
```

R_exact = readmatrix('Nodes1thru57_dist_exact.txt');
S_exact = R_exact.^2;
N = size(R_exact, 1);
[~,~,G_clean] = algo_1(R_exact, S_exact, N);

R_rounded = readmatrix('Nodes1thru57_dist.txt');
S_rounded = R_rounded.^2;
[~,~,G_noisy] = algo_1(R_rounded, S_rounded, N);

names = ['Noisy' + string(1:10) + 'Nodes1to57kn57_dist', string(1:10)
+ 'kn57Nodes1to57_exactdist'];
graphs = size(names, 2);
clean_errors = zeros([1, graphs]);
noisy_errors = zeros([1, graphs]);
epsilons = zeros([1, graphs]);
for j=1:graphs
    % Putting 2048 here because it's somewhat close to the epsilon
    % values that CVX deems 'good'
    [clean_errors(j), noisy_errors(j), epsilons(j)] = get_error(names(j),
    G_noisy, G_clean, 4096);
end

figure
hold on
plot(epsilons(1:10), 'ro-')
plot(epsilons(11:20), 'bo-')
title('Epsilon Values for Noisy and Clean data')
xlabel('Tenths of data given')
ylabel('Epsilon value')
legend(['Noisy Values'; 'Clean Values'])
hold off

figure
hold on
plot(noisy_errors(1:10), 'ro-')
plot(clean_errors(11:20), 'bo-')
title('Error Values for Noisy and Clean data')
xlabel('Tenths of data given')
ylabel('Error compared to Actual Gram')
legend(['Noisy Values'; 'Clean Values'])
hold off

m_list = 1./(((1:10)/10) * N*(N-1)/2);
b_noisy = (m_list' * m_list) \ (m_list' * noisy_errors(1:10));
b_clean = (m_list' * m_list) \ (m_list' * clean_errors(11:20));
figure
hold on
loglog(noisy_errors(1:10), 'ro-')
loglog(clean_errors(11:20), 'bo-')
axis tight

```

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title('Error Values for Noisy and Clean data')
xlabel('Tenths of data given')
ylabel('Log-scaled error')
legend(['Noisy Values'; 'Clean Values'])
hold off

```

```

figure
hold on
semilogx(noisy_errors(1:10), 'ro-')
semilogx(clean_errors(11:20), 'bo-')
axis tight
title('Error Values for Noisy and Clean data')
xlabel('Tenths of data given')
ylabel('SemiLog-scaled error')
legend(['Noisy Values'; 'Clean Values'])
hold off

```

```

% The epsilon looks like it has an error rate similar to  $1/m^\alpha$  instead of
%  $e^{-\beta m}$  as the rate seems to be significantly higher at  $X=1$  than at  $X=2$ .
% Epsilon seems to make a significant difference; when more and more
% restrictions are placed on the program it makes sense that it becomes harder
% to fully conform to every restriction. When epsilon is much larger, it
% becomes far more feasible.
% The feasibility vs accuracy tradeoff is a massive one. As mentioned before,
% the fact that epsilon has such a large difference on the program is evident
% given the disparity between epsilons where we've obtained results.
% When epsilon is low, the program has a more difficult time finding a
% solution; that is, the program determines that a solution would not be found
% in any reasonable amount of time.

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function [Error_clean, Error_noisy, epsilon] = get_error(name, G_noisy,
    G_clean, init_epsilon)
[N, ~, ~, E_ijs, D_tilde] = prep_distance('SparseGraphs\Sparse' + name
    + '.txt');

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G = NaN(N);
epsilon = init_epsilon / 2;

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while (isnan(G))
    epsilon = epsilon * 2;
    % TODO: FIX THIS PART FOR LATER
    % Want to get down the rest of the code before coming back
    G = perform_cvx(N, D_tilde, E_ijs, epsilon);
end

```

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Error_clean = norm(G - G_clean, 'fro');
Error_noisy = norm(G - G_noisy, 'fro');
end

```

```

function G = perform_cvx(N, D_tilde, E_ijs, epsilon)
m = size(D_tilde, 1);
cvx_begin sdp quiet
variable G(N,N) semidefinite symmetric;

```

```

minimize trace(G);
subject to
G*ones(N, 1) == 0;
% We're looking to minimize the inner product of Ge_ij and e_ij and
% the random d distances
% e_ij = 1 at i, -1 at j, 0 else
% E_ijs should be this M times for each possible edge

% Need to make sure it only counts constraints that make sense
% That is, we need to ensure it only counts constraints that we have data
% for
% Otherwise, we get that the distance between two points has to be zero (or
% close enough to zero as to be zero)

% By doing this, we're making sure that we're only getting the distances
% that are non-zero
abs(diag(E_ijs'*G*E_ijs) - D_tilde') <= epsilon * ones(m, 1);

cvx_end
end

function [E, D_tilde] = fetch_vals(N, D)
% First get the number of possible edges
m = (N*N - N) / 2;
% Initialize E to all zeros
E = zeros([N, m]);
% Create a variable to store the distances
D_tilde = zeros([1, m]);
j = 1;
index = 1;

% Do it all with a single for loop (makes D easier)
for k=1:m
    % Add 1 to J and loop if it's at N + 1
    j = mod(j + 1, N + 1);
    % If it's zero, we've hit the end of a column
    if (j == 0)
        index = index + 1;
        j = 1;
    end

    % We don't really want all the duplicate i-j pairs, so remove those by
    % only making the first pair of i-j matter (instead of j-i)
    % This way, we only get the right number of i-j
    while (j <= index)
        j = mod(j + 1, N + 1);
    end
    % At the end, we then fill the values we want to fill
    % i is 1 and j is 0
    E(index, k) = 1;
    E(j, k) = -1;

    % Then, we assign the right distance from our data to the distance
    % array

```

```

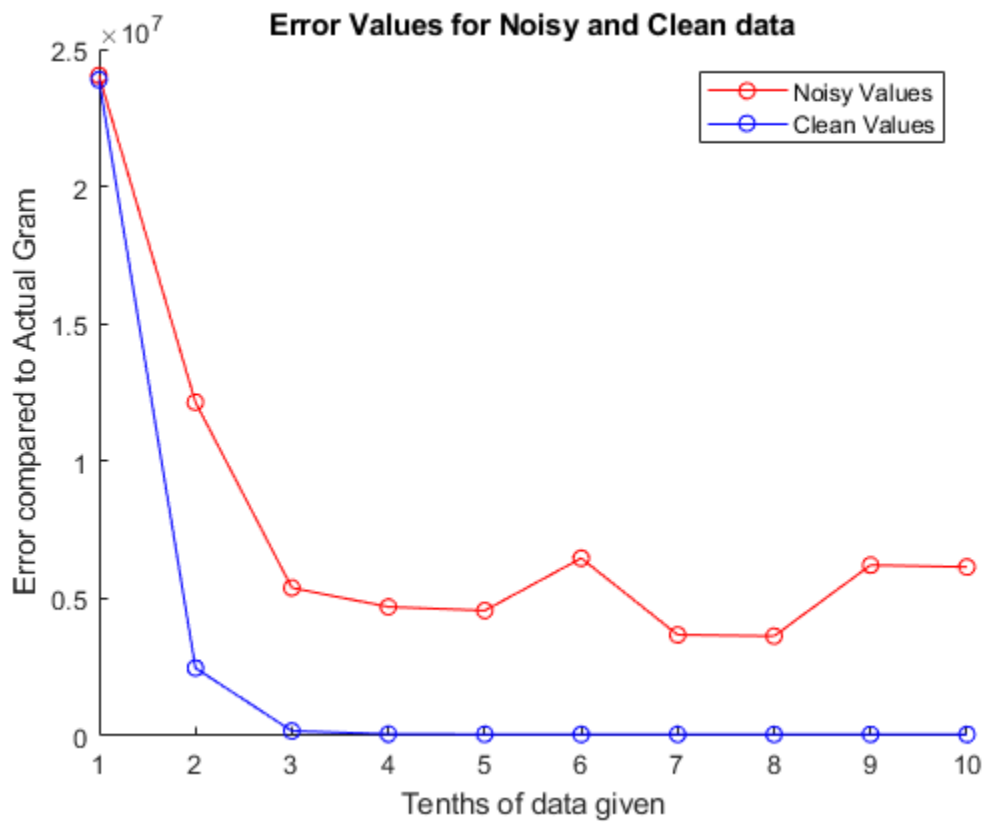
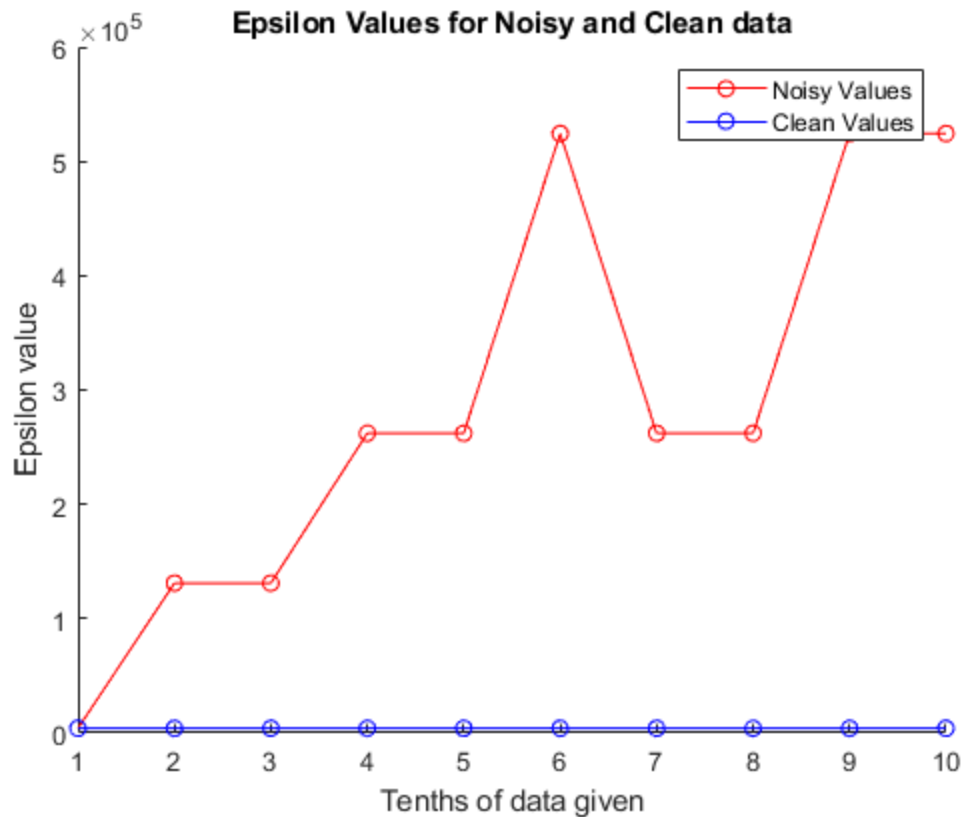
        D_tilde(k) = D(j, index);
end
end

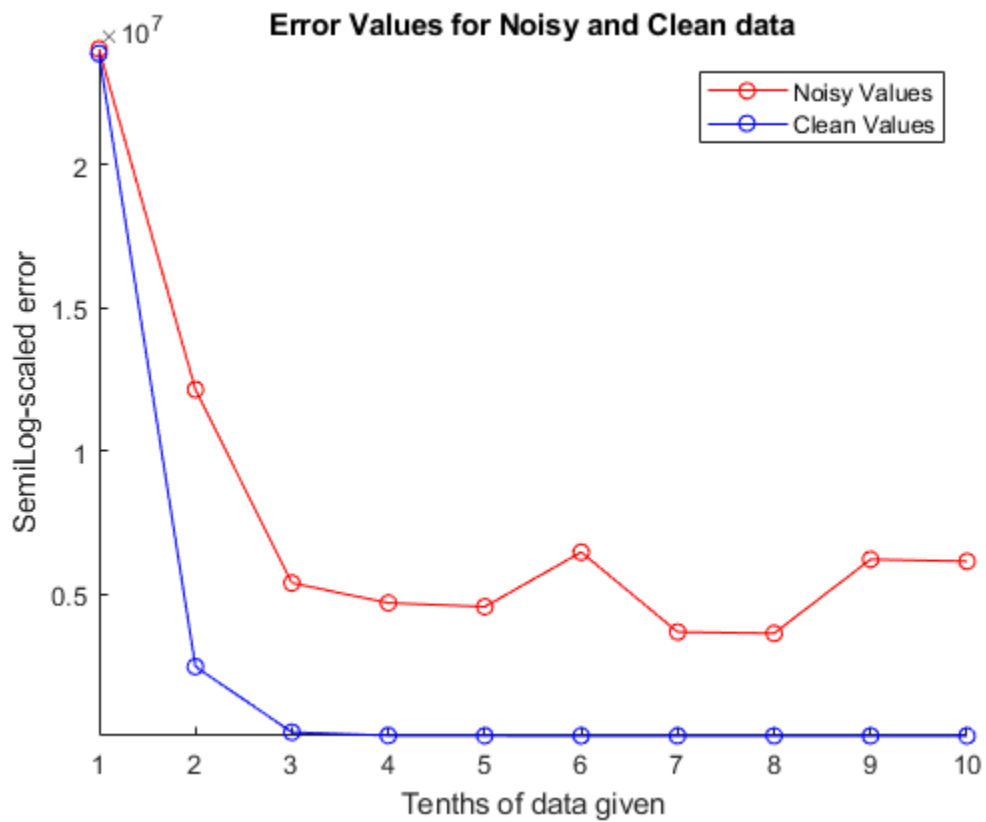
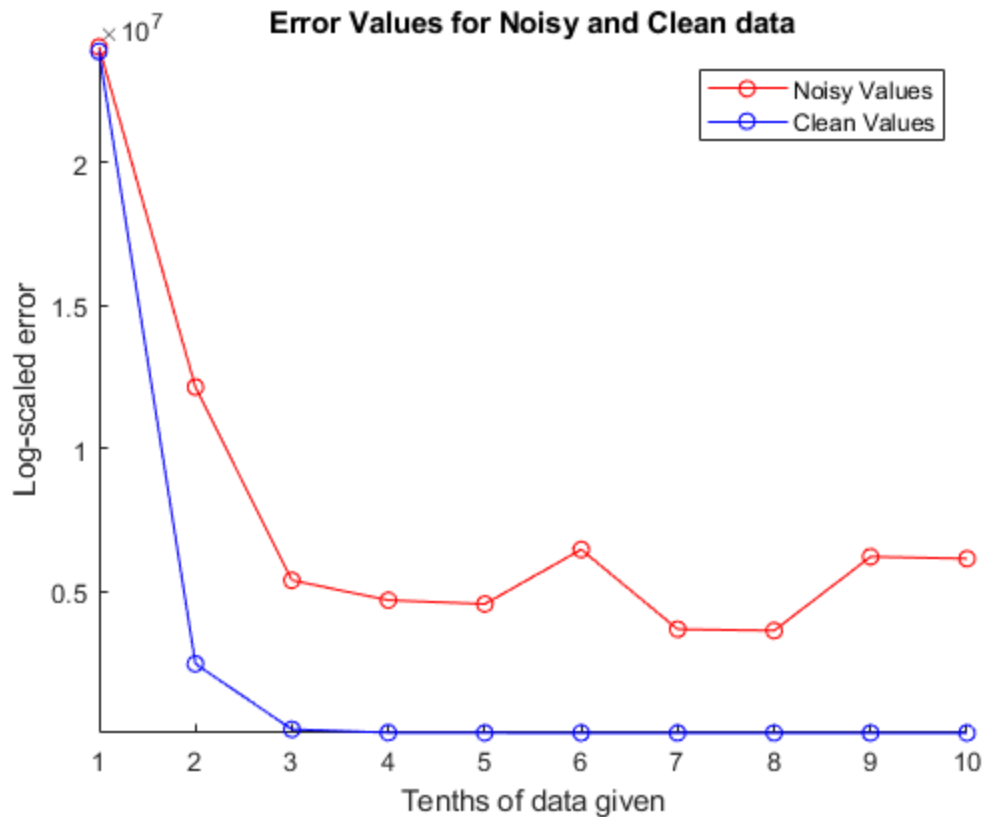
% Below are setup functions we don't really need to worry about.
function [N, M, arr] = read_to_arr(filename)
T = readtable(filename);
N = table2array(T(1, 1));
M = table2array(T(1, 2));
arr = table2array(T(2:M+1, :));
end

% Initially assign D to zero to allocate space and to make it easy to tell
% when a point has no value given (0 distances!??)
function [N, M, D, E, y] = prep_distance(filename)
[N, M, arr] = read_to_arr(filename);
D = zeros(N);
for index=1:M
    point = arr(index, 1:2);
    D(point(1), point(2)) = arr(index, 3);
    E(point(1), index) = 1;
    E(point(2), index) = -1;
    y(index) = arr(index, 3)^2;
end
D = D + D';
end

function [rho, upsilon, G] = algo_1(R, S, n)
% Use Algorithm 1 to create Gram G
rho = (1/(2 * n)) * sum(sum(S));
upsilon = (1/n) * (S - rho*eye(n)) * ones(n, 1);
G = 0.5*upsilon*ones(n, 1)' + 0.5*ones(n, 1)*upsilon' - 0.5*S;
end

```





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