

Exercise-A

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%% Load Data

% Load data from the provided CSV files into MATLAB

stable_data = readtable('/MATLAB Drive/all_stable - all_stable.csv'); % Stable nuclides data

stable2_data = readtable('/MATLAB Drive/all_stable2 - all_stable2.csv'); % Additional stable data with averages

unstable_data = readtable('/MATLAB Drive/all_unstable - all_unstable.csv'); % Unstable nuclides data


%% Define Column Names

% Assign column names for better code readability and reusability

neutron_column_stable = 'N'; % Column name for the number of neutrons

atomic_number_column = 'Z'; % Column name for atomic number (protons)

nat_abundance_column = 'Nat_abundance'; % Column name for natural abundance

neutron_avg_column = 'N_avg'; % Column name for average neutrons in stable2 data


%% Plot NZ1: Segre Chart of Known Nuclides

% Create a Segre chart displaying stable and unstable nuclides

figure;

% Plot stable nuclides in red squares

scatter(stable_data.(neutron_column_stable), stable_data.(atomic_number_column), 50, ...

    's', 'MarkerFaceColor', 'r', 'MarkerEdgeColor', 'k');

hold on;

% Plot neutron-rich unstable nuclides ( $N > Z$ ) in cyan triangles

neutron_rich = unstable_data(unstable_data.(neutron_column_stable) > unstable_data.(atomic_number_column), :);

scatter(neutron_rich.(neutron_column_stable), neutron_rich.(atomic_number_column), 50, ...

    '^', 'MarkerFaceColor', 'c', 'MarkerEdgeColor', 'w');

% Plot proton-rich unstable nuclides ( $N < Z$ ) in magenta inverted triangles

proton_rich = unstable_data(unstable_data.(neutron_column_stable) < unstable_data.(atomic_number_column), :);

scatter(proton_rich.(neutron_column_stable), proton_rich.(atomic_number_column), 50, ...

    'v', 'MarkerFaceColor', 'm', 'MarkerEdgeColor', 'w');
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% Plot symmetric radioactive isotopes (N = Z) as white circles

symmetric = unstable_data(unstable_data.(neutron_column_stable) == unstable_data.(atomic_number_column), :);

scatter(symmetric.(neutron_column_stable), symmetric.(atomic_number_column), 50, ...

    'o', 'MarkerFaceColor', 'w', 'MarkerEdgeColor', 'r');


% Add N/Z = 1 reference line

n_vals = 0:max([stable_data.(neutron_column_stable); unstable_data.(neutron_column_stable)]);

plot(n_vals, n_vals, 'g--', 'LineWidth', 1.75);


% Add titles, labels, and legend for NZ1

title('Segre Chart of Known Nuclides');

xlabel('Number of Neutrons (N)');

ylabel('Atomic Number (Z)');

grid on;

legend('Stable nuclides', 'Neutron-rich nuclides', 'Proton-rich nuclides', 'Symmetric RIs', 'N/Z = 1', 'Location', 'best');

hold off;


%% Average N/Z Calculation

% Calculate the average N/Z ratio for stable2 data

avg_NZ = mean(stable2_data.(neutron_avg_column) ./ stable2_data.(atomic_number_column));


% Determine the range of Z where N/Z is within 10% of 1

within_range = stable2_data( ...

    (stable2_data.(neutron_avg_column) ./ stable2_data.(atomic_number_column)) >= 0.9 & ...

    (stable2_data.(neutron_avg_column) ./ stable2_data.(atomic_number_column)) <= 1.1, :);

range_Z = [min(within_range.(atomic_number_column)), max(within_range.(atomic_number_column))];


% Find the maximum average N/Z ratio

max_NZ = max(stable2_data.(neutron_avg_column) ./ stable2_data.(atomic_number_column));


% Display average N/Z calculations in the Command Window

disp(['Average N/Z: ', num2str(avg_NZ)]);

disp(['Range where N/Z is within 10% of 1: Z = ', num2str(range_Z(1)), ' to Z = ', num2str(range_Z(2))]);

disp(['Maximum N/Z ratio: ', num2str(max_NZ)]);

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%% Add Max Average N/Z Line to NZ1

% Plot a line corresponding to the maximum average N/Z ratio on the Segre chart

hold on;

plot(n_vals, n_vals * max_NZ, 'b-', 'LineWidth', 1.75);

hold off;

%% Plot NZ2: N/Z vs. Z

% Create a plot showing the N/Z ratio against the atomic number Z

figure;

% Plot N/Z without considering natural abundance

plot(stable2_data.(atomic_number_column), stable2_data.(neutron_avg_column) ./ stable2_data.(atomic_number_column), ...

    'r-', 'LineWidth', 1.5);

hold on;

% Plot N/Z considering natural abundance

weighted_NZ = stable2_data.N_avg_weighted_by_abundance ./ stable2_data.(atomic_number_column);

plot(stable2_data.(atomic_number_column), weighted_NZ, ...

    'b--', 'LineWidth', 1.5);

% Customize line style for N/Z without abundance

set(findobj(gca, 'Type', 'line', 'Color', 'r'), 'LineStyle', '-');

% Add titles, labels, and legend for NZ2

title('N/Z vs Z (Step 6 & 7)');

xlabel('Atomic Number (Z)');

ylabel('N/Z Ratio');

grid on;

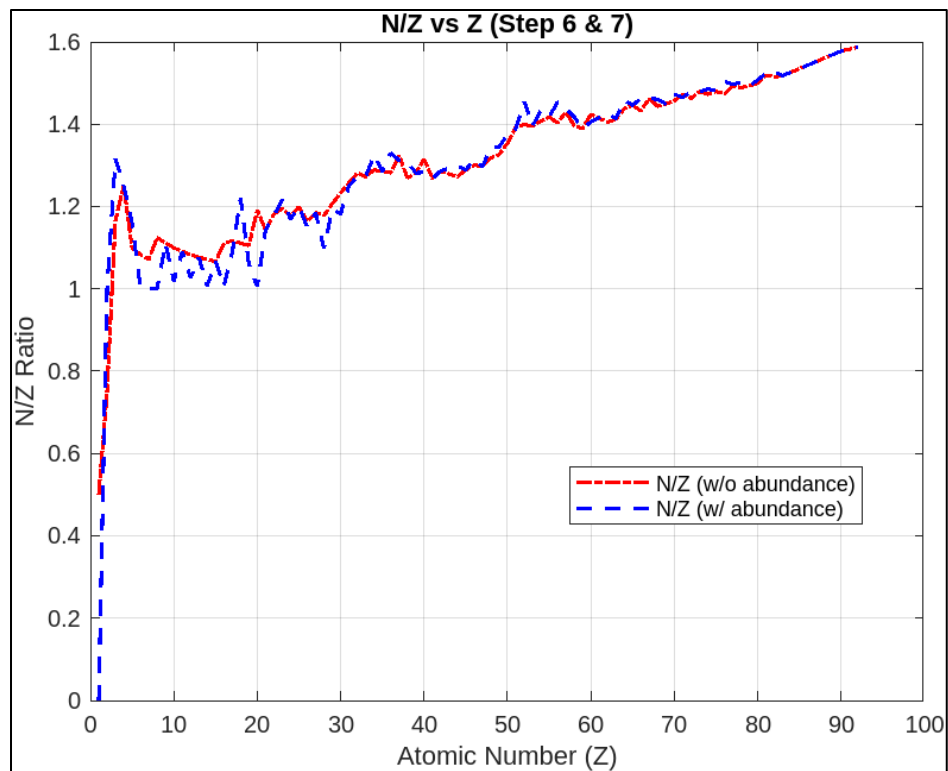
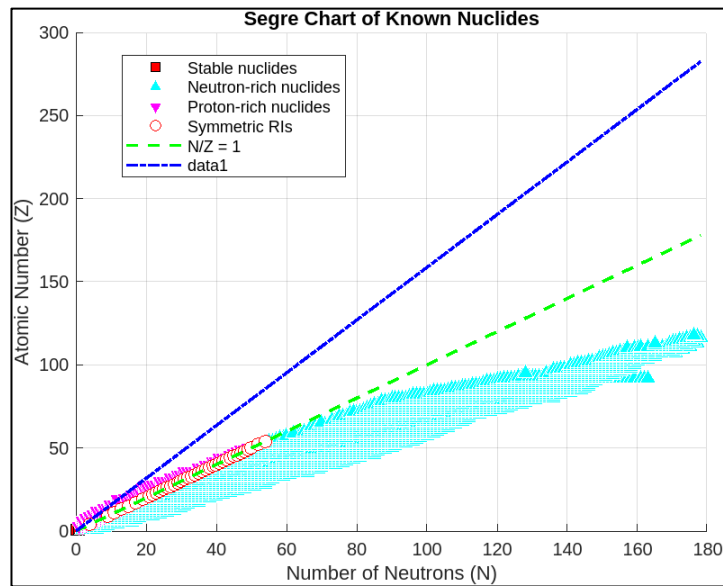
legend('N/Z (w/o abundance)', 'N/Z (w/ abundance)', 'Location', 'best');

hold off;

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Command Window

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Average N/Z: 1.2923  
Range where N/Z is within 10% of 1: Z = 5 to Z = 15  
Maximum N/Z ratio: 1.587  
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Exercise-B:

%Code for Calculating Mass Defect, Binding Energy, and BE/A Curve

% Conversion factor: 1 atomic mass unit (amu) to MeV

amu_to_MeV = 931.5;

% Given masses in atomic mass units (amu)

m_U235 = 235.043924; % Mass of Uranium-235

m_n = 1.008665; % Mass of a neutron

m_Rb93 = 92.91699; % Mass of Rubidium-93

m_Cs140 = 139.90910; % Mass of Cesium-140

m_n_products = 3 * m_n; % Total mass of 3 emitted neutrons

% Compute total mass of reactants and products

mass_reactants = m_U235 + m_n; % Mass of Uranium-235 and one neutron

mass_products = m_Rb93 + m_Cs140 + m_n_products; % Mass of fission products and neutrons

% Calculate mass defect (Δm)

mass_defect = mass_reactants - mass_products;

% Compute binding energy (B.E.) in MeV

binding_energy = mass_defect * amu_to_MeV;

% Calculate binding energy per nucleon (B.E./A)

A = 235; % Total number of nucleons in Uranium-235

binding_energy_per_nucleon = binding_energy / A;

% Display results

disp('Mass defect (amu):');

disp(mass_defect);

disp('Binding energy (MeV):');

disp(binding_energy);

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disp('Binding energy per nucleon (MeV):');
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disp(binding_energy_per_nucleon);
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% Generate and plot the Binding Energy per Nucleon (B.E./A) Curve
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atomic_numbers = 1:240; % Range of mass numbers for hypothetical nuclides
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binding_energies = 15.75 .* atomic_numbers - 17.8 * (atomic_numbers.^(2/3)); % Binding energy model
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binding_energies = binding_energies ./ atomic_numbers; % Normalize to calculate B.E./A
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Command Window

Mass defect (amu):
    0.2005

Binding energy (MeV):
    186.7695

Binding energy per nucleon (MeV):
    0.7948

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