```
function [LL, W, s2] = ppca_nsp(X, D)
% This function was written by Jonathan Kao
  for use in EE239AS.2, UCLA.
%
% Inputs:
% - X:
         N x K matrix, where N is the dimensionality of the data and K is
% the number of data points.
% - D: the latent dimensionality
% Outputs:
% - LL:
         the log-likelihoods over all iterations
   - W:
         the W parameter of PPCA
   - s2: the sigma^2 parameter of PPCA
         the mu parameter of PPCA
   % Check for convergence (tol) or set a max number of cycles (cyc)
   tol = 1e-8;
   cyc = 1e8;
   % Set the random seed so everyone's code executes the same thing.
   randn('state', 0);
   [N, K] = size(X);
   % Initialization ofparameters
   cX = cov(X', 1);
      = randn(N,D);
   mu = mean(X, 2);
   LL = [];
   LL_prev
             = -Inf;
   % do a maximum of cyc cycles
   for i = 1:cyc
       % E-STEP -- compute E[s], E[ss^T]
       % Denote E[s] as:
       % Denote E[ss^T] as: Ess
       %%%%% START YOUR CODE HERE TO CALCULATE Es and Ess %%%%%
       mu mat = repmat(mu,1,K);
       % identity matrix needs to be N x N to match dimensions of W*W'
       Es = W'*inv(W*W' + s2*eye(N))*(X-mu_mat);
       Ess = eye(D) - W'*inv(W*W' + s2*eye(N))*W + (1/K)*Es*Es';
       % Ess = cov(s_k) + Es * Es'
       %%%% END YOUR CODE HERE TO CALCULATE Es and Ess %%%%
```

```
% Compute log likelihood
       % ============
       % We'll do this one for you.
              = W*W' + s2 * eye(N);
              = bsxfun(@minus, X, mu);
       Xm
               = -N*K/2*log(2*pi) - K/2*log(det(Sx))-1/2*trace(Xm' * inv(Sx) * Xm')
       LLi
       T.T.
               = [LL LLi];
       % ===========
       % M-STEP -- update W, s2
       %%%%% START YOUR CODE HERE TO CALCULATE W and s2 %%%%%
       W = (X-mu_mat)*Es'*(1/K)*inv(Ess');
       s2 = 1/(N*K) * trace((X-mu_mat)*(X-mu_mat)' - W*Es*(X-mu_mat)');
       %%%% END YOUR CODE HERE TO CALCULATE W and s2 %%%%%
       % ===========
       % Check for convergence
       % First iteration, set the base likelihood that we'll measure
       % comparisons vs.
       if i == 1
           LLbase = LLi;
       end
       if LLi < LL_prev</pre>
           disp('LIKELIHOOD VIOLATION!');
       elseif ((LLi-LLbase) < (1+tol)*(LL_prev-LLbase))</pre>
           break;
       end
       LL_prev = LLi;
   end
end
Error using ppca_nsp (line 23)
Not enough input arguments.
```

% =============

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