Report for Lab 2-Part 1

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_get_jt_clique_and_edges()

In this function, clique sets and their edges with each other are calculated. To be more detailed, nx.Graph() is used to represent the original graph. Then, we can get jt_cliques via nx.find_cliques().

According to these cliques, we can build a clique graph, and here I add a helper function build_clique_graph_and_weight(jt_cliques). The clique sets are needed as its input and nx.Graph() represents the clique graph, where cliques' indexes are nodes, and if 2 cliques have intersections, an edge between them will be added to the graph, with the size of their intersection being the edge's weight.

After getting the clique graph, nx.maximum_spanning_tree() is used to get the maximum spanning tree, then we can get jt_edges from this maximum spanning tree.

_get_clique_factors()

In this function, we get clique nodes' factors by using factor_product() and iteration.

_update_mrf_w_evidence()

In this function, <code>updated_factors</code> are obtained by using <code>factor_evidence()</code> and iteration.

<code>query_nodes</code> are obtained by removing observed nodes from factor sets. <code>updated_edges</code> are obtained by removing edges related to observed nodes.

_get_clique_potentials()

In this function, the sum-product algorithm is used to get clique potentials. nx.Graph() is used to represent clique graph. Four helper functions:

collect(g,i,j,msg,cliques), sendmessage(g,j,i,msg,cliques), distribute(g,i,j,msg,cliques)
and computeMarginal(g,i,msg) are implemented like Lab 1.

_get_node_marginal_probabilities()

In this function, we get the marginal probability of each query node by marginal the clique potentials. Considering a node may appear in multiple cliques of different sizes, we can first find the smallest clique related to it and then marginalize other nodes, which can avoid extra computation.