

MSc Project IIS1 | BWFLnet_Water Analysis

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```
clc  
clear  
close all
```

Model Setup and Initialisation

Initialisation of EPANET Matlab toolkit and loading of .net file.

```
epanet_path = 'E:\Program Files\MATLAB\R2021b\toolbox\epanet\EPANET-Matlab-Toolkit-master';  
net_id = 'BWFLnet_MSc_2022_calibrated';  
run([epanet_path, '\start_toolkit'])
```

EPANET-MATLAB Toolkit Paths Loaded.

```
net = epanet([net_id, '.inp']);
```

```
EPANET version {20200} loaded (EMT version {v2.2.0}).  
Loading File "BWFLnet_MSc_2022_calibrated.inp"...  
Input File "BWFLnet_MSc_2022_calibrated.inp" loaded sucessfuly.
```

```
load("wq_data.mat");  
load("R_CData.mat");
```

Network Data Loading

Load general network data, noting that this network has already been hydraulically calibrated, thus we can directly perform the quality analysis.

```
% Save element count variables  
nn = net.NodeJunctionCount;  
n0 = net.NodeReservoirCount;  
np = net.LinkCount; % count of pipes and valves  
D = net.getLinkDiameter';  
R = net.getLinkRoughnessCoeff';  
  
% Save node and link index vectors  
Reservoir_Idx = net.getNodeReservoirIndex;  
Junction_Idx = net.getNodeJunctionIndex;  
Link_Idx = double(net.getLinkIndex);  
  
% Specify hydraulic and quality time steps (in seconds)  
net.setTimeQualityStep(15*60); % 15 minute quality time steps  
net.setTimeHydraulicStep(15*60); % 15 minute hydraulic time steps  
nt = net.getTimeSimulationDuration./3600; % number of time steps in hours  
days = 7; % simulation duration in days  
net.setTimeSimulationDuration(nt*3600*days/7); % Set the simulation duration by changing days  
nt = net.getTimeSimulationDuration./3600; % get the new nt  
  
% Nodes name and chlorine concentrations
```

```

Junction_All_Name = {'BW9','BW7','BW1','BW2','BW3','BW4','BW5','BW6','BW12'};
Junction_SelectIdx = net.getNodeIndex(wq_data.node_ids([1,4,7,9]));
Junction_Name = Junction_All_Name([1,4,7,9]);
Junction_HydrantIdx = net.getNodeIndex(wq_data.node_ids([2,5,8]));
Junction_Hydrant_Name = Junction_All_Name([2,5,8]);
Junction_Reservoir_Name = Junction_All_Name([6,3]);
cext = wq_data.chlorine([6,3],:);
c_nodes_HydrantObs = wq_data.chlorine([2,5,8],:);
c_nodes_Obs = wq_data.chlorine([1,4,7,9],:);

```

Graph Creation Preparation

Use graph theory to plot network connectivity and spatial coordinates.

```

% Create A12 and A10 incidence matrices
LinkNodesList = net.getLinkNodesIndex;

A = zeros(np,nn+n0);
for k=1:np
    i = LinkNodesList(k,1);
    j = LinkNodesList(k,2);
    A(k,i) = -1;
    A(k,j) = 1;
end

A12 = A(:,Junction_Idx);
A10 = A(:,Reservoir_Idx);
A12 = sparse(A12);
A10 = sparse(A10);

% Obtain node XY and elevation information
XY = zeros(nn+n0,2);
XY(:,1) = net.getNodeCoordinates{1};
XY(:,2) = net.getNodeCoordinates{2};
elev = double(net.getNodeElevations(Junction_Idx))';

% Create adjacency matrix
A = [A12,A10];
AdjA = sparse(size(A,2),size(A,2));

for k = 1:size(A,1)
    node_in = find(A(k,:) == -1);
    node_out = find(A(k,:) == 1);
    AdjA(node_in,node_out) = 1;
    AdjA(node_out,node_in) = 1;
end
gr = graph(AdjA);

```

Graph Plotting

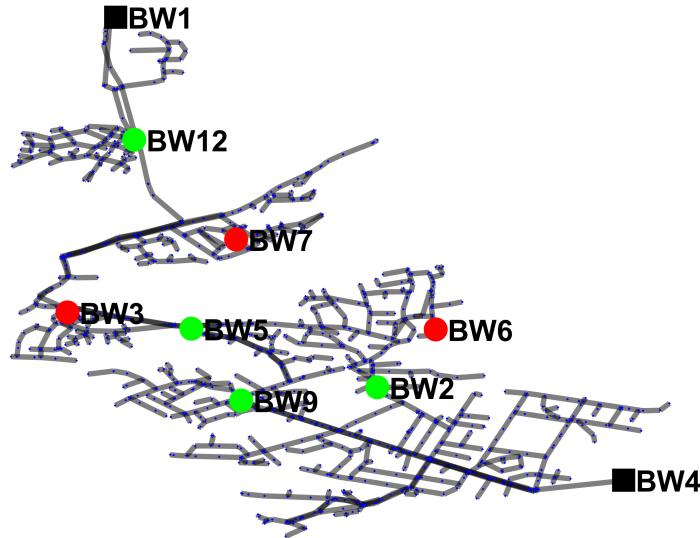
Setup general chemical simulation parameters.

```

% Plot the sensor location on the network map
figure
p1 = plot(gr);
p1.XData = XY(:,1);
p1.YData = XY(:,2);
p1.LineWidth = 2;
p1.EdgeColor = 'k';
p1.MarkerSize = 0.01;
p1.NodeColor = 'b';
p1.NodeLabel = '';
highlight(p1,Junction_SelectIdx,'NodeColor','green','MarkerSize',8);
labelnode(p1,Junction_SelectIdx,Junction_Name);
highlight(p1,Junction_HydrantIdx,'NodeColor','red','MarkerSize',8);
labelnode(p1,Junction_HydrantIdx,Junction_Hydrant_Name);
highlight(p1,Reservoir_Idx,'NodeColor','k','Marker','s','MarkerSize',10);
labelnode(p1,Reservoir_Idx,Junction_Reservoir_Name);
p1.NodeFontSize = 11;
p1.NodeLabelColor = 'k';
p1.NodeFontWeight = 'bold';
p1.EdgeFontSize = 11;
p1.EdgeLabelColor = 'k';
p1.EdgeFontWeight = 'bold';
% p1.EdgeCData = R_CData;
% hcb_R = colorbar;
% colormap('jet');
% hcb_R.Label.String = 'Pipe Roughness Group';
% hcb_R.Label.FontSize = 12;
axis('off')
title('Sensor Locations in Network')

```

Sensor Locations in Network



```
% Plot the Roughness group on the network map
figure
p1 = plot(gr);
p1.XData = XY(:,1);
p1.YData = XY(:,2);
p1.LineWidth = 3;
p1.EdgeColor = 'k';
p1.MarkerSize = 1e-10;

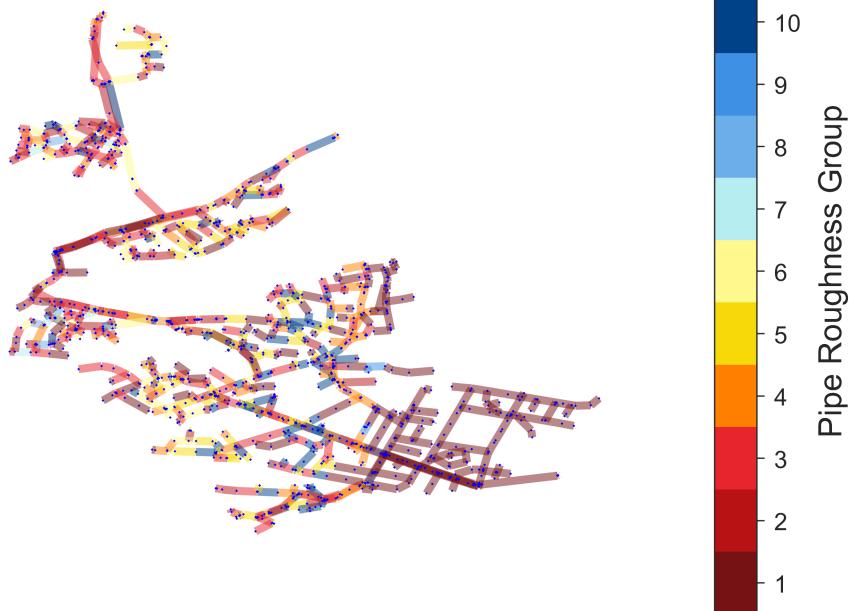
p1.NodeColor = 'b';
p1.NodeLabel = '';
p1.NodeFontSize = 11;
p1.NodeLabelColor = 'k';
p1.NodeFontWeight = 'bold';
p1.EdgeFontSize = 11;
p1.EdgeLabelColor = 'k';
p1.EdgeFontWeight = 'bold';
p1.EdgeCData = R_CData;
cmap=[118 18 19;
      184 18 21;
      230 38 44;
      255 127 0;
      247 217 9;
      255 248 140;
      182 237 240;
      107 174 232;
      61 144 227];
```

```

0 66 135;
0 31 103];
cmap=cmap/255;
colormap(cmap);
hcb = colorbar;
hcb.TicksMode = 'manual';
set(hcb,'Tickdir','out');
caxis([1,11]);
set(hcb,'YTick',1+5/11:10/11:11-5/11);
set(hcb,'TickLabels',{'1','2','3','4','5','6','7','8','9','10','11'});
hcb.Label.String = 'Pipe Roughness Group';
hcb.Label.FontSize = 12;
axis('off')
title('Roughness-based groups in Network')

```

Roughness-based groups in Network



Chlorine Simulation

Setup general chemical simulation parameters.

```

% Initialise EPANET simulation type
net.setQualityType('Chemical','mg/L');
net.setNodeSourceQuality(1:nn+n0,zeros(nn+n0,1)); % set all node source quality to zero

% % Link bulk reaction coefficients
% lambda = 0.5*ones(np,1); % units of days^-1
% net.setLinkBulkReactionCoeff(1:np, -lambda);
% net.setOptionsPipeBulkReactionOrder(1);

```

```
% Initial concentrations at nodes (mg/L)
c0 = zeros(nn+n0,nt);
net.setNodeInitialQuality(net.NodeIndex,c0);
base_cext = ones(n0,1);
pattern_cext = cext./(base_cext*ones(1,size(cext,2))); % extend vector over nt columns

% For loop to assign new patterns to source contrations at reservoirs and
% source type
for i=1:n0
    patternId = sprintf('Res_C_%d',i);
    net.addPattern(patternId,pattern_cext(i,:));
    net.setNodeSourcePatternIndex(net.NodeReservoirIndex(i),net.getPatternIndex(patternId));
    net.setNodeSourceQuality(net.NodeReservoirIndex(i),base_cext(i));
    net.setNodeSourceType(net.NodeReservoirIndex(i),'CONCEN');
end
```

Results from EPANET Simulation

Simulate hydraulic and quality analyses using EPANET's solvers.

```
hydraulic_res = net.getComputedHydraulicTimeSeries;
quality_res = net.getComputedQualityTimeSeries;
```

```
% Assign hydraulic results to network elements
h = hydraulic_res.Head(1:1+4*nt,1:nn).';
q = 1e-3*hydraulic_res.Flow(1:1+4*nt,:).';

% Assign quality results to network elements
c_nodes = quality_res.NodeQuality';
c_pipes = quality_res.LinkQuality';

% Get the simulated and observed data
c_nodes_Sim = c_nodes(Junction_SelectIdx,:);
c_nodes_Sim = c_nodes_Sim(:,1:end-1);
c_nodes_HydrantSim = c_nodes(Junction_HydrantIdx,:);
c_nodes_HydrantSim = c_nodes_HydrantSim(:,1:end-1);
c_nodes_Reservoir = pattern_cext;

% split the data
Sim_train = c_nodes_Sim(:,2*96+1:3*96);
Sim_val = c_nodes_Sim(:,3*96+1:end);
Sim_Hydrant_train = c_nodes_HydrantSim(:,2*96+1:3*96);
Sim_Hydrant_val = c_nodes_HydrantSim(:,3*96+1:end);

Obs_train = c_nodes_Obs(:,2*96+1:3*96);
Obs_val = c_nodes_Obs(:,3*96+1:end);
Obs_Hydrant_train = c_nodes_HydrantObs(:,2*96+1:3*96);
Obs_Hydrant_val = c_nodes_HydrantObs(:,3*96+1:end);

% Combine the two data sets
Junction_CombinedIdx = [Junction_SelectIdx,Junction_HydrantIdx];
Obs_Combined_train = [Obs_train; Obs_Hydrant_train];
```

```

Obs_Combined_val = [Obs_val; Obs_Hydrant_val];
Sim_Combined_train = [Sim_train; Sim_Hydrant_train];
Sim_Combined_val = [Sim_val; Sim_Hydrant_val];

% Calculate the original mse
mse_function = @(sim,obs) (1/size(obs,1)/size(obs,2))*sum(sum((sim - obs).^2));
mse_before_train = mse_function(Sim_Combined_train, Obs_Combined_train);
mse_before = mse_function(Sim_Combined_val, Obs_Combined_val);

```

GA calibration

Assign an uniform bulk decay coefficient to the whole network based on literatures.

```

% bulk reaction coefficient
lambda = 1.0*ones(np,1); % units of days^-1
net.setLinkBulkReactionCoeff(1:np, -lambda);
net.setOptionsPipeBulkReactionOrder(1);

```

Single wall decay coefficient GA calibration

Start the GA process with one single wall decay coefficient, regardless of the pipe roughness and diameter.

```

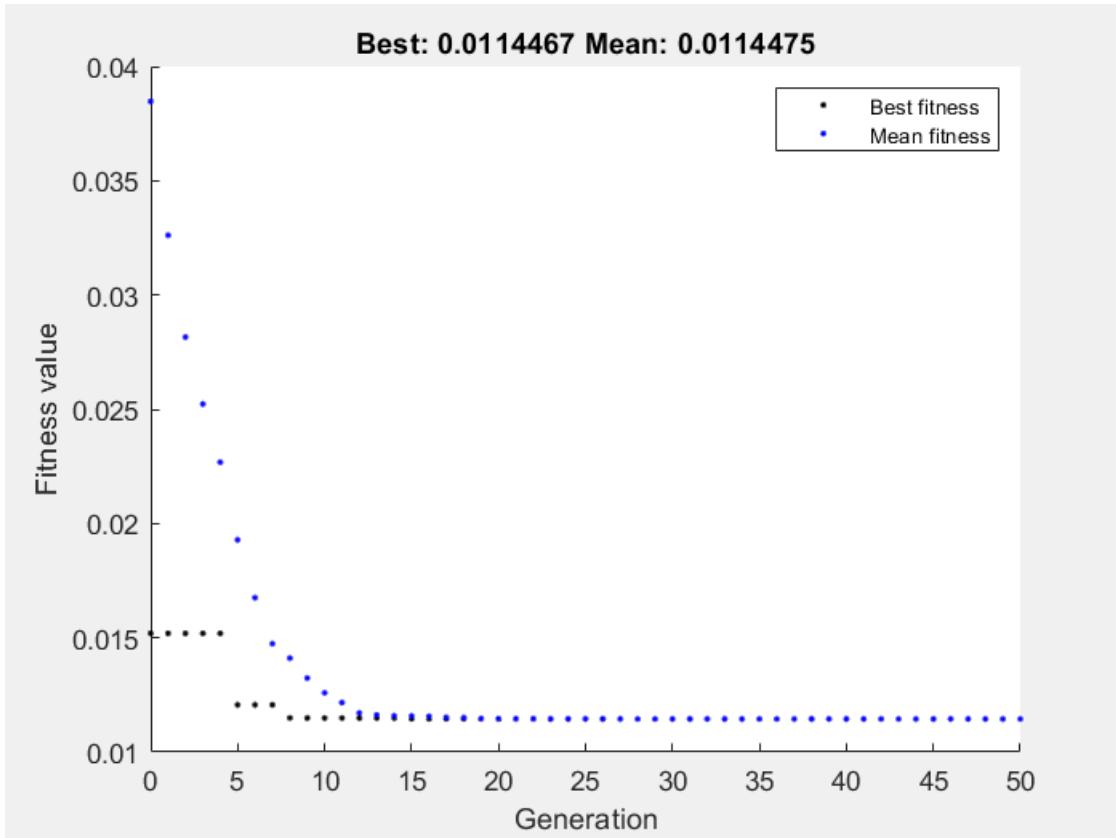
% Consider the whole network together and assign the same decay parameter

rng default
options = optimoptions('ga','Display','none','PopulationType','DoubleVector','PopulationSize',50000,'Generations',50,"ConstraintTolerance",1e-6,"PlotFcn","gaplotbestf");

lb = -10;
ub = 0;

tic;
Mse_function = @(x) Mse_NoGroup(x,Obs_Combined_train,Junction_CombinedIdx);
[x_NoGroup, fval_NoGroup] = ga(Mse_function,1,[],[],[],lb,ub,[],options);

```



```
t_NoGroup = toc;
fprintf('The computation time is %.2fs',t_NoGroup);
```

The computation time is 729.16s

```
% Test the ga result
theta_NoGroup = ones(2281,1); % units of m/day
for i = 1:2281
    theta_NoGroup(i) = x_NoGroup(1);
end
net.setLinkWallReactionCoeff(1:np, theta_NoGroup);
net.setOptionsPipeWallReactionOrder(1);
quality_res_NoGroup = net.getComputedQualityTimeSeries;

c_nodes_NoGroup = quality_res_NoGroup.NodeQuality';
c_nodes_Sim_NoGroup = c_nodes_NoGroup(Junction_SelectIdx,1:end-1);
c_nodes_Sim_Hydrant_NoGroup = c_nodes_NoGroup(Junction_HydrantIdx,1:end-1);
c_nodes_Sim_Combined_NoGroup = [c_nodes_Sim_NoGroup;c_nodes_Sim_Hydrant_NoGroup];

Sim_Combined_NoGroup_val = c_nodes_Sim_Combined_NoGroup(:,3*96+1:end);
mse_after_NoGroup = mse_function(Sim_Combined_NoGroup_val,Obs_Combined_val);

Sim_main_NoGroup_val = c_nodes_Sim_NoGroup(:,3*96+1:end);
mse_main_after_NoGroup = mse_function(Sim_main_NoGroup_val,Obs_val);
fprintf('The validation mse of nodes in the main network after calibration is %.4f',mse_main_at
```

The validation mse of nodes in the main network after calibration is 0.0280

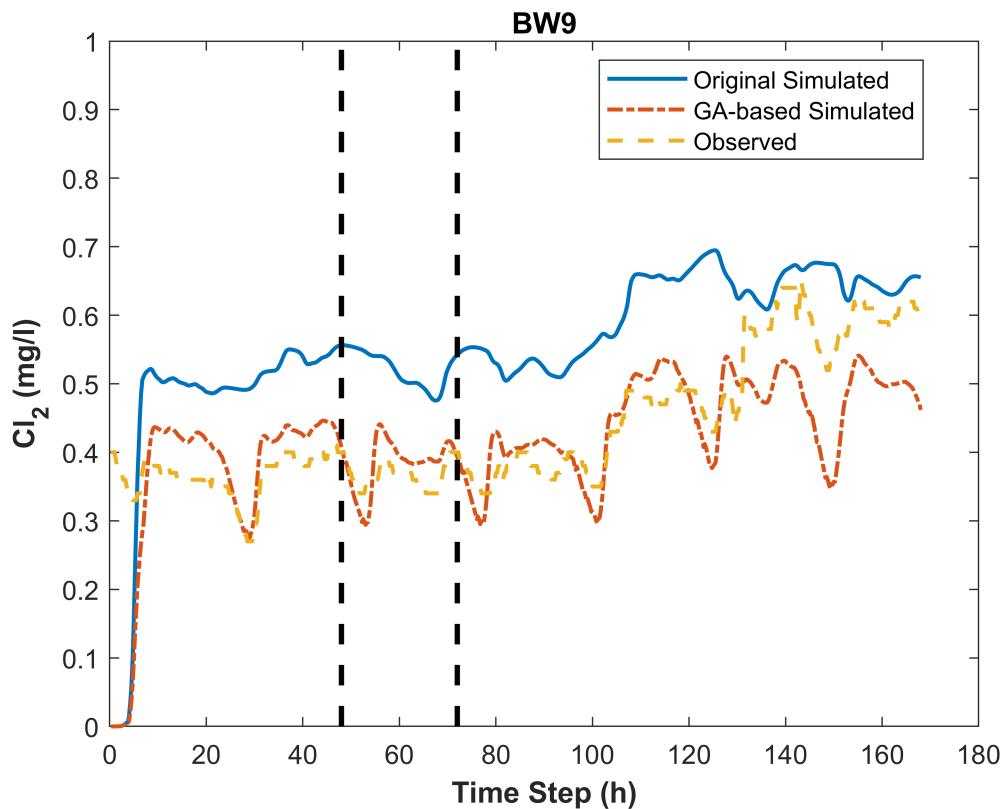
```
fprintf('The train mse before calibration is %.4f, after calibration it is %.4f',mse_before_tr
```

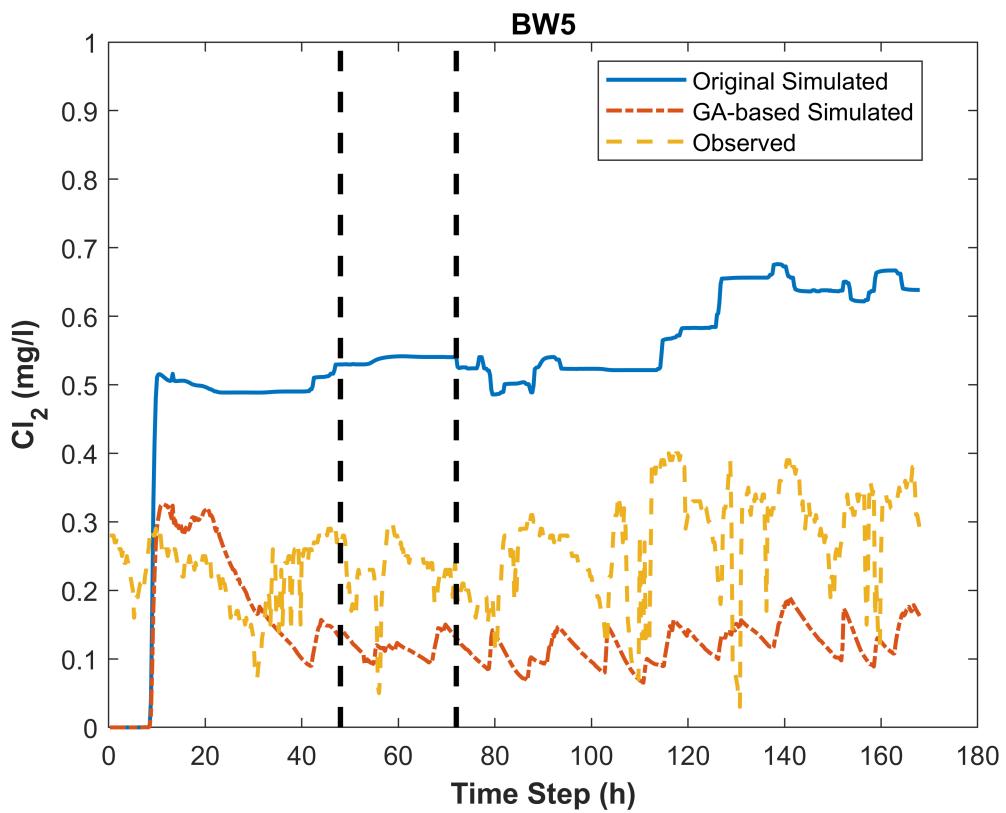
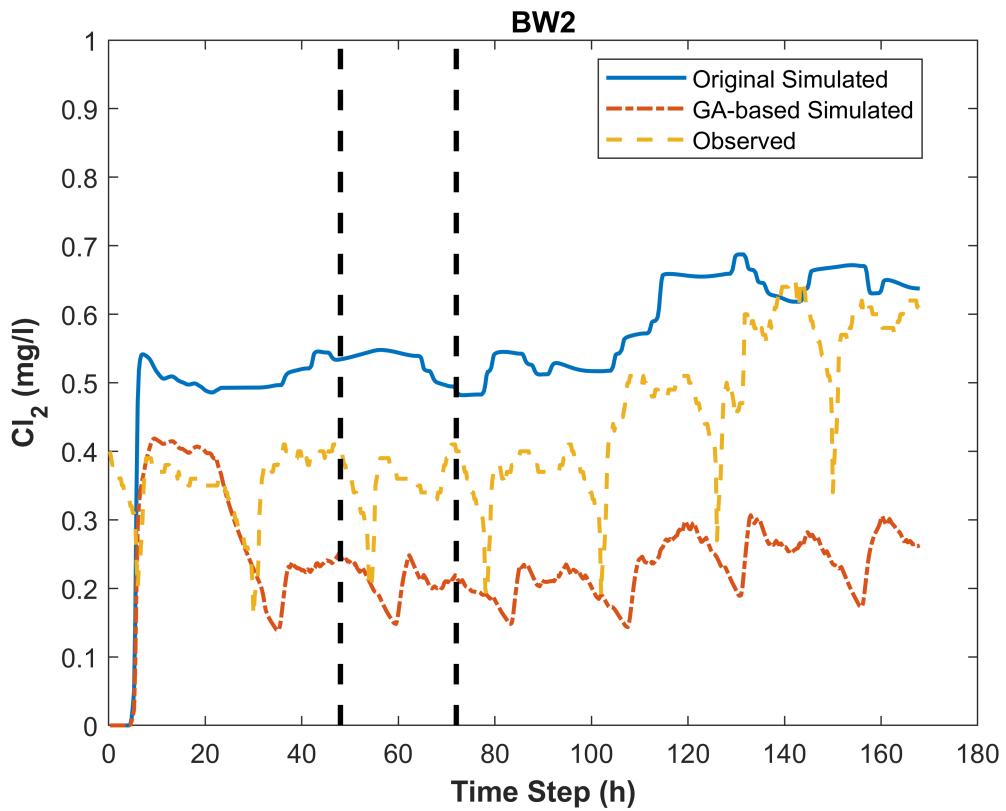
The train mse before calibration is 0.0634, after calibration it is 0.0114

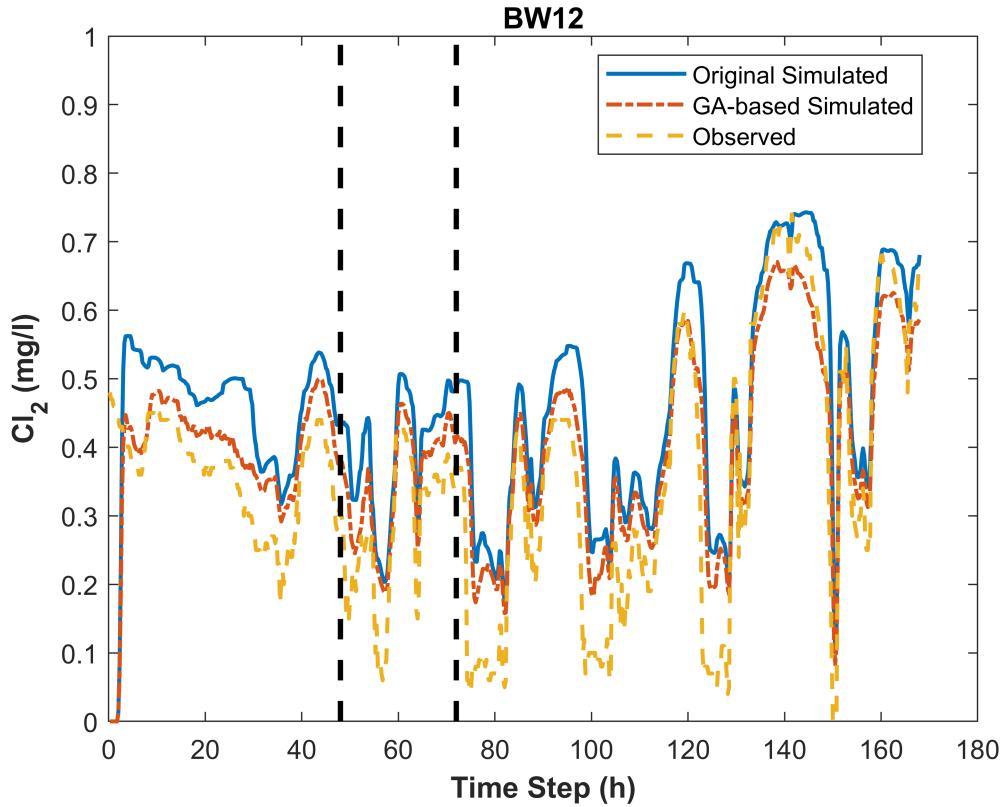
```
fprintf('The validation mse before calibration is %.4f, after calibration it is %.4f',mse_befor
```

The validation mse before calibration is 0.0631, after calibration it is 0.0311

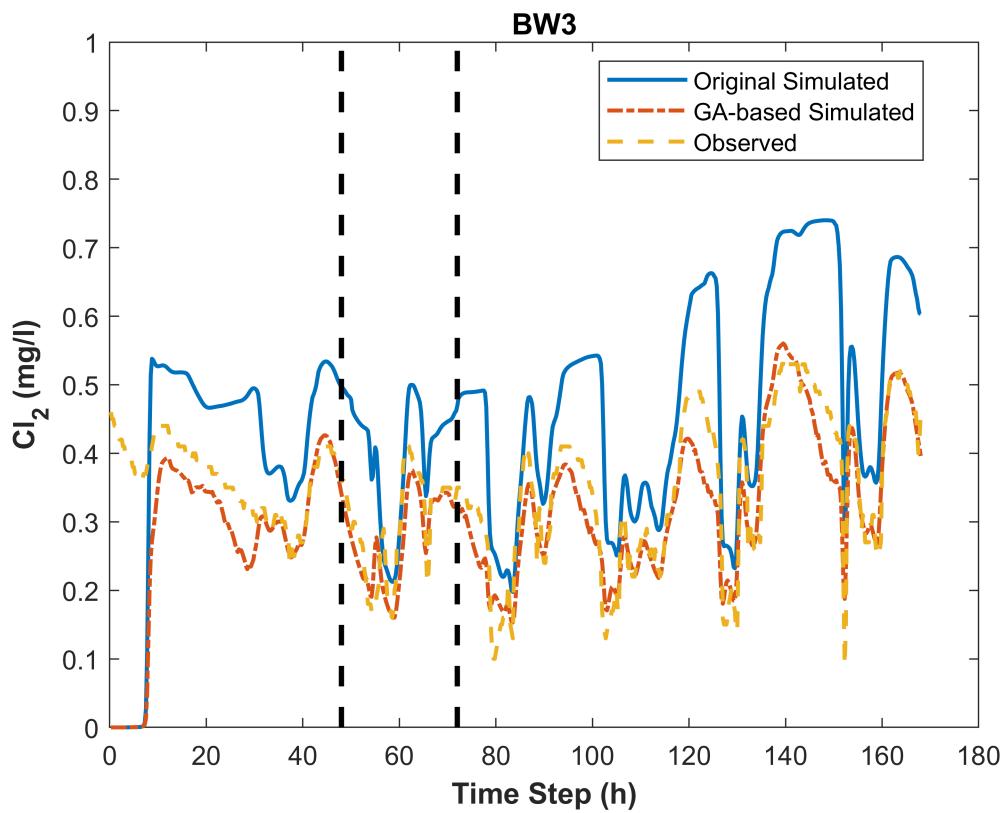
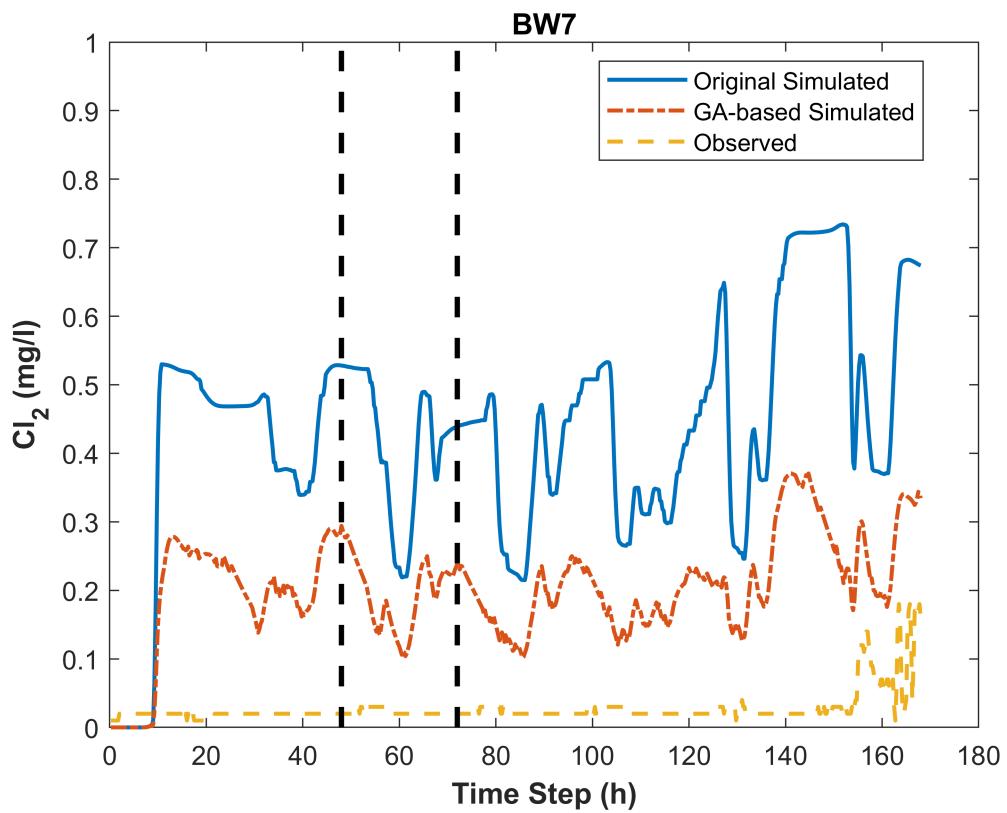
```
% figures for simulated and observed data
for i = 1:size(c_nodes_Obs,1)
    figure,
    plot(0.25:0.25:days*24,c_nodes_Sim(i,:),'-','LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Sim_NoGroup(i,:),'-','LineStyle', '-.', 'LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Obs(i,:),'-','LineStyle', '--', 'LineWidth',1.5)
    hold on
    plot([48,48],[0,1],'Color','k','LineStyle', '--', 'LineWidth',2)
    hold on
    plot([72,72],[0,1],'Color','k','LineStyle', '--', 'LineWidth',2)
    legend({'Original Simulated','GA-based Simulated','Observed'},'location','best');
    xlabel('Time Step (h)', 'fontweight', 'bold')
    ylabel('Cl2 (mg/l)', 'fontweight', 'bold')
    title(Junction_Name{i})
end
```

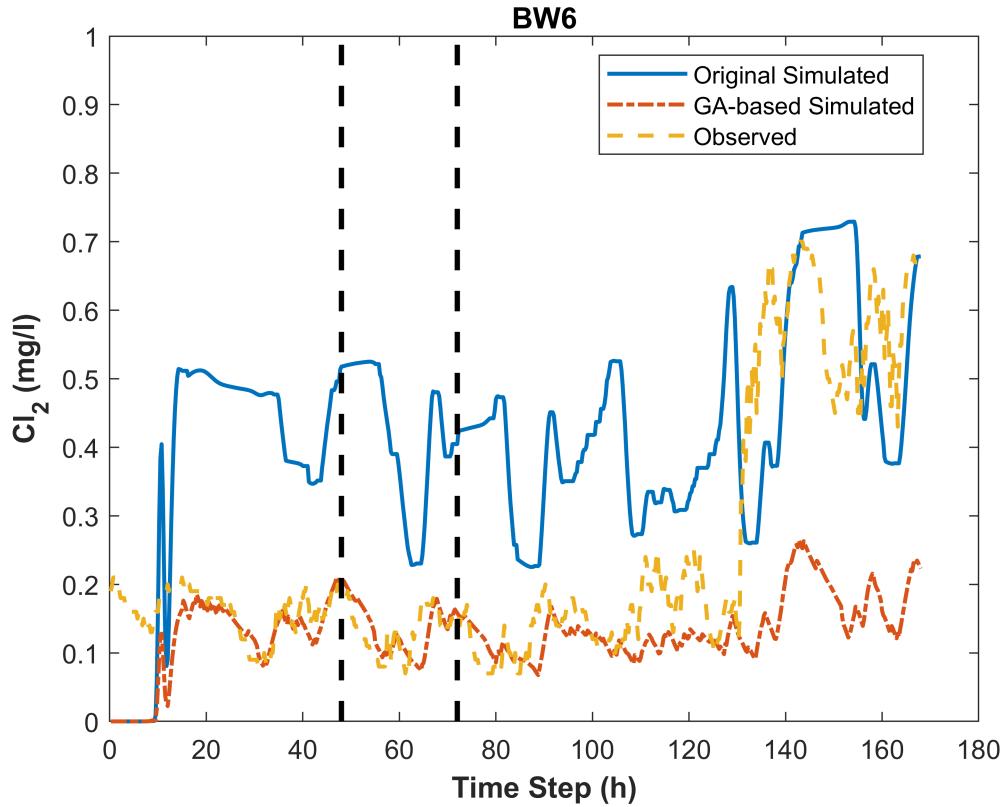






```
% figures for simulated and observed data (Hydrants)
for i = 1:size(c_nodes_HydrantObs,1)
figure,
plot(0.25:0.25:days*24,c_nodes_HydrantSim(i,:),'-','LineWidth',1.5)
hold on
plot(0.25:0.25:days*24,c_nodes_Sim_Hydrant_NoGroup(i,:),'LineStyle', '-.', 'LineWidth', 1.5)
hold on
plot(0.25:0.25:days*24,c_nodes_HydrantObs(i,:),'LineStyle', '--', 'LineWidth', 1.5)
hold on
plot([48,48],[0,1],'Color','k','LineStyle', '--', 'LineWidth', 2)
hold on
plot([72,72],[0,1],'Color','k','LineStyle', '--', 'LineWidth', 2)
legend({'Original Simulated','GA-based Simulated','Observed'},'location','best');
xlabel('Time Step (h)', 'fontweight', 'bold')
ylabel('Cl₂ (mg/l)', 'fontweight', 'bold')
title(Junction_Hydrant_Name{i})
end
```





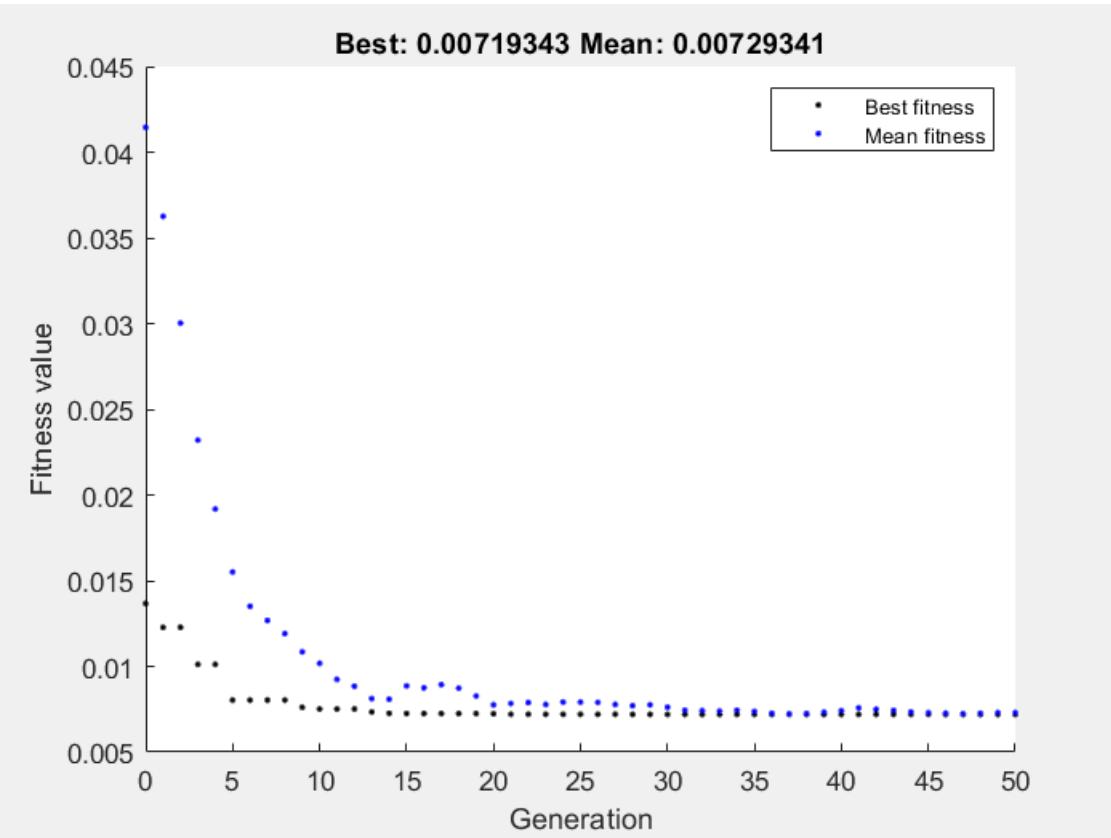
Diameter based Group Genetic Algorithm Implementation

Implement genetic algorithm to solve the parameter optimization problem based on diameter groups.

```
% % genetic algorithm implementation
rng default
options = optimoptions('ga','Display','none','PopulationType','DoubleVector','PopulationSize',50,'Generations',50,"ConstraintTolerance",1e-6,"PlotFcn","gaplotbestf");

lb = [-10,-10,-10,-10];
ub = [0,0,0,0];

tic;
Mse_function = @(x) Mse_D(x,Obs_Combined_train,Junction_CombinedIdx,D);
[x_D, fval_D] = ga(Mse_function,4,[],[],[],lb,ub,[],options);
```



```
t_D = toc;
fprintf('The computation time is %.2fs',t_D);
```

The computation time is 602.73s

```
% Test the ga result
theta_D = ones(2281,1); % units of m/day
for i = 1:2281
    if D(i) <= 75
        theta_D(i) = x_D(1);
    elseif D(i) > 75 && D(i) <= 150
        theta_D(i) = x_D(2);
    elseif D(i) > 150 && D(i) <= 250
        theta_D(i) = x_D(3);
    else
        theta_D(i) = x_D(4);
    end
end
net.setLinkWallReactionCoeff(1:np, theta_D);
net.setOptionsPipeWallReactionOrder(1);
quality_res_D = net.getComputedQualityTimeSeries;
c_nodes_D = quality_res_D.NodeQuality';
c_nodes_Sim_D = c_nodes_D(Junction_SelectIdx,1:end-1);
c_nodes_Sim_Hydrant_D = c_nodes_D(Junction_HydrantIdx,1:end-1);
c_nodes_Sim_Combined_D = [c_nodes_Sim_D;c_nodes_Sim_Hydrant_D];
```

```

Sim_Combined_D_val = c_nodes_Sim_Combined_D(:,3*96+1:end);
mse_after_D = mse_function(Sim_Combined_D_val,Obs_Combined_val);

Sim_main_D_val = c_nodes_Sim_D(:,3*96+1:end);
mse_main_after_D = mse_function(Sim_main_D_val,Obs_val);
fprintf('The validation mse of nodes in the main network after calibration is %.4f',mse_main_a

```

The validation mse of nodes in the main network after calibration is 0.0121

```

fprintf('The train mse before calibration is %.4f, after calibration it is %.4f',mse_before_trai

```

The train mse before calibration is 0.0634, after calibration it is 0.0072

```

fprintf('The validation mse before calibration is %.4f, after calibration it is %.4f',mse_befor

```

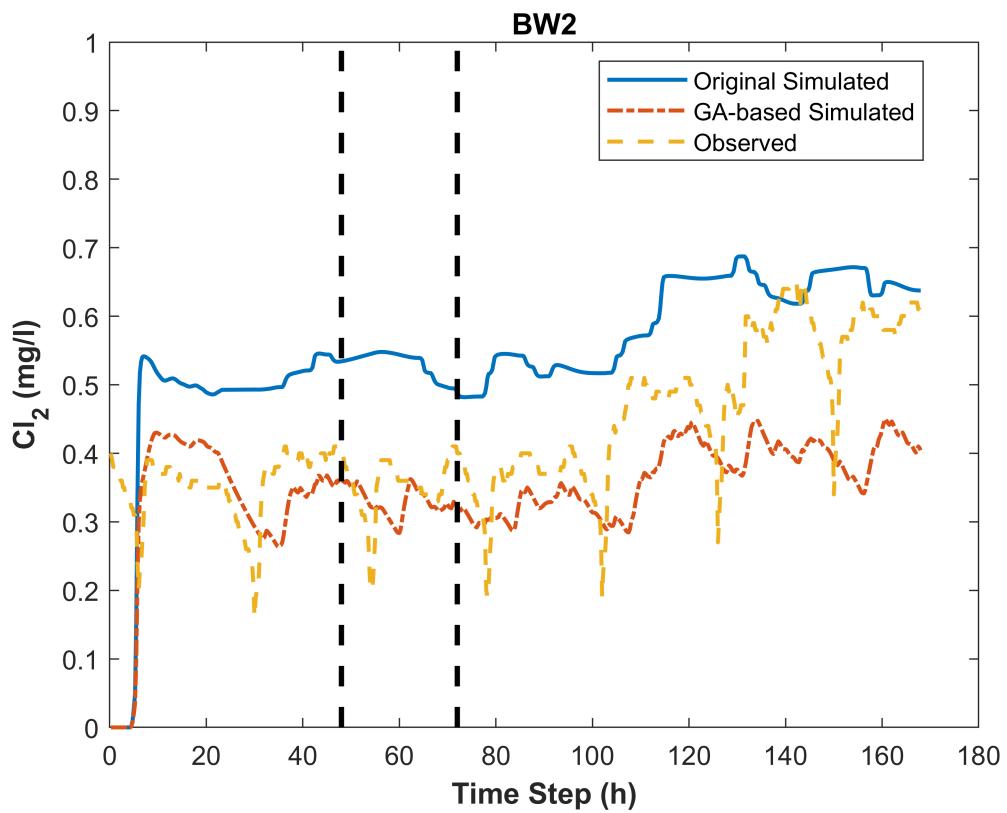
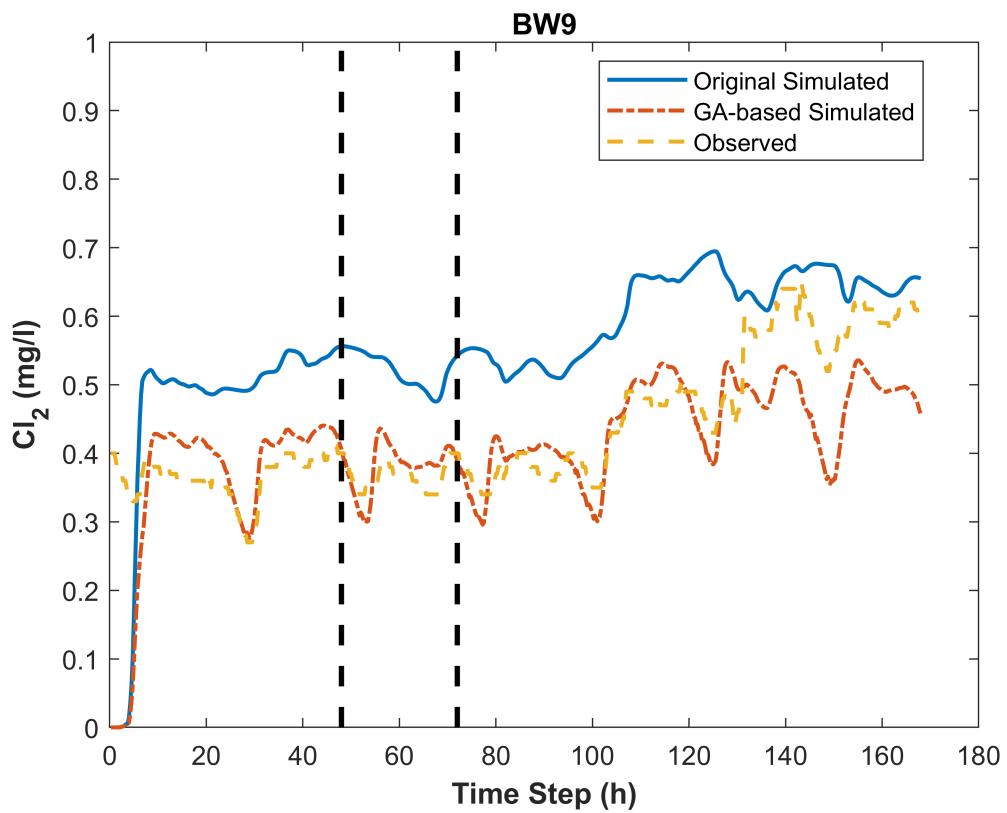
The validation mse before calibration is 0.0631, after calibration it is 0.0227

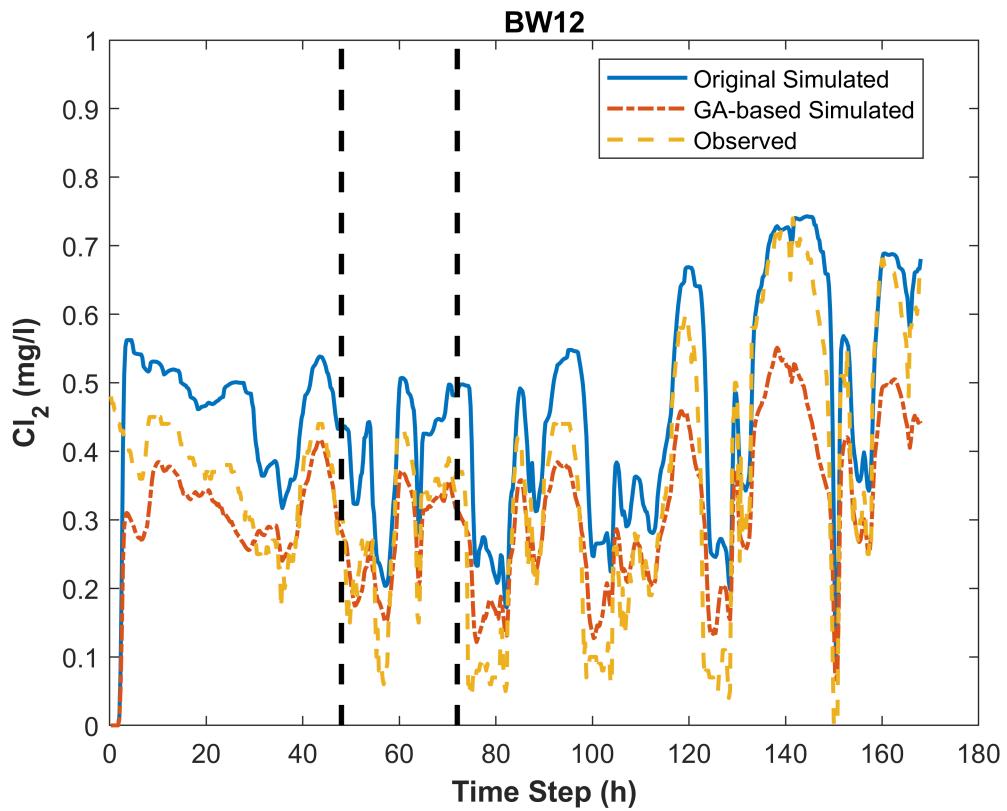
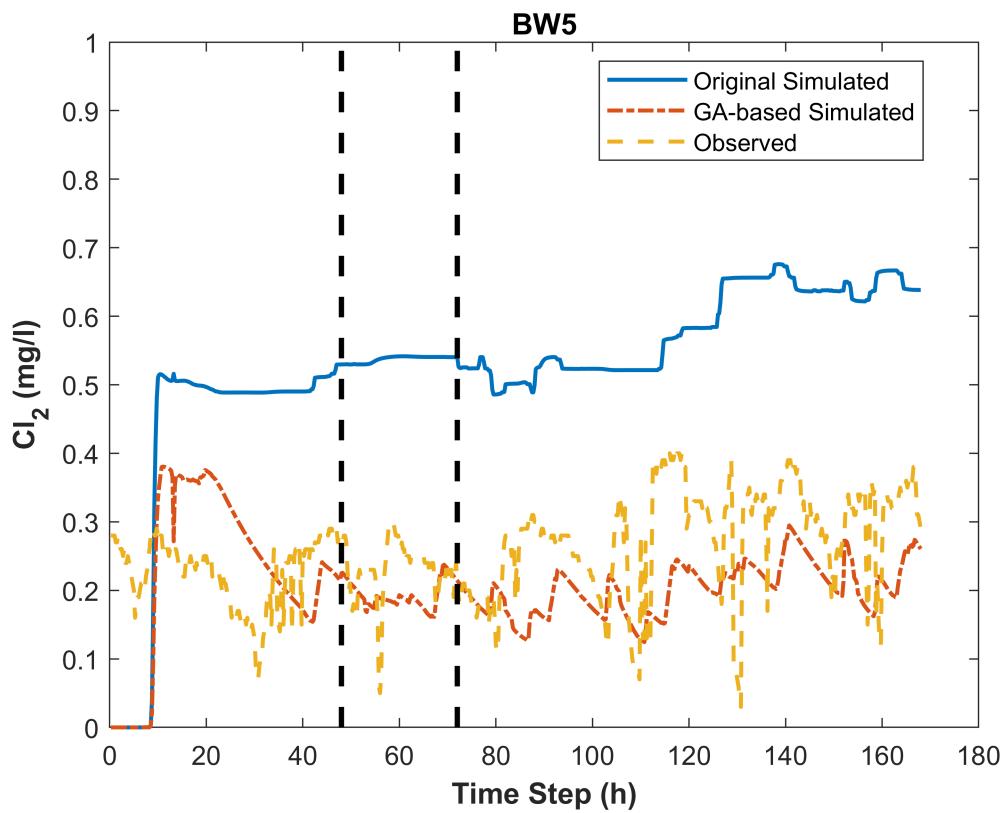
% figures for simulated and observed data

```

for i = 1:size(c_nodes_Obs,1)
    figure,
    plot(0.25:0.25:days*24,c_nodes_Sim(i,:),'-','LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Sim_D(i,:),'LineStyle','-.','LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Obs(i,:),'LineStyle','--','LineWidth',1.5)
    hold on
    plot([48,48],[0,1],'Color','k','LineStyle','--','LineWidth',2)
    hold on
    plot([72,72],[0,1],'Color','k','LineStyle','--','LineWidth',2)
    legend({'Original Simulated','GA-based Simulated','Observed'},'location','best');
    xlabel('Time Step (h)','fontweight','bold')
    ylabel('Cl_2 (mg/l)','fontweight','bold')
    title(Junction_Name{i})
end

```



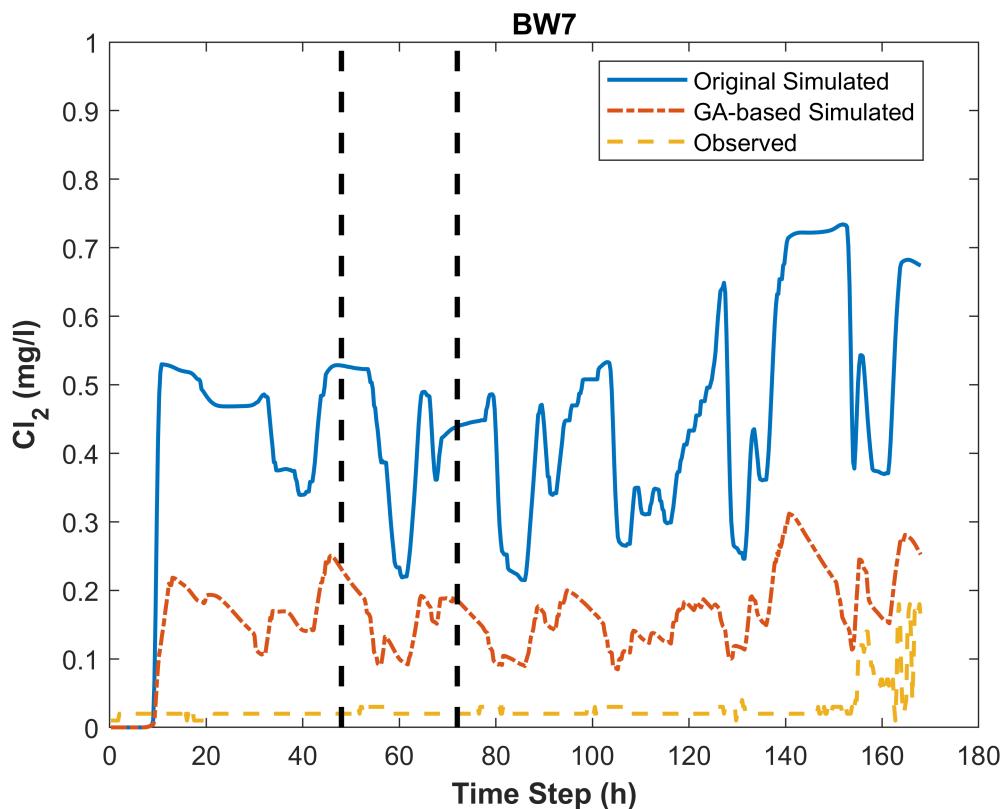


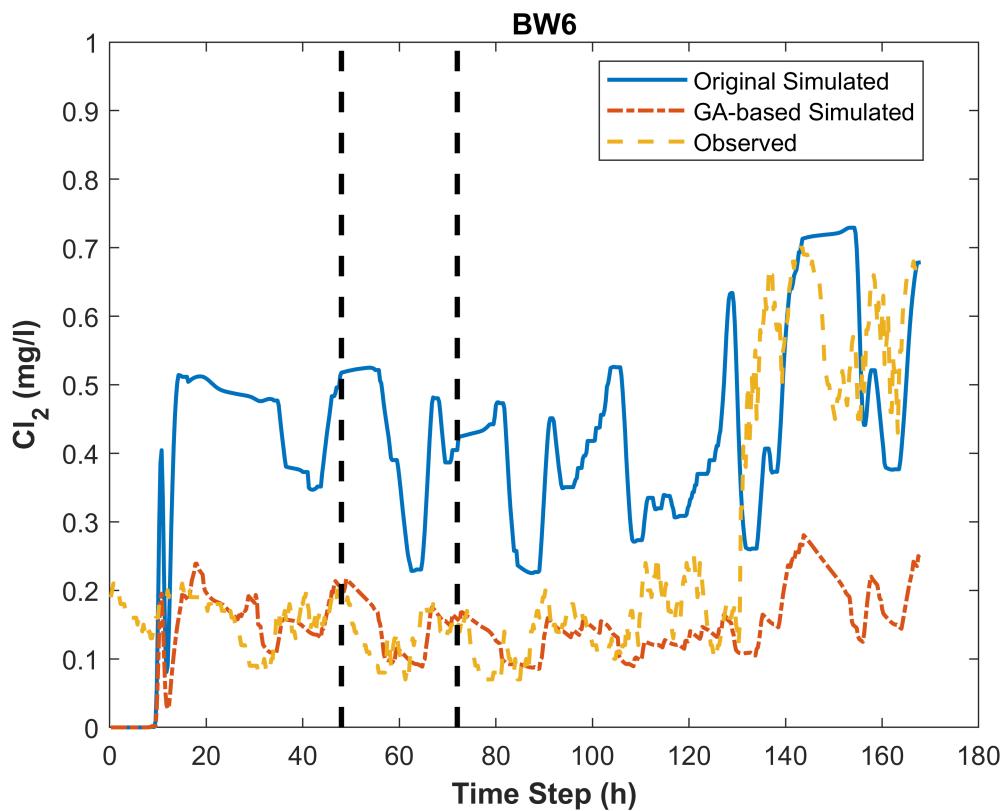
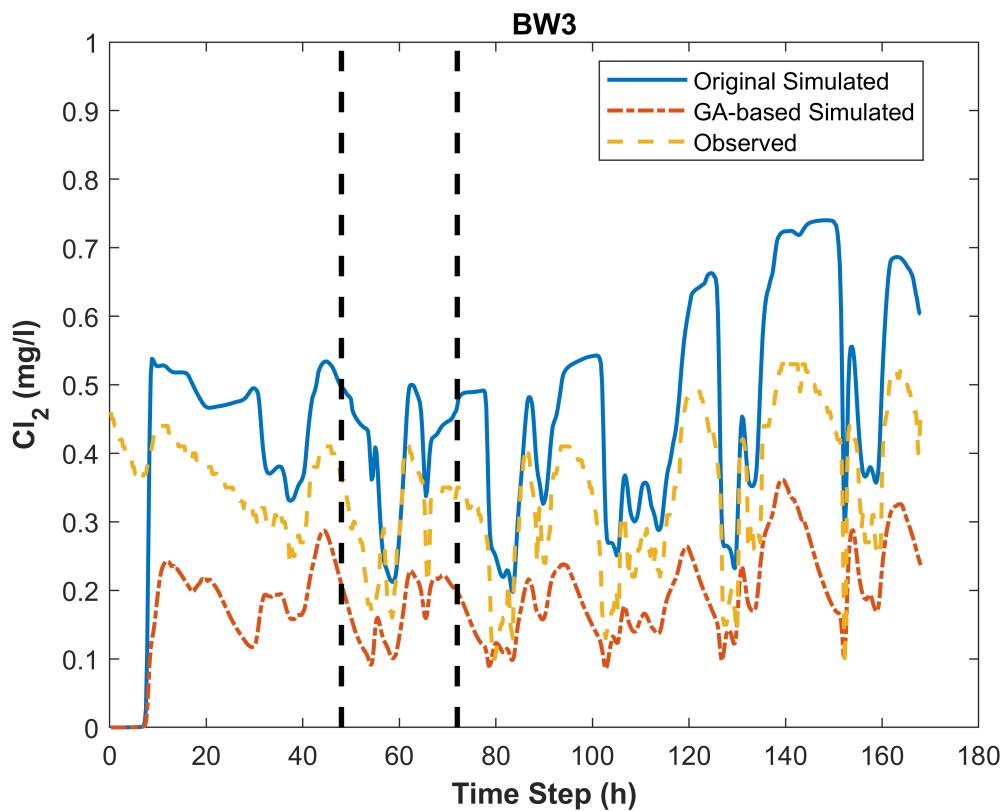
```
% figures for simulated and observed data (hydrants)
for i = 1:size(c_nodes_HydrantObs,1)
```

```

figure,
plot(0.25:0.25:days*24,c_nodes_HydrantSim(i,:),'-','LineWidth',1.5)
hold on
plot(0.25:0.25:days*24,c_nodes_Sim_Hydrant_D(i,:),'LineStyle','-.','LineWidth',1.5)
hold on
plot(0.25:0.25:days*24,c_nodes_HydrantObs(i,:), 'LineStyle','--','LineWidth',1.5)
hold on
plot([48,48],[0,1], 'Color','k','LineStyle','--','LineWidth',2)
hold on
plot([72,72],[0,1], 'Color','k','LineStyle','--','LineWidth',2)
legend({'Original Simulated','GA-based Simulated','Observed'},'location','best')
xlabel('Time Step (h)', 'fontweight','bold')
ylabel('Cl2 (mg/l)', 'fontweight','bold')
title(Junction_Hydrant_Name{i})
end

```



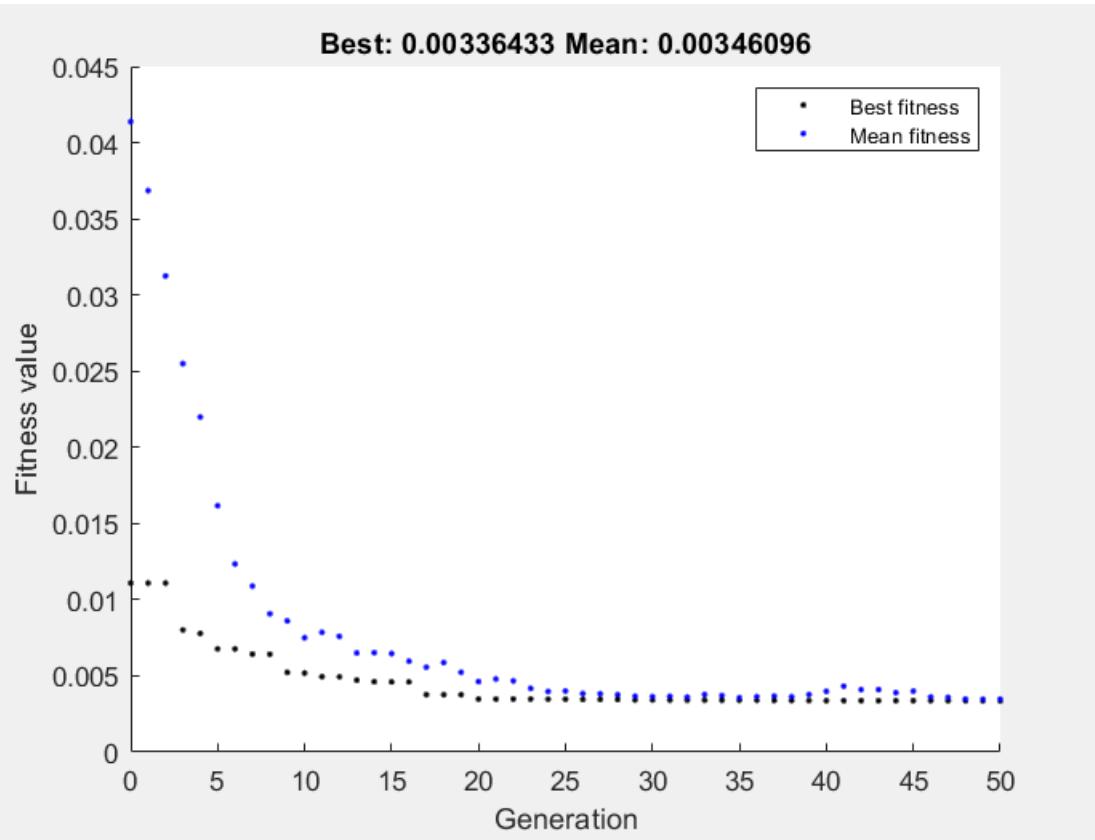


Roughness based Group Genetic Algorithm Implementation

Implement genetic algorithm to solve the parameter optimization problem based on roughness groups.

```
% genetic algorithm implementation
rng default
options = optimoptions('ga','Display','none','PopulationType','DoubleVector','PopulationSize',50,'Generations',50,"ConstraintTolerance",1e-6,"PlotFcn","gaplotbestf");
lb = -10*ones(11,1);
ub = 0*ones(11,1);

tic;
Mse_function = @(x) Mse_R(x,Obs_Combined_train,Junction_CombinedIdx,R);
[x_R, fval_R] = ga(Mse_function,11,[],[],[],[],lb,ub,[],options);
```



```
t_R = toc;
fprintf('The computation time is %.2fs',t_R);
```

The computation time is 647.56s

```
% Test the ga result
theta_R = ones(2281,1); % units of m/day
for i = 1:2281
    if abs(R(i) - 0) <= 0.01
        theta_R(i) = x_R(1);
    elseif abs(R(i) - 0.1000) <= 0.01
        theta_R(i) = x_R(2);
    elseif abs(R(i) - 0.1399) <= 0.01
        theta_R(i) = x_R(3);
```

```

elseif abs(R(i) - 1.5267) <= 0.01
    theta_R(i) = x_R(4);
elseif abs(R(i) - 2.1576) <= 0.01
    theta_R(i) = x_R(5);
elseif abs(R(i) - 9.0114) <= 0.01
    theta_R(i) = x_R(6);
elseif abs(R(i) - 10.3677) <= 0.01
    theta_R(i) = x_R(7);
elseif abs(R(i) - 11.6809) <= 0.01
    theta_R(i) = x_R(8);
elseif abs(R(i) - 14.8687) <= 0.01
    theta_R(i) = x_R(9);
elseif abs(R(i) - 16.2377) <= 0.01
    theta_R(i) = x_R(10);
else
    theta_R(i) = x_R(11);
end
end

net.setLinkWallReactionCoeff(1:np, theta_R);
net.setOptionsPipeWallReactionOrder(1);
quality_res_R = net.getComputedQualityTimeSeries;
c_nodes_R = quality_res_R.NodeQuality';
c_nodes_Sim_R = c_nodes_R(Junction_SelectIdx,1:end-1);
c_nodes_Sim_Hydrant_R = c_nodes_R(Junction_HydrantIdx,1:end-1);

c_nodes_Sim_Combined_R = [c_nodes_Sim_R;c_nodes_Sim_Hydrant_R];
Sim_Combined_R_val = c_nodes_Sim_Combined_R(:,3*96+1:end);
mse_after_R = mse_function(Sim_Combined_R_val,Obs_Combined_val);

Sim_main_R_val = c_nodes_Sim_R(:,3*96+1:end);
mse_main_after_R = mse_function(Sim_main_R_val,Obs_val);
fprintf('The validation mse of nodes in the main network after calibration is %.4f',mse_main_at

```

The validation mse of nodes in the main network after calibration is 0.0112

```
fprintf('The train mse before calibration is %.4f, after calibration it is %.4f',mse_before_trai
```

The train mse before calibration is 0.0634, after calibration it is 0.0034

```
fprintf('The validation mse before calibration is %.4f, after calibration it is %.4f',mse_befor
```

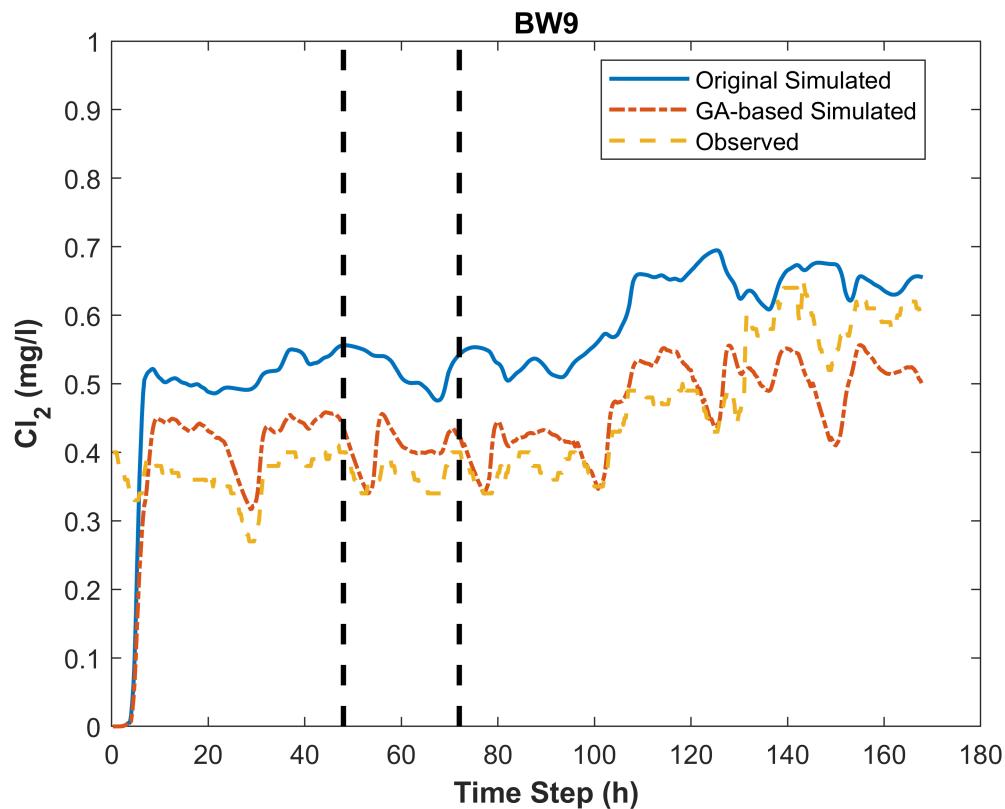
The validation mse before calibration is 0.0631, after calibration it is 0.0171

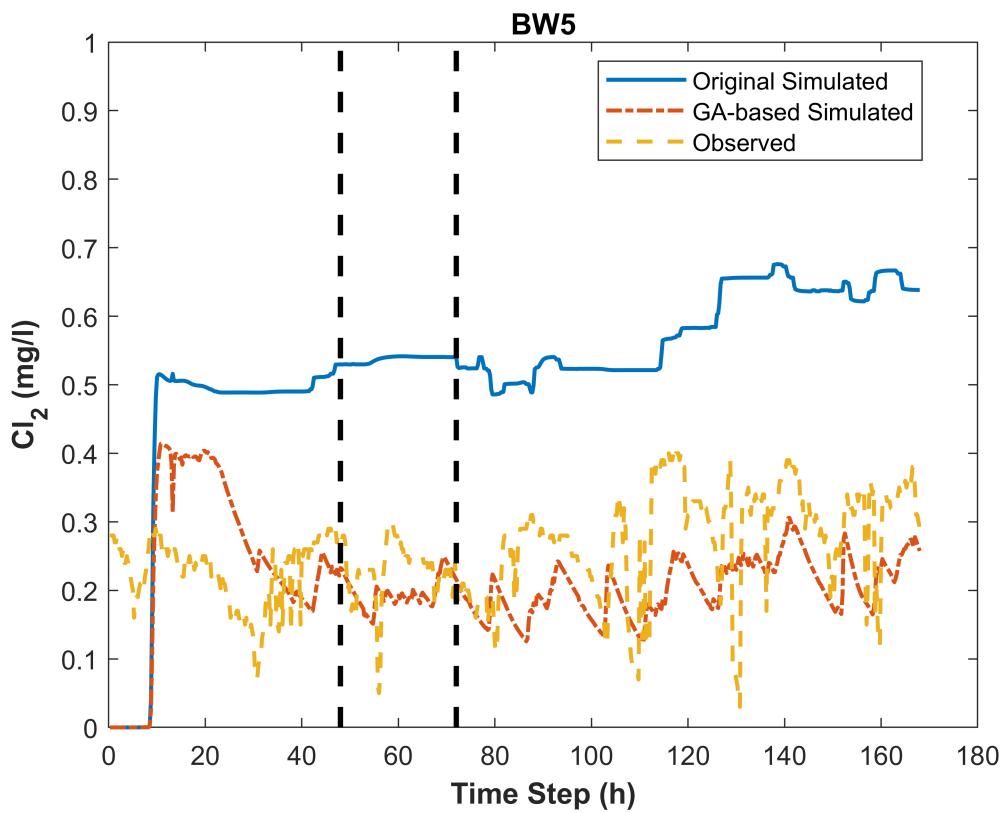
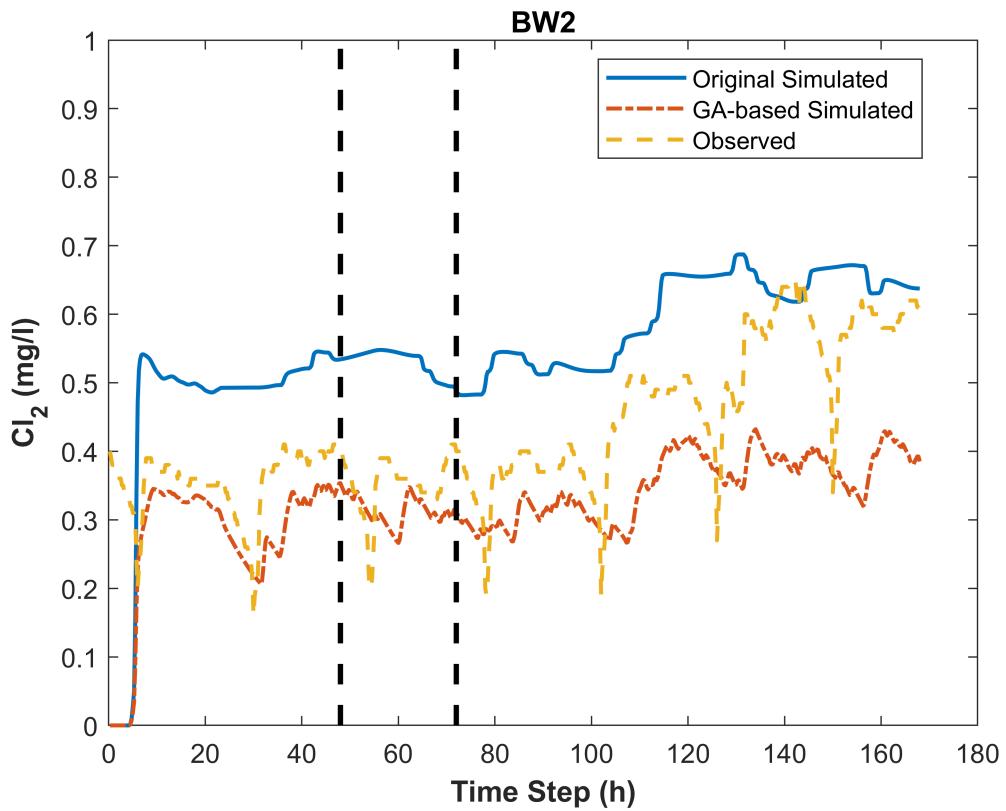
```
% figures for simulated and observed data
for i = 1:size(c_nodes_Obs,1)
    figure,
    plot(0.25:0.25:days*24,c_nodes_Sim(i,:),'-','LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Sim_R(i,:),'LineStyle','-.','LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Obs(i,:), 'LineStyle','--','LineWidth',1.5)
    hold on
    plot([48,48],[0,1], 'Color','k','LineStyle','--','LineWidth',2)
```

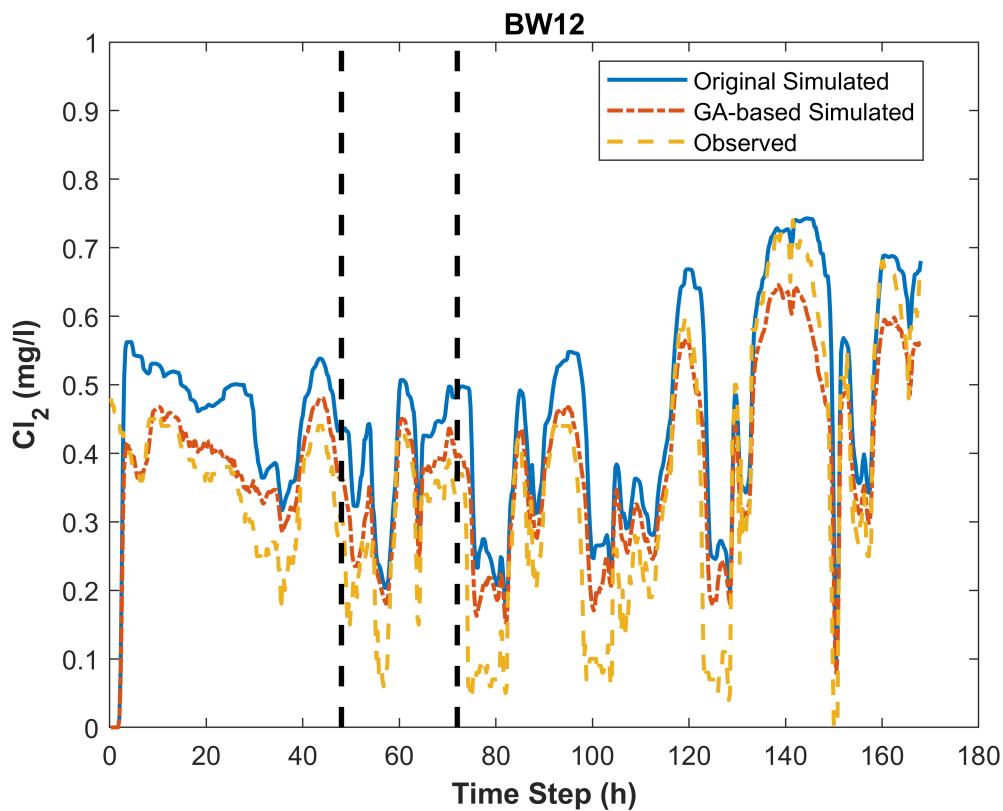
```

hold on
plot([72,72],[0,1], 'Color','k', 'LineStyle', '--', 'LineWidth', 2)
legend({'Original Simulated', 'GA-based Simulated', 'Observed'}, 'location', 'best');
xlabel('Time Step (h)', 'fontweight', 'bold')
ylabel('Cl2 (mg/l)', 'fontweight', 'bold')
title(Junction_Name{i})
end

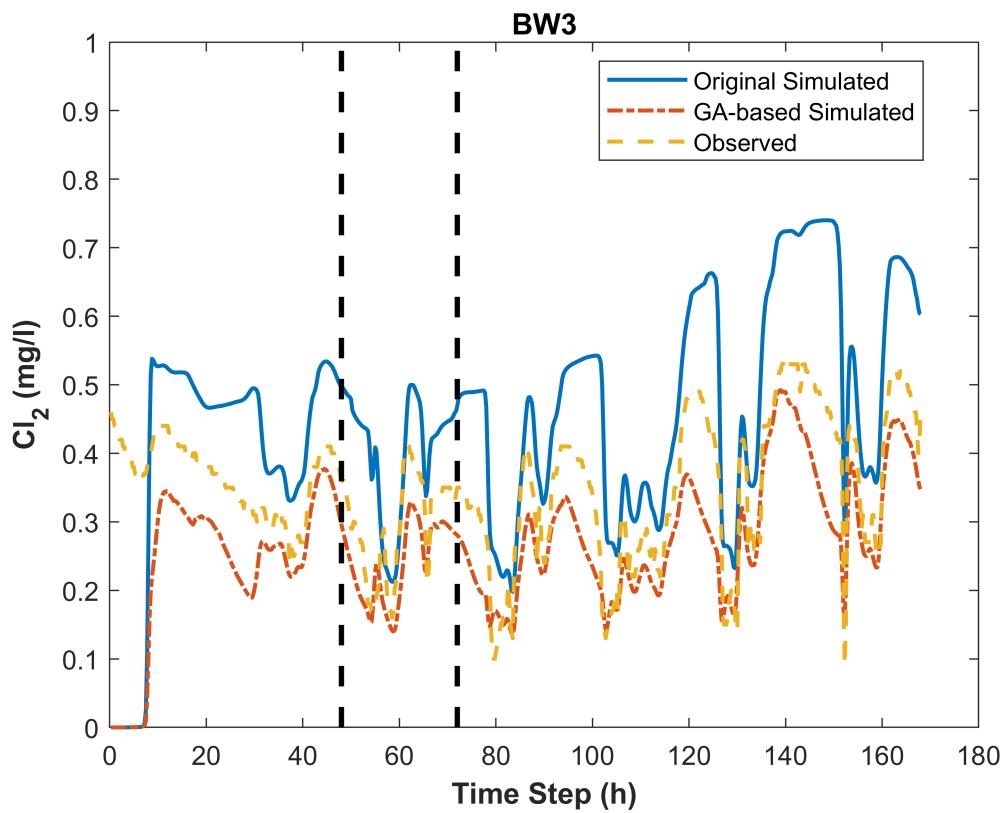
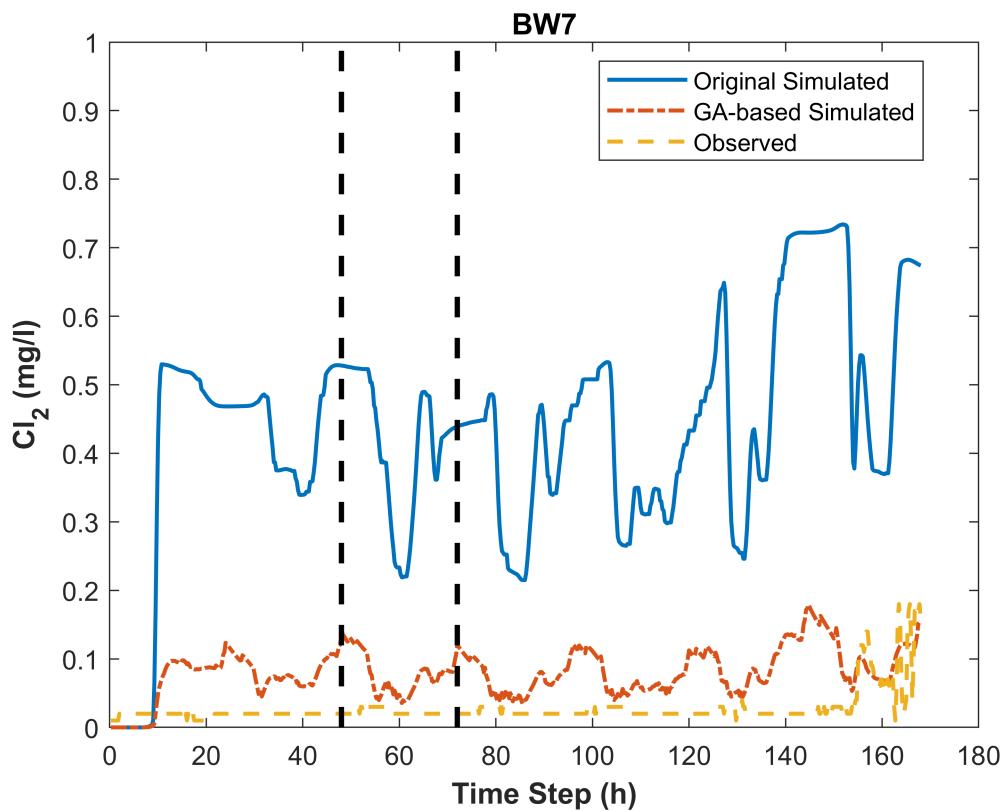
```

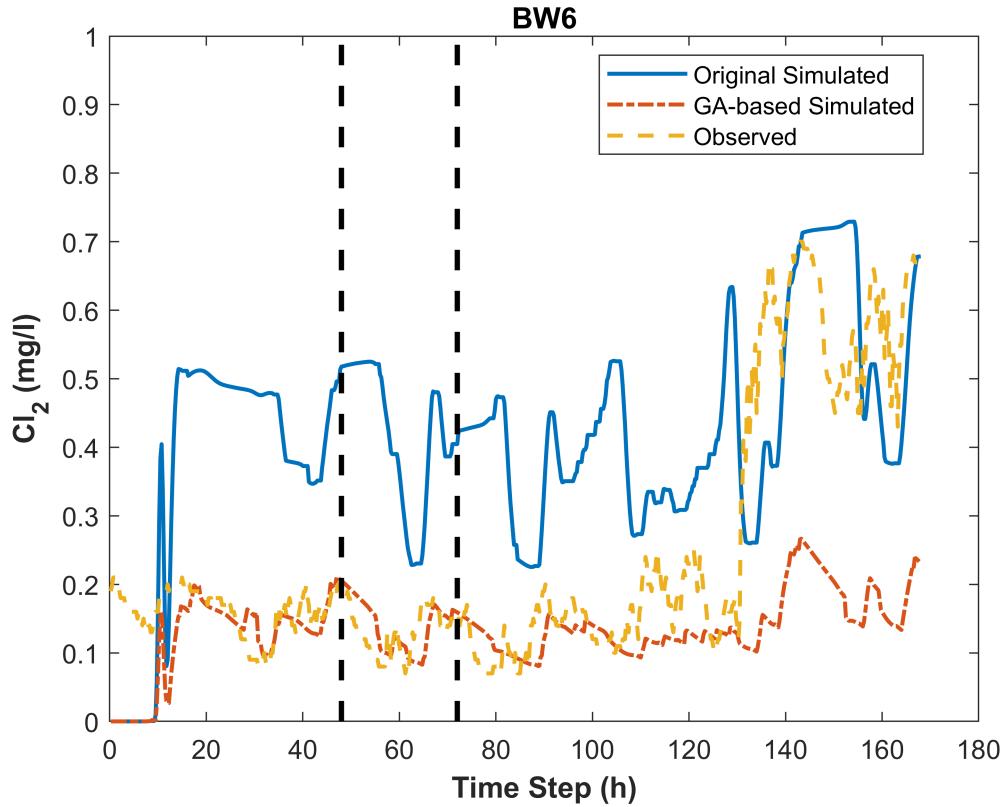






```
% figures for simulated and observed data (hydrants)
for i = 1:size(c_nodes_HydrantObs,1)
    figure,
    plot(0.25:0.25:days*24,c_nodes_HydrantSim(i,:),'-','LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Sim_Hydrant_R(i,:),'LineStyle','-.','LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_HydrantObs(i,:),'LineStyle','--','LineWidth',1.5)
    hold on
    plot([48,48],[0,1],'Color','k','LineStyle','--','LineWidth',2)
    hold on
    plot([72,72],[0,1],'Color','k','LineStyle','--','LineWidth',2)
    legend({'Original Simulated','GA-based Simulated','Observed'},'location','best');
    xlabel('Time Step (h)', 'fontweight','bold')
    ylabel('Cl₂ (mg/l)', 'fontweight','bold')
    title(Junction_Hydrant_Name{i})
end
```





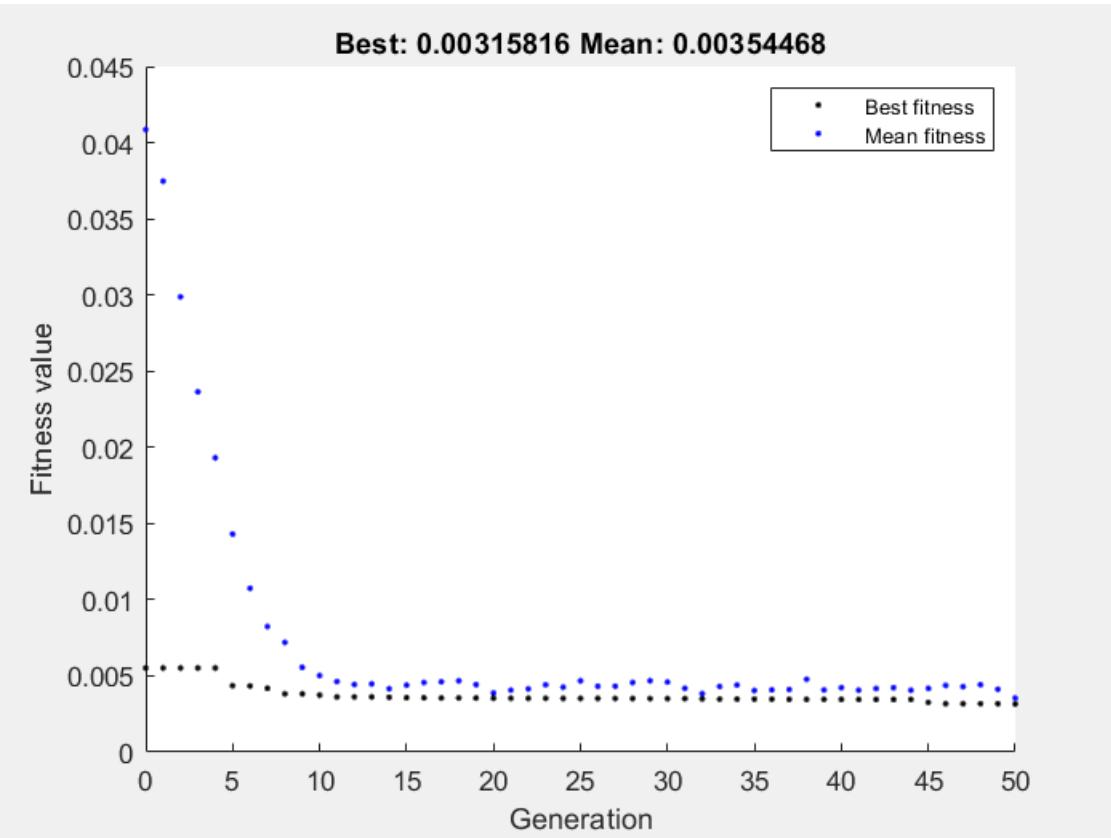
Roughness and Diameter based Group Genetic Algorithm Implementation

Implement genetic algorithm to solve the parameter optimization problem based on roughness and diameter groups, in this we consider the roughness groups and diameter groups affect the bulk decay together.

```
% genetic algorithm implementation
rng default
options = optimoptions('ga','Display','none','PopulationType','DoubleVector','PopulationSize',50
    'Generations',50,"ConstraintTolerance",1e-6,"PlotFcn","gaplotbestf");

lb = -10*ones(22,1);
ub = 0*ones(22,1);

tic;
Mse_function = @(x) Mse_DandR(x,Obs_Combined_train,Junction_CombinedIdx,D,R);
[x_DandR, fval_DandR] = ga(Mse_function,22,[],[],[],[],lb,ub,[],options);
```



```
t_DandR = toc;
fprintf('The computation time is %.2fs',t_DandR);
```

The computation time is 623.11s

```
% Test the ga result
thetaDandR = ones(2281,1); % units of m/day
for i = 1:2281
    if abs(R(i) - 0) <= 0.01
        if D(i) <= 150
            thetaDandR(i) = x_DandR(1);
        else
            thetaDandR(i) = x_DandR(2);
        end
    elseif abs(R(i) - 0.1000) <= 0.01
        if D(i) <= 150
            thetaDandR(i) = x_DandR(3);
        else
            thetaDandR(i) = x_DandR(4);
        end
    elseif abs(R(i) - 0.1399) <= 0.01
        if D(i) <= 150
            thetaDandR(i) = x_DandR(5);
        else
            thetaDandR(i) = x_DandR(6);
        end
    end
```

```

elseif abs(R(i) - 1.5267) <= 0.01
if D(i) <= 150
    thetaDandR(i) = x_DandR(7);
else
    thetaDandR(i) = x_DandR(8);
end
elseif abs(R(i) - 2.1576) <= 0.01
if D(i) <= 150
    thetaDandR(i) = x_DandR(9);
else
    thetaDandR(i) = x_DandR(10);
end
elseif abs(R(i) - 9.0114) <= 0.01
if D(i) <= 150
    thetaDandR(i) = x_DandR(11);
else
    thetaDandR(i) = x_DandR(12);
end
elseif abs(R(i) - 10.3677) <= 0.01
if D(i) <= 150
    thetaDandR(i) = x_DandR(13);
else
    thetaDandR(i) = x_DandR(14);
end
elseif abs(R(i) - 11.6809) <= 0.01
if D(i) <= 150
    thetaDandR(i) = x_DandR(15);
else
    thetaDandR(i) = x_DandR(16);
end
elseif abs(R(i) - 14.8687) <= 0.01
if D(i) <= 150
    thetaDandR(i) = x_DandR(17);
else
    thetaDandR(i) = x_DandR(18);
end
elseif abs(R(i) - 16.2377) <= 0.01
if D(i) <= 150
    thetaDandR(i) = x_DandR(19);
else
    thetaDandR(i) = x_DandR(20);
end
else
if D(i) <= 150
    thetaDandR(i) = x_DandR(21);
else
    thetaDandR(i) = x_DandR(22);
end
end
end

net.setLinkWallReactionCoeff(1:np, thetaDandR);
net.setOptionsPipeWallReactionOrder(1);

```

```

quality_res_DandR = net.getComputedQualityTimeSeries;
c_nodes_DandR = quality_res_DandR.NodeQuality';
c_nodes_Sim_DandR = c_nodes_DandR(Junction_SelectIdx,1:end-1);
c_nodes_Sim_Hydrant_DandR = c_nodes_DandR(Junction_HydrantIdx,1:end-1);

c_nodes_Sim_Combined_DandR = [c_nodes_Sim_DandR;c_nodes_Sim_Hydrant_DandR];
Sim_Combined_DandR_val = c_nodes_Sim_Combined_DandR(:,3*96+1:end);
mse_after_DandR = mse_function(Sim_Combined_DandR_val,Obs_Combined_val);

Sim_main_DandR_val = c_nodes_Sim_DandR(:,3*96+1:end);
mse_main_after_DandR = mse_function(Sim_main_DandR_val,Obs_val);
fprintf('The validation mse of nodes in the main network after calibration is %.4f',mse_main_at

```

The validation mse of nodes in the main network after calibration is 0.0094

```

fprintf('The train mse before calibration is %.4f, after calibration it is %.4f',mse_before_trai

```

The train mse before calibration is 0.0634, after calibration it is 0.0032

```

fprintf('The validation mse before calibration is %.4f, after calibration it is %.4f',mse_befor

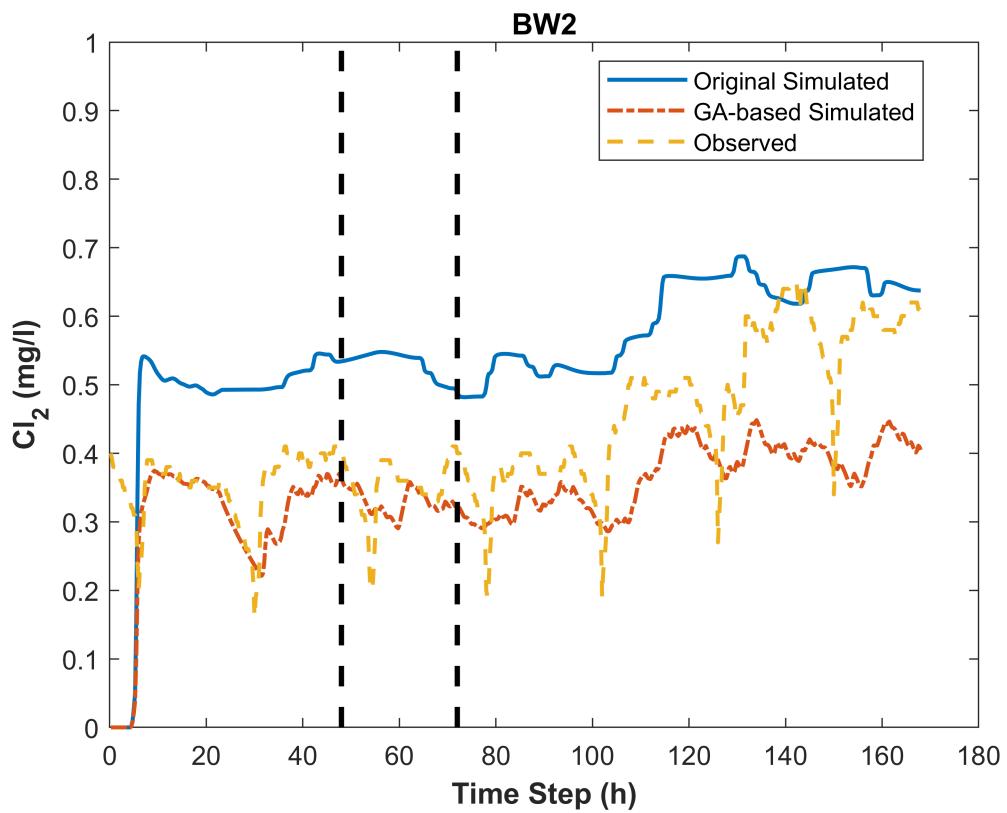
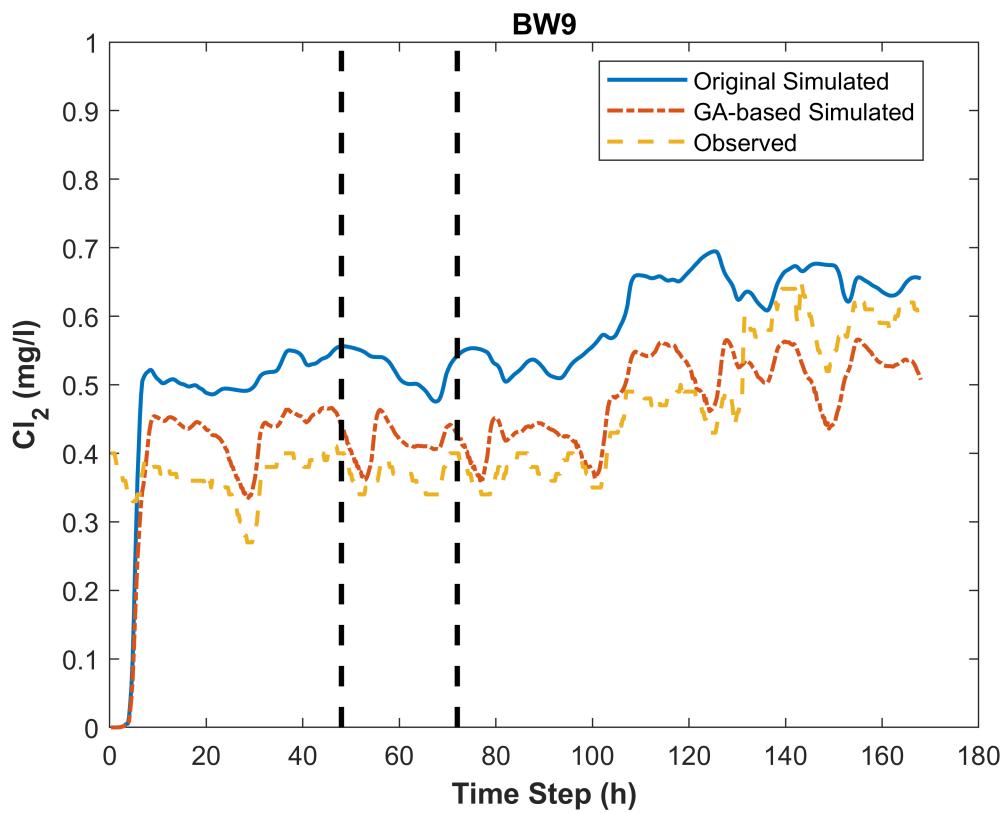
```

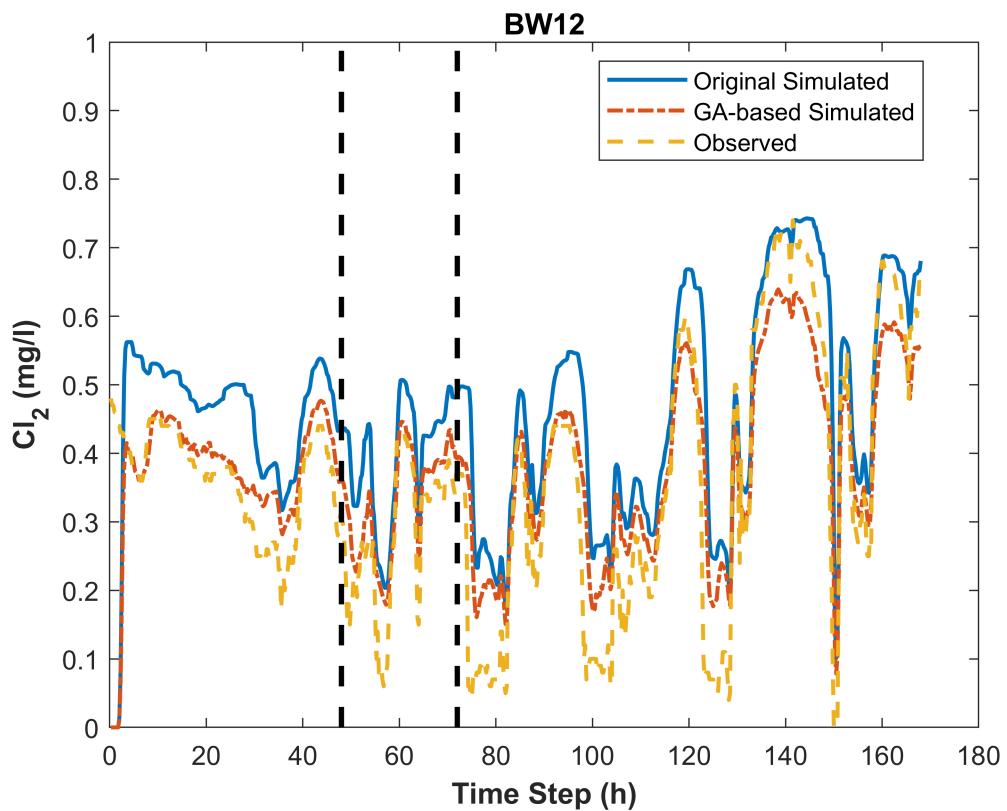
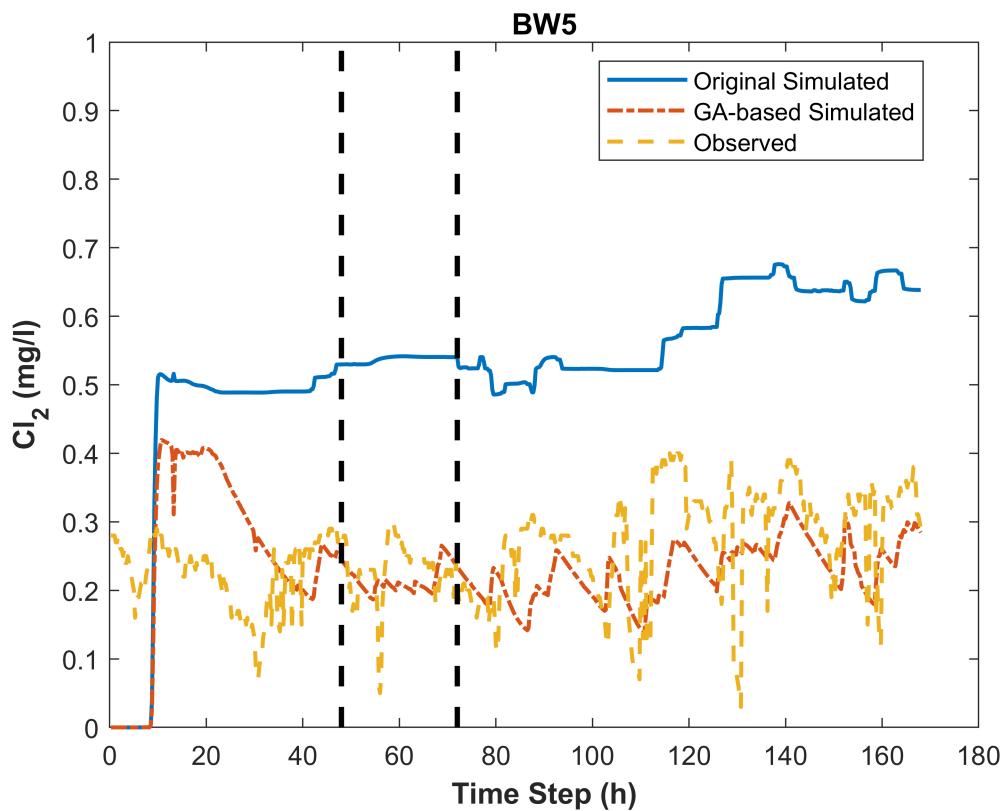
The validation mse before calibration is 0.0631, after calibration it is 0.0160

```

% figures for simulated and observed data
for i = 1:size(c_nodes_Obs,1)
    figure,
    plot(0.25:0.25:days*24,c_nodes_Sim(i,:),'-', 'LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Sim_DandR(i,:),'LineStyle','-.','LineWidth',1.5)
    hold on
    plot(0.25:0.25:days*24,c_nodes_Obs(i,:),'LineStyle','--','LineWidth',1.5)
    hold on
    plot([48,48],[0,1],'Color','k','LineStyle','--','LineWidth',2)
    hold on
    plot([72,72],[0,1],'Color','k','LineStyle','--','LineWidth',2)
    legend({'Original Simulated','GA-based Simulated','Observed'},'location','best');
    xlabel('Time Step (h)', 'fontWeight','bold')
    ylabel('Cl₂ (mg/l)', 'fontWeight','bold')
    title(Junction_Name{i})
end

```



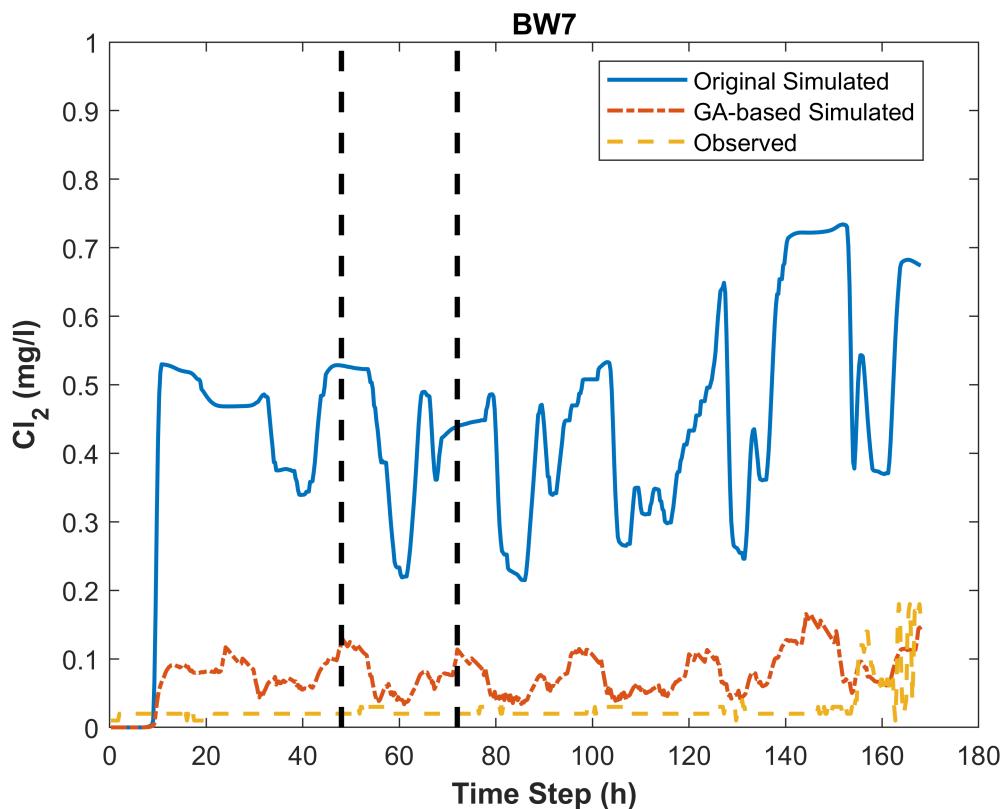


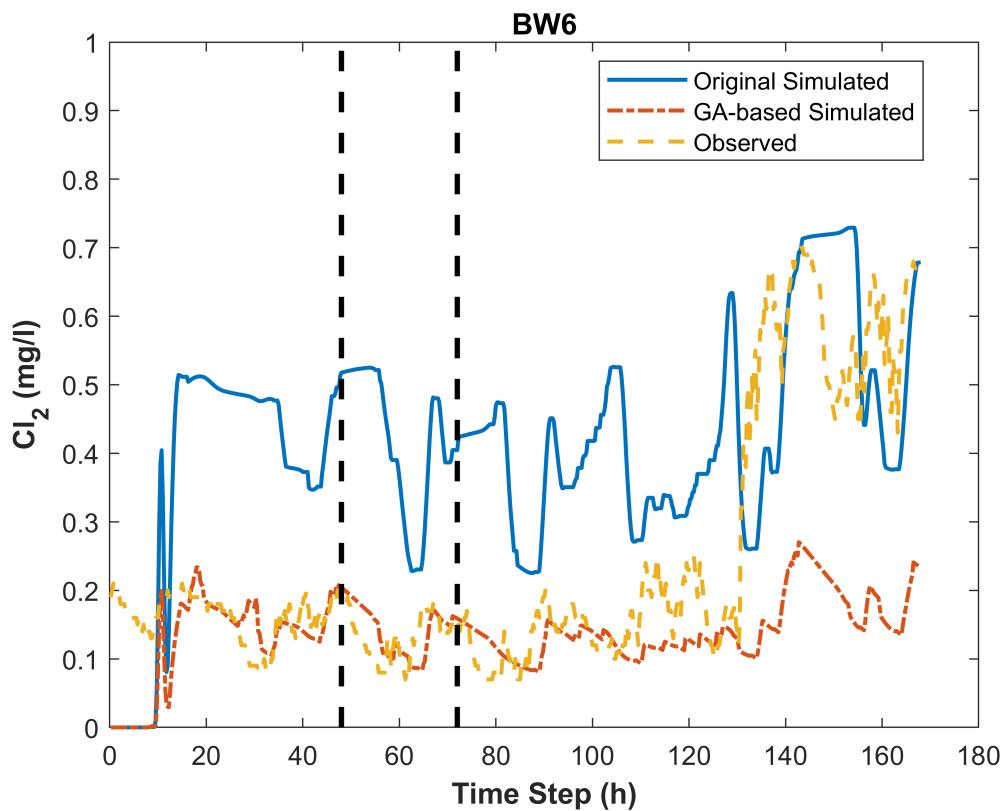
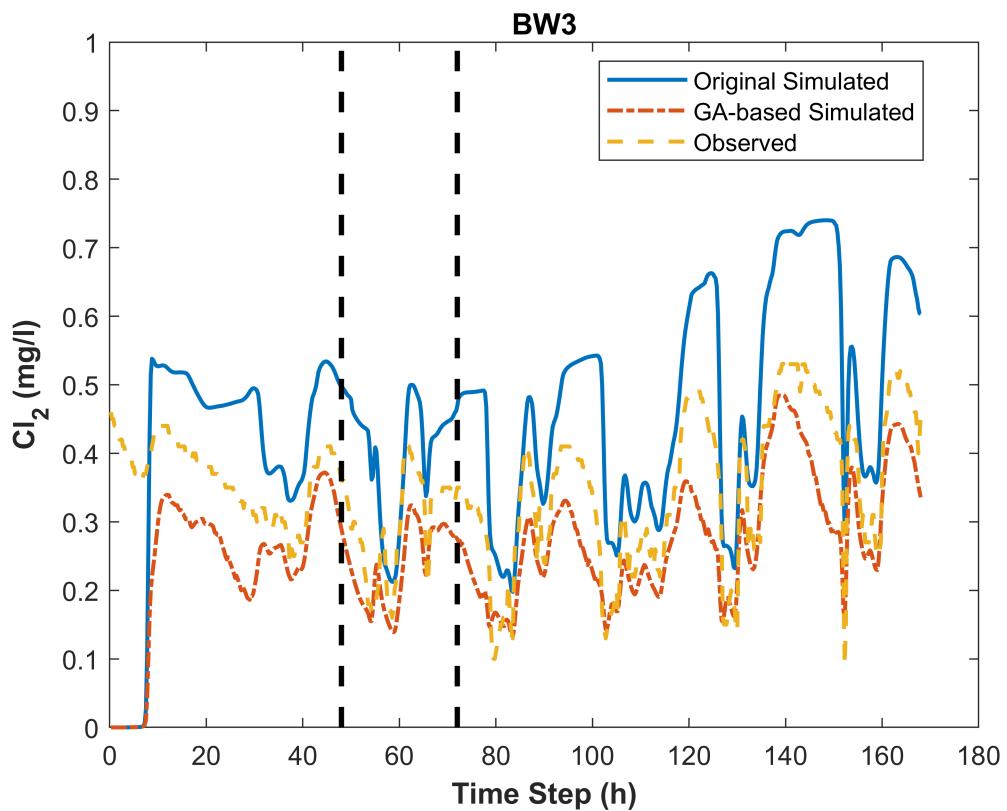
```
% figures for simulated and observed data (hydrants)
for i = 1:size(c_nodes_HydrantObs,1)
```

```

figure,
plot(0.25:0.25:days*24,c_nodes_HydrantSim(i,:),'-','LineWidth',1.5)
hold on
plot(0.25:0.25:days*24,c_nodes_Sim_Hydrant_DandR(i,:),'LineStyle','-.','LineWidth',1.5)
hold on
plot(0.25:0.25:days*24,c_nodes_HydrantObs(i,:),'LineStyle','--','LineWidth',1.5)
hold on
plot([48,48],[0,1],'Color','k','LineStyle','--','LineWidth',2)
hold on
plot([72,72],[0,1],'Color','k','LineStyle','--','LineWidth',2)
legend({'Original Simulated','GA-based Simulated','Observed'},'location','best')
xlabel('Time Step (h)','fontweight','bold')
ylabel('Cl2 (mg/l)', 'fontweight','bold')
title(Junction_Hydrant_Name{i})
end

```





```
fprintf('The single group k_w value is %.4f.',x_NoGroup);
```

The single group k_w value is -0.1126.

```
fprintf('The D-based group k_w value is %.4f, %.4f, %.4f, %.4f.',x_D);
```

The D-based group k_w value is -0.4180, 0.0000, -0.1045, -0.7969.

```
fprintf(['The R-based group k_w value is %.4f, %.4f, %.4f, %.4f, %.4f,\n %.4f, %.4f' ...
    ', %.4f, %.4f, %.4f, %.4f'],x_R);
```

The R-based group k_w value is -1.9972, 0.0000, -0.3270, -5.8649, 0.0000,
-0.0504, -0.1836, 0.0000, 0.0000, -1.0921.

```
fprintf(['The DR-based group k_w value is %.4f, %.4f, %.4f, %.4f, %.4f, %.4f, %.4f\n' ...
    ', %.4f, %.4f, %.4f, %.4f, %.4f, %.4f, %.4f\n' ...
    ', %.4f, %.4f, %.4f, %.4f, %.4f, %.4f, %.4f.'],x_DandR);
```

The DR-based group k_w value is -3.1952, -9.4605, -0.0088, -1.8750, 0.0000, -0.3077, -7.2759
, -2.0625, 0.0000, 0.0000, -8.6920, -0.1250, -0.1250, -0.1560, 0.0000
, -0.1425, -0.1241, -7.5744, 0.0000, 0.0000, 0.0000, -1.6450,

Test the hydrants

Find out the regulations in the node with hydrants.

```
% % Boxplot to see the residual
% c_nodes_hydrant_Residual = c_nodes_hydrantObs - c_nodes_hydrantSim;
% c_nodes_hydrant_Residual = c_nodes_hydrant_Residual';

% Calculate the residuals at BW nodes.
c_nodes_NoGroup_Residual = [c_nodes_Obs;c_nodes_HydrantObs] - c_nodes_NoGroup([Junction_SelectIdx]);
c_nodes_NoGroup_Residual = c_nodes_NoGroup_Residual';

c_nodes_D_Residual = [c_nodes_Obs;c_nodes_HydrantObs] - c_nodes_D([Junction_SelectIdx,Junction_Hydrant_Name]);
c_nodes_D_Residual = c_nodes_D_Residual';

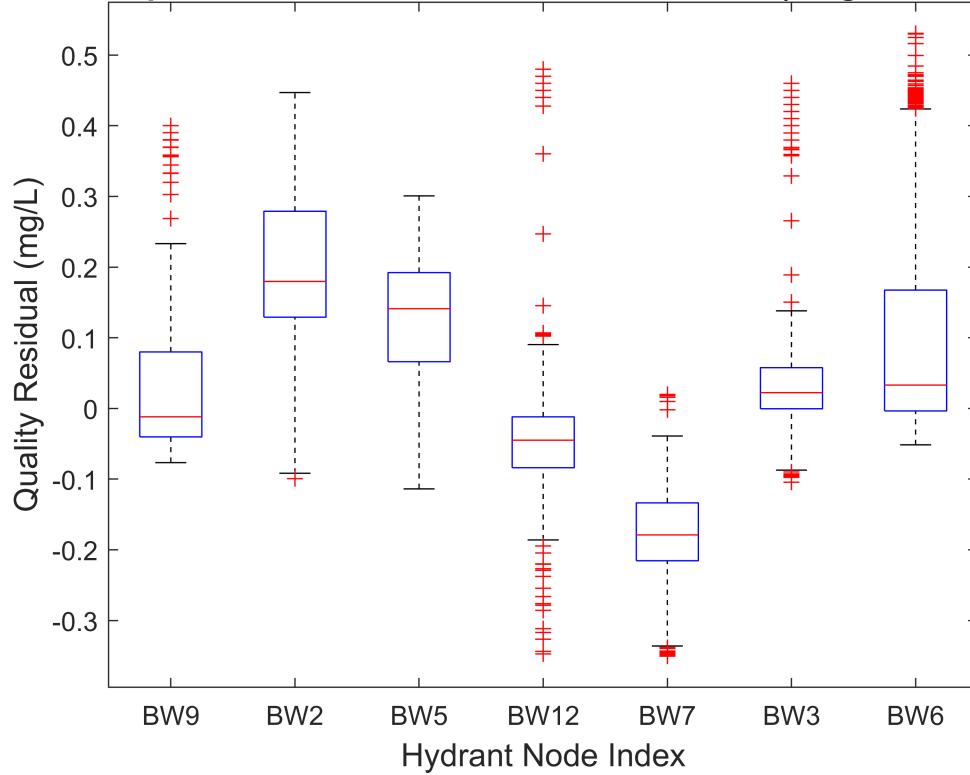
c_nodes_R_Residual = [c_nodes_Obs;c_nodes_HydrantObs] - c_nodes_R([Junction_SelectIdx,Junction_Hydrant_Name]);
c_nodes_R_Residual = c_nodes_R_Residual';

c_nodes_DandR_Residual = [c_nodes_Obs;c_nodes_HydrantObs] - c_nodes_DandR([Junction_SelectIdx,Junction_Hydrant_Name]);
c_nodes_DandR_Residual = c_nodes_DandR_Residual';

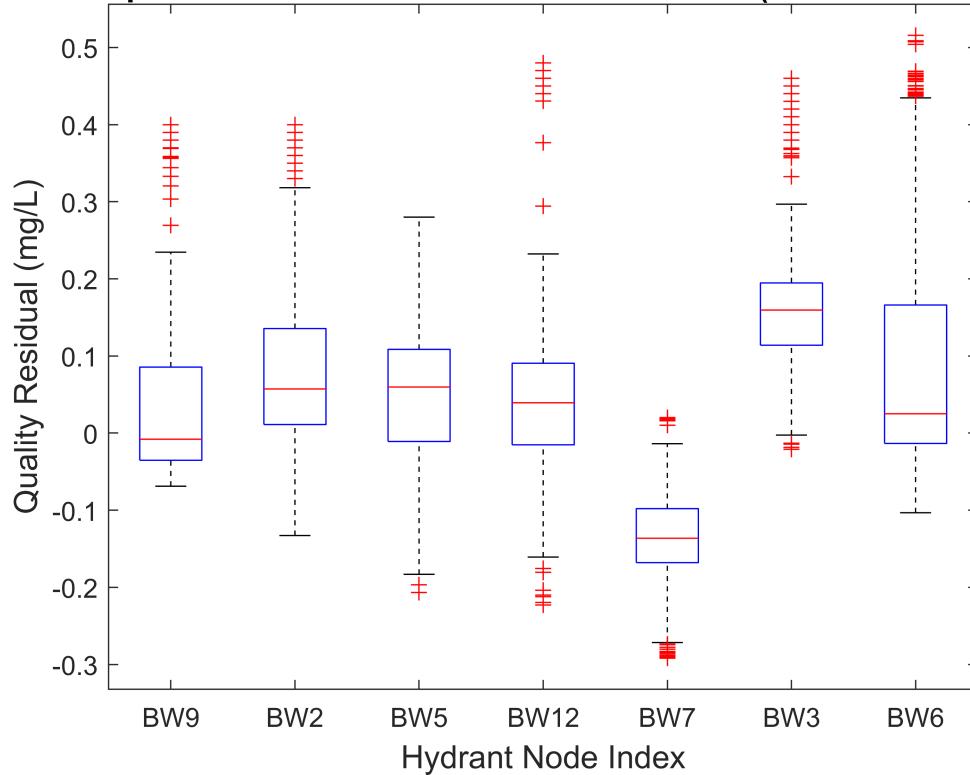
c_nodes_Residual = {c_nodes_NoGroup_Residual,c_nodes_D_Residual,c_nodes_R_Residual,... 
    c_nodes_DandR_Residual};
Residual_Name = {'(Single Group)', '(Diameter Based)', '(Roughness Based)', '(DR Based)'};

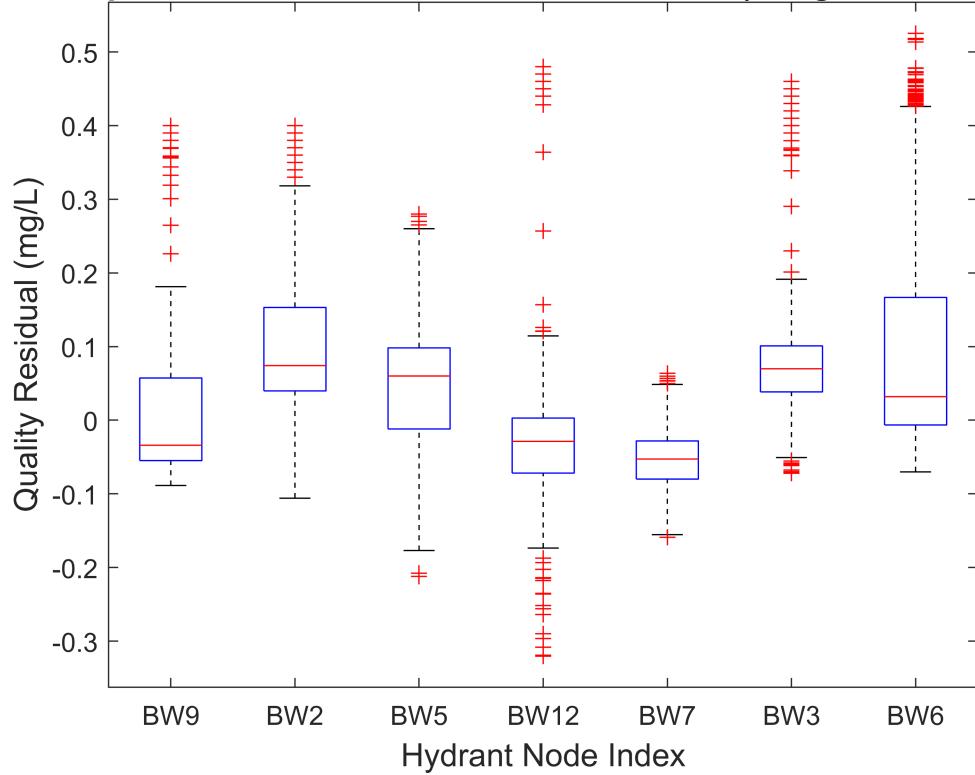
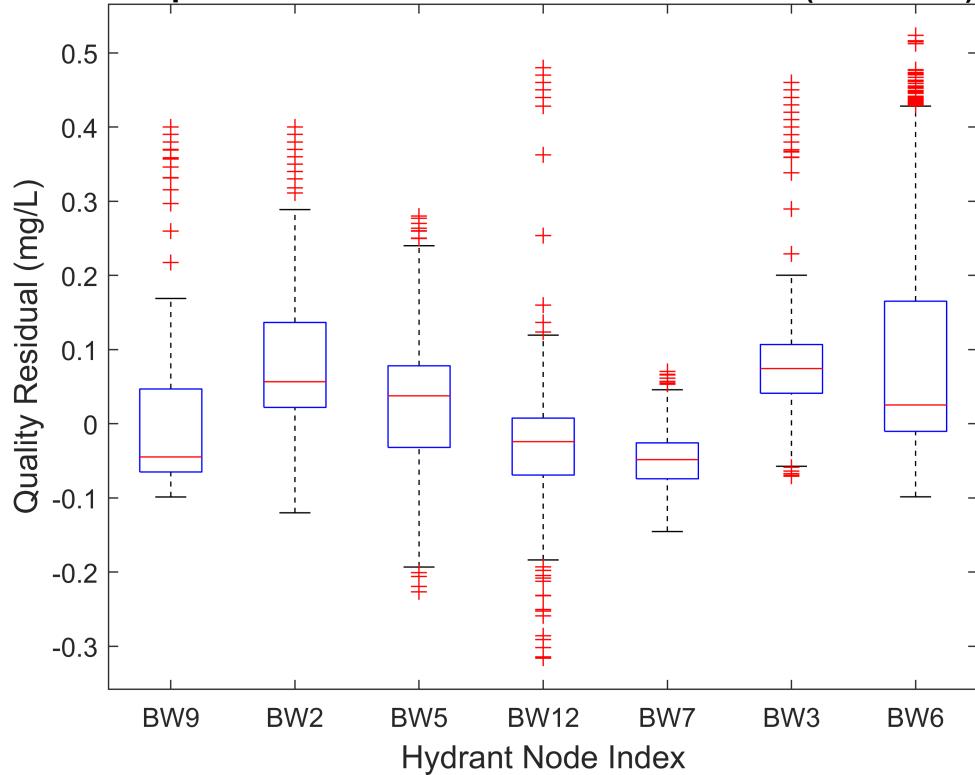
for i = 1:length(c_nodes_Residual)
    figure,
    boxplot(c_nodes_Residual{i});
    xlabel('Hydrant Node Index','FontSize',12)
    ylabel('Quality Residual (mg/L)','FontSize',12)
    title(['Boxplot of Chlorine Residuals at Main Nodes ',Residual_Name{i}], 'FontSize',12)
    xticklabels([Junction_Name,Junction_Hydrant_Name]);
end
```

Boxplot of Chlorine Residuals at Main Nodes (Single Group)



Boxplot of Chlorine Residuals at Main Nodes (Diameter Based)



Boxplot of Chlorine Residuals at Main Nodes (Roughness Based)**Boxplot of Chlorine Residuals at Main Nodes (DR Based)**

```
% Whole plot
```

```
c_nodes_Original_Residual = [c_nodes_Obs;c_nodes_HydrantObs] - c_nodes([Junction_SelectIdx,Junc
```

```

c_nodes_Original_Residual = c_nodes_Original_Residual';

c_nodes_Residual_combined = [c_nodes_Original_Residual,c_nodes_NoGroup_Residual,c_nodes_D_Residual
    c_nodes_R_Residual,c_nodes_DandR_Residual];

```

```

% figure,
% g2 = repmat(1:7,1,5);
% boxplot(c_nodes_Residual_combined,'ColorGroup',g2,'BoxStyle','filled');
% xlabel('Hydrant Node Index','FontSize',12)
% ylabel('Quality Residual (mg/L)','FontSize',12)
% title('Boxplot of Chlorine Residuals of different grouping method','FontSize',12)

Name_box = repmat({'BW9' 'BW2' 'BW5' 'BW12' 'BW7' 'BW3' 'BW6'},1,5);

simobs = [repmat({'O'},1,7),repmat({'S'},1,7),repmat({'D'},1,7),repmat({'R'},1,7),...
    repmat({'DR'},1,7)];
```

```

boxplot(c_nodes_Residual_combined,{Name_box,simobs}, 'Colors', repmat('ymcrg',1,7), ...
    'factorgap',[5 2], 'labelverbosity','minor', 'BoxStyle','filled');
xlabel('Node / Grouping method','FontWeight','bold')
ylabel('Chlorine residual error (mg/L)', 'FontWeight','bold')
title('Chlorine Residual Rrrors for All the Nodes Using Different Grouping Methods')
```

Chlorine Residual Rrrors for All the Nodes Using Different Grouping Methods

