])])
                 function_tables += ['', '2', prob_str]
             # edge potentials
             for i, j in mst_edge_pairs:
                 assert i > j
                 prob00 = edge_potentials[i][j][0][0]
                 prob01 = edge_potentials[i][j][0][1]
                 prob_str1 = ' '.join([' ', str(prob00), str(prob01)])
                 prob10 = edge_potentials[i][j][1][0]
                 prob11 = edge_potentials[i][j][1][1]
                prob_str2 = ' '.join([' ', str(prob10), str(prob11)])
                 function_tables += ['', '4', prob_str1, prob_str2]
             return '\n'.join(preamble + function_tables + [''])
         uai_str = create_graph_as_uai_file(mst_edge_pairs, node_potentials, edge_potentials)
         with open ('hw5_prob2.uai', 'w') as f:
             f.write(uai_str)
In [15]: # Debug output - Edge potentials when both items are present for the first 25 edges.
         # Format:
         # item 1 index, item 2 index, item 1 label -- item 2 label : edge potential when
         # both are present.
         for i, j in mst_edge_pairs[:25]:
             print i, j, labels[i], '--', labels[j], ":", edge_potentials[i][j][1][1]
```

```
13 11 books -- book: 17.8188951664
13 12 books -- bookcase : 32.2502421308
15 14 bottles -- bottle : 21.8532110092
18 17 boxes -- box : 13.7630003152
20 2 building -- awning : 2.83260887878
20 4 building -- balcony : 3.15318242536
22 16 cabinet -- bowl : 15.1463651498
25 24 cars -- car : 7.9865994979
32 22 countertop -- cabinet : 19.6495829675
35 8 cushion -- bed : 12.1792726461
36 26 desk -- chair : 5.25427127617
36 28 desk -- clock : 6.07160236357
39 20 dome -- building : 2.81421531463
40 6 door -- bars : 3.19225146199
41 22 drawer -- cabinet : 15.4129411765
45 1 fireplace -- armchair : 15.1631944444
46 1 floor -- armchair : 2.60851983636
46 13 floor -- books : 2.42753607988
46 20 floor -- building : 0.0921954638603
46 26 floor -- chair : 2.34104808146
46 27 floor -- chandelier: 2.52800622804
47 23 flowers -- candle : 12.0635359116
51 46 ground -- floor: 0.0828942173358
52 40 handrail -- door : 2.77395644283
53 50 headstone -- grass : 7.11527494908
```

#### 2.1Question 3

I collaborated with Israel Malkin and Peter Li on this problem.

Define kernel function  $p(x_i, x_j) = C_{i,j}^{-1}$ , where C is a covariance matrix and  $C_{i,j} = k(x_i, x_j)$ , and  $x_i, x_j \in$  $\mathbf{x_n}$ , a sequence of i.i.d random variables.

Drawing the first variable,  $x_1$ , results in a covariance matrix  $C_1 = (k(x_1, x_1))$ , which implies that  $C_1^{-1} = \left(\frac{1}{k(x_1, x_1)}\right)$  which implies  $p(x_1, x_1) = \frac{1}{k(x_1, x_1)}$ . Drawing a second variable,  $x_2$ , results in a covariance matrix

$$C_2 = \begin{pmatrix} k(x_1, x_1) & k(x_1, x_2) \\ k(x_1, x_2) & k(x_2, x_2) \end{pmatrix}$$

Since  $C_2$  is a covariance matrix, it must be symmetric, so  $k(x_1, x_2) = k(x_2, x_1)$ , and its inverse is given by

$$C_2^{-1} = \frac{1}{k(x_1,x_1)k(x_2,x_2) - k(x_1,x_2)^2} \left( \begin{array}{cc} k(x_2,x_2) & -k(x_1,x_2) \\ -k(x_1,x_2) & k(x_1,x_1) \end{array} \right)$$

Since the first element of  $C_2^{-1} = p(x_1, x_1)$ , this implies

$$p(x_1, x_1) = \frac{k(x_2, x_2)}{k(x_1, x_1)k(x_2, x_2) - k(x_1, x_2)^2}$$

Plugging in  $p(x_1, x_1) = \frac{1}{k(x_1, x_1)}$  from above, this turns into

$$\frac{1}{k(x_1,x_1)} = \frac{k(x_2,x_2)}{k(x_1,x_1)k(x_2,x_2) - k(x_1,x_2)^2}$$

Rearranging, this becomes,

$$k(x_1, x_1)k(x_2, x_2) = k(x_1, x_1)k(x_2, x_2) - k(x_1, x_2)^2$$

which implies

$$k(x_1, x_2) = k(x_2, x_1) = 0$$

This means that  $p(x_i, x_j)$  only exists if C is a diagonal matrix. Since C is not required to be diagonal, this implies that  $p(x_i, x_j)$  does not exist generally.

### 2.2 Question 4

I collaborated with Peter Li on this problem.

I've graphed four covariance functions:

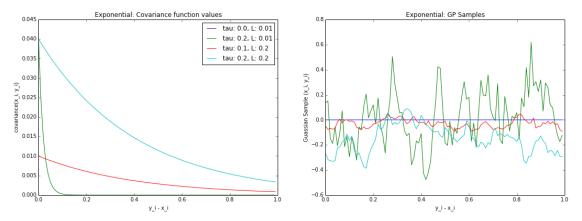
- 1) Exponential (not differentiable, not compact)
- 2) Squared Exponential (differentiable, not compact)
- 3) Spherical (not differentiable, compact)
- 4) Linear (differentiable, not compact)

```
In [435]: # Covariance functions
          import itertools
          import math
          import matplotlib.pyplot as plt
          import numpy as np
          from mpl_toolkits.mplot3d import Axes3D
          from matplotlib import cm
          %matplotlib inline
          # 1) Exponential (not differentiable, not compact)
          def make_exp_covar_fn(tau, 1):
              def exp_covar_fn(x, y):
                  return tau**2 * math.exp(-np.linalg.norm(x - y) / (2*1))
              return exp_covar_fn
          # 2) Squared Exponential (differentiable, not compact)
          def make_squared_exp_covar_fn(tau, 1):
              def squared_exp_covar_fn(x, y):
                  diff = x - y
                  return tau**2 * math.exp(-np.inner(diff, diff) / (2*(1**2)))
              return squared_exp_covar_fn
          # 3) Spherical (not differentiable, compact)
          def make_spherical_covar_fn(tau, theta):
              def spherical_covar_fn(x, y):
                  dist = np.linalg.norm(x - y)
                  if dist <= theta:
                      return tau**2 * (1 - 3*dist/(2*theta) + dist**3/(2*theta**3))
                  else:
                      return 0.0
              return spherical_covar_fn
          # 4) Linear (differentiable, not compact)
```

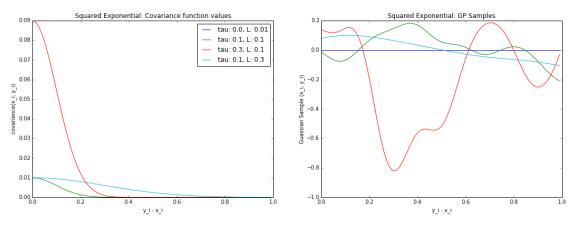
```
def make_linear_covar_fn(sigma, tau, c):
              def linear_covar_fn(x, y):
                  return sigma**2 + tau**2 * np.inner(x - c, y - c)
              return linear_covar_fn
In [317]: # This samples from a multivariate Gaussian distribution, with a zero mean
          # value, and a covariance matrix determined by the convariance function
          # and the x values parameters.
          def sample_gp(xs, covar_fn):
              mean_vec = np.zeros(len(xs))
              gram_matrix = np.zeros((len(xs), len(xs)))
              for i in range(len(xs)):
                  x_i = xs[i]
                  for j in range(0, i):
                      x_j = xs[j]
                      gram_matrix[i][j] = covar_fn(x_i, x_j)
                      gram_matrix[j][i] = gram_matrix[i][j]
                  gram_matrix[i][i] = covar_fn(x_i, x_i)
              samples = np.random.multivariate_normal(mean_vec, gram_matrix)
              return samples
In [491]: def plot_covar_fn(title, covar_fn_maker, param_list, param_names):
              covar_fn_list = [covar_fn_maker(*params) for params in param_list]
              ys = np.arange(0.0, 1.0, 0.01)
              covar_vals = []
              sample_vals = []
              for covar_fn in covar_fn_list:
                  covar_vals.append([covar_fn(0.0, y) for y in ys])
                  sample_vals.append(sample_gp(ys, covar_fn))
              fig = plt.figure(1, figsize=(18, 6))
              # Covariance values
              plt.subplot(1, 2, 1)
              plt.title(title + ': Covariance function values')
              plt.xlabel('y_i - x_i')
              plt.ylabel('covariance(x_i, y_i)')
              lines = []
              for params, covar_val in zip(param_list, covar_vals):
                  param_str = ", ".join([name + ": " + str(param)
                                         for name, param in zip(param_names, params)])
                  line, = plt.plot(ys, covar_val)
                  lines.append(param_str)
              plt.legend(lines)
              # Gaussian Process samples
              plt.subplot(1, 2, 2)
              plt.title(title + ': GP Samples')
              for sample_val in sample_vals:
                  plt.plot(ys, sample_val)
              plt.xlabel('y_i - x_i')
              plt.ylabel('Guassian Sample (x_i, y_i)')
              plt.show()
In [492]: # L can't be zero or you get a divide by zero error
```

```
exp_params = [(0.0, 0.01), (0.2, 0.01), (0.1, 0.2), (0.2, 0.2)]
plot_covar_fn("Exponential", make_exp_covar_fn, exp_params, ['tau', 'L'])
```

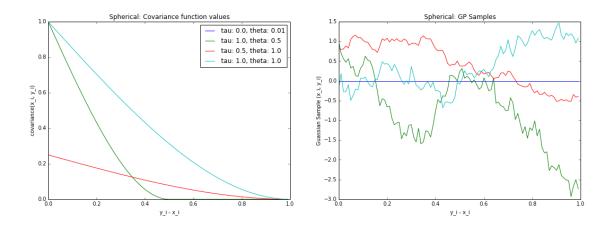
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/Library/Python/2.7/site-packages/ipykernel/\_main\_.py:13: DeprecationWarning: using a non-integer number



/Library/Python/2.7/site-packages/ipykernel/\_main\_..py:13: DeprecationWarning: using a non-integer number



/Library/Python/2.7/site-packages/ipykernel/\_main\_.py:13: DeprecationWarning: using a non-integer number

