	Boxed Particles The purpose of this exam specification is to implement a 2D simulation of elastic collisions between circular particles inside a rectangular box. The simulation includes collisions with the walls of the box and collisions between the particles themselves. The main objective of the assignment is to study the energy evolution of the system and to analyze the particle velocity and height distributions. In the code, the positions and velocities of the particles inside the box are initialized. The particles are then evolved over time under the effects of gravity and collisions, while particle positions, velocities and average kinetic energies are measured. Average energies over time are plotted and an animation is created to visualize particle motion.
In []:	Utility Class The ProgressBar class is a utility for displaying a progress bar in the console to track the progress of a long-running operation. It's initialized with the total number of iterations that the operation will take. Each time the step method is called, it increments an internal counter and updates the display of the progress bar. import sys import time as clock
	class ProgressBar: """ A class used to represent a progress bar in the console. Attributes total_iterations: int The total number of iterations that the operation will take.
	current_iteration : int The current iteration of the operation. start_time : float The time when the operation started. Methods
	definit(self, total_iterations): Constructs all the necessary attributes for the progress bar object. Parameters total iterations : int
	The total number of iterations that the operation will take. """ self.total_iterations = total_iterations self.current_iteration = 0 self.start_time = clock.time() def step(self): """ Increments the current iteration and updates the progress bar. If the operation is complete, it also prints the total time taken for the operation.
	<pre>self.current_iteration += 1 self.show_progress() if self.current_iteration == self.total_iterations: self.end_time = clock.time() print(f'\nTotal time taken: (self.end_time - self.start_time) seconds') def show_progress(self): """ Updates the display of the progress bar in the console.</pre>
	The progress bar is displayed as a line of 20 characters, with '=' characters representing completed iterations and ' ' characters representing remaining iterations. The percentage of completion is also displayed next to the progress bar. """ percent_complete = self.current_iteration / self.total_iterations num_bars = int(percent_complete * 20) # 20 bars on the progress bar sys.stdout.write('\r') sys.stdout.write('\r') sys.stdout.write("\r') sys.stdout.flush() sys.stdout.flush()
	 NumPy: NumPy is one of the fundamental libraries for scientific computing in Python. It provides a flexible and powerful multidimensional array object, as well as tools for working with these arrays. In this code, it is used for vector and matrix calculus, random number generation, and other mathematical operations. This library makes your code more readable and optimized. Matplotlib.pyplot: matplotlib.pyplot is a module in the matplotlib library in Python. It's a collection of functions that provides a MATLAB-like interface for creating visualizations. Each pyplot function makes some change to a figure, such as creating a figure, creating a plotting area in a figure, plotting some lines in a plotting area, decorating the plot with labels, and so on. Matplotlib.animation: This is a submodule of Matplotlib that provides tools for creating animations that can be embedded in Jupyter notebooks, saved as GIF or MP4 files, or displayed directly in a view window. In the code, it is used to animate the temporal evolution of the position
In []:	• Matplottlib inline import numpy as np import numpy nas np import numpy n
	import matplotlib. animation as animation import matplotlib. patches as mpatches from IPython.display import HTML, clear_output Translating Physics into Code Particle Movement
	In this simulation, the motion of particles is computed using a time-stepping approach with considerations for collisions and gravity . The system starts by initializing particles' positions and velocities randomly within the defined box. Each step of the simulation involves updating the positions of particles based on their current velocities and the effects of gravity. The position update is done using the equations of motion under constant acceleration, where the only acceleration considered is due to gravity. The velocity is updated by considering the gravitational acceleration. The equations used are: $\begin{cases} x(t+\Delta t) &= x(t)+v_x(t)\Delta t\\ y(t+\Delta t) &= y(t)+v_y(t)\Delta t-\frac{1}{2}g(\Delta t)^2 \end{cases}$ $\begin{cases} v_x(t+\Delta t) &= v_x(t)\\ v_y(t+\Delta t) &= v_y(t)-y_y(t)-y_z(t) = v_y(t)-y_z(t) = v_y(t) $
	Finally, the system state is sampled at specific intervals to observe the motion and other properties (like kinetic and potential energy) of the particles over time. The system also provides an animation method to visualize the motion of the particles. Particle Collision The handle_collisions_between_particles function is called to handle collisions between all particles in the system. The collisions are <i>treated as elastic</i> , meaning that the total kinetic energy of the system is conserved. The new velocities after the collision are calculated using the conservation of momentum and kinetic energy , resulting in the following equations:
	$\vec{v}_1' = \vec{v}1 - \frac{2M_2}{M_1 + M_2} \frac{(\vec{v}_{rd} \cdot \vec{r}_{red})}{(\vec{r}_{rd} \cdot \vec{r}_{red})} \vec{v}_{red} \cdot \vec{v}_{red}' = \vec{v}2 + \frac{2M_1}{M_1 + M_2} \frac{(\vec{v}_{rd} \cdot \vec{r}_{red})}{(\vec{r}_{rd} \cdot \vec{r}_{red})} \vec{r}_{red}$ These formulas reflect the fact that the colliding disks repel each other along the direction which joins their centers. The formulas are valid for arbitrary masses M_1, M_2 , and of course simplify a bit for those collisions involving particles with equal mass. Wall Collision For each particle in the simulation, it checks if the particle's position on the x and y coordinates is outside the bounds of the box, taking into account the particle radius self.R. If a particle is found to be in contact with the wall, its velocity is reversed, simulating an elastic collision with the wall.
	Energy Considerations In the simulation, the average kinetic and potential energies of each species are calculated at each sampling step. The kinetic energy is calculated using the formula: $K_i = \frac{1}{2} M_i v_i^2 = \frac{1}{2} M_i \left(\vec{v}_i \cdot \vec{v}_i \right)$ and the potential energy is calculated using the formula:
In []:	$U_i = M_i g y_i \tag{2}$ Full Code for the Simulation
	definit(self, N_A, N_B, M_A, M_B, R, v_0, seed=None): Initialize the simulation. Parameters N_A : int Number of particles of type A. N_B : int
	Number of particles of type B. M.A: float Mass of particles of type A. M.B: float Mass of particles of type B. R: float Radius of the particles. V.O: float Maximum initial velocity of the particles. seed: int, optional
	Random'seed for reproducibility. """ self.N_A = N_A self.N_B = N_B self.M_A = M_B self.M_B = M_B self.M_B = M_B self.Y_B = N_B self.Y_B = N
	np.random.seed(seed) self.box_width = 8 self.box_height = 16 self.initialize_particles() def initialize_particles(self): "" Initialize the positions and velocities of all particles.
	The particles are randomly distributed within the box, and their initial velocities are also chosen randomly. """ total_particles = self.N_A + self.N_B self.positions = np.random.uniform(low=0, high=self.box_width, size=(total_particles, 2)) self.positions[:, 1] = np.minimum(self.positions[:, 1], 8) # Limit y position to 8 self.velocities = np.random.uniform(low=-self.v_0, high=self.v_0, size=(total_particles, 2)) self.mass = np.concatenate([np.full(self.N_A, self.M_A), np.full(self.N_B, self.M_B)])
	self.colors = np.concatenate([np.full(self.N_A, 'blue'), np.full(self.N_B, 'red')]) def evolve(self, T, dt, dt_s): """ Evolve the simulation for a given time period. Parameters
	dt: float Time step for the evolution of the system. dt_s: float Time step for the evolution of the system. dt_s: float Time step for sampling the system's state. Returns tuple Tuple containing arrays of positions and velocities samples and average kinetic and potential energy of particles A and B. """ Netnos = int(I / dt)
	<pre>N_steps = int(T / dt) N_samples = int(T / dt_s) positions_samples = np.zeros((N_samples, self.N_A + self.N_B, 2)) velocities_samples = np.zeros((N_samples, self.N_A + self.N_B, 2)) avg_kinetic_energy_A = np.zeros(N_samples) avg_kinetic_energy_B = np.zeros(N_samples) avg_potential_energy_A = np.zeros(N_samples) avg_potential_energy_B = np.zeros(N_samples) pb = ProgressBar(N_steps) pb = ProgressBar(N_steps)</pre>
	<pre>pu = rtogresser(m_steps) for step in range(N_steps): self.update_positions(dt) self.handle_collisions_with_walls() self.handle_collisions_between_particles() if step % (dt_s / dt) == 0: sample_index = int(step / (dt_s / dt)) positions_sample(spample_index) = self.positions.copy() velocities_samples[sample_index] = self.velocities.copy()</pre>
	avg_kinetic_energy_A[sample_index] = self.get_average_kinetic_energy(self.N_A, self.M_A) avg_kinetic_energy_B[sample_index] = self.get_average_kinetic_energy(self.N_A, self.M_B) avg_potential_energy_A[sample_index] = self.get_average_potential_energy(self.N_A, self.M_B) avg_potential_energy_B[sample_index] = self.get_average_potential_energy(self.N_B, self.M_B) pb.step() return positions_samples, velocities_samples, avg_kinetic_energy_A, avg_kinetic_energy_B, avg_potential_energy_A, avg_potential_energy_B def update_positions(self, dt):
	Update the positions of all particles. Parameters dt : float Time step for the position update. """ self.positions += self.velocities * dt self.positions[:, 1] -= 0.5 * self.g * dt**2
	<pre>self.velocities[:, 1] -= self.g * dt def handle_collisions_with_walls(self): """ Handle collisions of particles with the walls of the box. """ self.velocities[(self.positions - self.R < 0) (self.positions + self.R > self.box_width)] *= -1 def handle_collisions_between_particles(self): """</pre>
	<pre>Handle collisions between particles. """ distances = self.distFast(self.positions) for i in range(len(self.positions)): for j in range(i+1, len(self.positions)): if distances[i, j] = 2 * self.R: self.handle_particle_collision(i, j) def distFast(self, pos):</pre>
	Calculate the pairwise distance between points for an array of points. Parameters pos: numpy.ndarray Array containing the positions of the points. Returns
	<pre>numpy.ndarray Array containing the pairwise distances. """ R = np.sqrt(np.einsum('ijl->ij', (pos[:,np.newaxis,:] - pos[np.newaxis,:,:])**2.)) return R def handle_particle_collision(self, i, j): """ Handle collision between two particles.</pre>
	Parameters i: int Index of the first particle. j: int Index of the second particle. """ ri = self.positions[i] rj = self.positions[j] vi = self.velocities[i] vj = self.velocities[i]
	<pre>vj = self.mass[j] mi = self.mass[j] rij = ri - rj vij = vi - vj rij_dot_vij = np.dot(rij, vij) rij_dot_vij = np.dot(rij, rij) vi_prime = vi - (2 * mj * rij_dot_vij / ((mi + mj) * rij_dot_rij)) * rij vj_prime = vj + (2 * mi * rij_dot_vij / ((mi + mj) * rij_dot_rij)) * rij</pre>
	self.velocities[i] = vi_prime self.velocities[j] = vj_prime def get_average_kinetic_energy(self, N_particles, mass): """ Calculate the average kinetic energy of a set of particles. Parameters
	N_particles: int Number of particles. mass: float Mass of the particles. Returns float Average kinetic energy of the particles.
	<pre>velocity_squared = np.sum(self.velocities[:N_particles]**2, axis=1) total_kinetic_energy = 0.5 * mass * np.sum(velocity_squared) return total_kinetic_energy / N_particles def get_average_potential_energy(self, N_particles, mass): """ Calculate the average potential energy of a set of particles. Parameters</pre>
	N_particles: int Number of particles. mass: float Mass of the particles. Returns float Average potential energy of the particles.
	total_potential_energy = mass * self.g * np.sum(self.positions[:N_particles, 1]) return total_potential_energy / N_particles def animate(self, positions_samples, interval, frame_number): """ Create an animation of the motion of the particles. Parameters
	<pre>positions_samples : numpy.ndarray Array containing the positions of all particles at different time steps. interval : int Time interval between frames in the animation. """ if (frame_number < 0): frame_number = len(positions_samples) fig, ax = plt.subplots() ax.set_xlim(0, self.box_width) ax.set_xlim(0, self.box_width) ax.set_xlim(0, self.box_width)</pre>
	# Initialize scatter plot with initial positions and colors initial_positions = positions_samples[0] colors = [particle for particle in self.colors] particles_scatter = ax.scatter(initial_positions[:, 0], initial_positions[:, 1], c=colors, s=10) # Add legend outside of plot blue_patch = mpatches.Patch(color='blue', label='Species A') red_patch = mpatches.Patch(color='red', label='Species B') ax.legend(handles=[blue_patch, red_patch, red_patch, loc='center left', bbox_to_anchor=(1, 0.5))
	<pre>pb = ProgressBar(frame_number) # Start progress bar def update(frame): x = positions_samples[frame, :, 0] y = positions_samples[frame, :, 1] particles_scatter.set_offsets(np.c_[x, y]) pb.step()</pre>
	return particles_scatter, anim = animation.FuncAnimation(fig, update, frames=frame_number, interval=interval, blit=True) # Use to_html5_video to convert the animation to a JavaScript HTML video html_vid = anim.to_html5_video() # Clean up the plot plt.close(fig) # Use IPython.display.HTML to display the animation
In []:	Parameters Initialization N.A = 30 # Number of type A particles N.B = 30 # Number of type B particles M.A = 0.025 # Mass of type A particles (kg) M.B = 0.025 # Mass of type B particles (kg)
	R = 0.04 # Radius of the particles (m) v_0 = 0.5 # Initial velocity of the particles (m/s) seed = 42 # Random seed for particles's position generation Avarage Energies The average kinetic and potential energies for particles of types A and B are computed at each time step during the simulation. The kinetic energy of a particle is given by the equation $\frac{1}{2}mv^2$, where m is the mass of the particle and v is its velocity. The potential energy of a particle in a gravitational field is given by mgh (in this simulation mgh), where m is the mass, g is the acceleration due to gravity, and h is the height or vertical position of the particle. These energy values are calculated for each particle, summed up , and then divided by the total number of particles.
	to get the average energy. The calculation is done for both particle types A and B, and the results are stored in arrays avg_kinetic_energy_A, avg_kinetic_energy_B, avg_potential_energy_A, and avg_potential_energy_B. These arrays are then plotted against time to see how the average energies of the particles evolve over the course of the simulation. # Create the simulation simulation = ParticleSimulation(N_A, N_B, M_A, M_B, R, v_0, seed) T = 100 # Total time for simulation
	dt_s = 0.04 # Sampling time step dt = 0.081 positions_samples, velocities_samples, avg_kinetic_energy_A, avg_potential_energy_A, avg_potential_energy_B = simulation.evolve(T, dt, dt_s) times = np.arange(0, T, dt_s) plt.figure(figsize=(10,6)) plt.plot(times, avg_kinetic_energy_A, label='Kinetic Energy A', color='blue') plt.plot(times, avg_kinetic_energy_B, label='Kinetic Energy B', color='red')
	plt.plot(times, avg_potential_energy_A, label='Potential Energy A', color='cyan') plt.plot(times, avg_potential_energy_B, label='Potential Energy B', color='magenta') plt.legend(loc='upper right') plt.label('Time') plt.ylabel('Time') plt.ylabel('Time') plt.title('Average Energies Over Time') plt.show() Considerations
	The plot of average kinetic and potential energy over time should provide key insights into the dynamical properties of the particle system under simulation. The kinetic energy, which is associated with the motion of the particles, is expected to fluctuate over time as the particles with each other and the walls of the box, thereby exchanging energy. Depending on the initial conditions and particle properties (mass, radius, initial velocity), the kinetic energy may trend towards an equilibrium value or exhibit chaotic behavior. The potential energy of a particle in this simulation is associated with its height in the box due to gravity. The potential energy should decrease when a particle descends and increase when a particle ascends. If there is no loss of energy (e.g., due to friction), the sum of the kinetic and potential energy for each type of particle should remain constant over time (the law of conservation of energy). By examining the relative changes in these energies, we can understand various phenomena, such as how energy is transferred between different types of particles (A and B) and how the system's behavior evolves over time. This kind of analysis is fundamental in statistical mechanics and thermodynamics, where one is often interested in the macroscopic behavior of systems composed of a large number of particles.
	Position and Velocity Plots The positions and velocities of all particles at specific time points are plotted using a quiver plot . A quiver plot displays vector fields, which are useful in this case for visualizing the direction and magnitude of the particle's position at the given time is represented by a point on the plot, and its velocity is represented by an arrow originating from that point. The <i>direction</i> of the arrow indicates the direction of the particle's velocity , and the <i>length of the arrow</i> represents the magnitude of the velocity. The plots are color-coded to distinguish between particle types A and B. These plots provide a visual understanding of how the particles' positions and velocities change over time. The position plot helps visualize the distribution and movement of the particles in the box, while the velocity plot helps visualize the speed and direction of the particles.
In []:	<pre>times_to_plot = np.linspace(0, len(positions_samples) - 1, 10).astype(int) # 10 points from 0 to 100% of simulation time fig, axs = plt.subplots(3, 4, figsize=(15,10)) # Create a 3x4 grid of subplots for i, ax in enumerate(axs.flat): # If we've reached the end of our data, remove the extra subplots and break the loop if i>= len(times_to_plot): fig.delaxes(ax) continue</pre>
	time = times_to_plot[i] positions = positions_samples[time] velocities = velocities_samples[time] colors = simulation.colors ax.quiver(positions[:, 0], positions[:, 1], velocities[:, 0], velocities[:, 1], color=colors, scale=1, scale_units='xy') ax.set_xlim(0, simulation.box_width) ax.set_ylim(0, simulation.box_height) ax.set_title(f'Iteration: {time}')
	plt.tight_layout() plt.show() Considerations The plot presented here is a series of quiver plots illustrating the progression of the particle simulation over time. These plots will provide a visual representation of both the positions and velocities of particles at various points in time. At each instance, the quiver plot shows the particles of both species, with their location in the box corresponding to their spatial position. The arrows originating from each point represent the direction and magnitude of the particle's velocity. The length of the arrow corresponds to the
	speed of the particle, and the direction of the arrow indicates the direction of movement. We expect to observe a few things from these plots. Initially, when the simulation just starts, the particles might be distributed unevenly and have high velocities (long arrows), depending on the initial conditions. As the simulation progresses, we hope to see the system approach equilibrium. This would manifest as the particles becoming more evenly distributed across the space, and their velocities becoming more uniform (arrows of similar length). By the end of the simulation, if equilibrium has been achieved, we expect to see a uniform distribution of particles with a similar spread of velocities in various directions. Total Energy at Different Time Values
	The plots allow us to visualize how energy distributes in the system over time. The total energy plot should remain approximately constant if the simulation is accurate, since total energy should be conserved in an isolated system. We also monitor the average kinetic energies of the two species. According to the equipartition theorem, even if the species start with different average kinetic energies, they should become equal over time , showing that energy distributes <i>evenly among all degrees of freedom</i> in a system at equilibrium . The simulation checks when the average kinetic energies of the two species become equal within a certain tolerance (<i>here, 10%</i>), indicating the system has reached thermal equilibrium . This is represented as T_{therm} . This information helps us understand the time scale at which the system tends towards equilibrium and whether certain parameters (T , dt , v_0) affect this time scale.
	Finally, we take precautions to ensure accurate simulation results, particularly considering that the chosen speed should be small compared to the ratio of the size of the disks and the simulation step. Otherwise, the detection of collisions <i>could be inaccurate</i> , compromising the reliability of the simulation. # Simulation parameters T = 300 # Time interval (in seconds) dt_values = [0.804, 0.802, 0.801] # Simulation steps (in seconds) dt_s = 0.84 # Sampling step (in seconds) vol = 2 # Initial velocities (in m/s) time = np.arange(0, T, dt_s)
	tolerance = 0.1 # Tolerance for equilibrium check # Set initial conditions simulation = ParticleSimulation(N_A, N_B, M_A, M_B, R, v0, seed) for dt in dt_values: # Run simulation positions_samples, velocities_samples, avg_kinetic_energy_A, avg_potential_energy_A, avg_potential_energy_B = simulation.evolve(T, dt, dt_s)
	# Compute total energy total_energy = simulation.N_A * (avg_kinetic_energy_A + avg_potential_energy_A) + simulation.N_B * (avg_kinetic_energy_B + avg_potential_energy_B) # Plot total energy as a function of time plt.figure(figsize=(10, 4)) plt.subplot(1, 2, 1) plt.plot(time, total_energy) plt.title(f'Total Energy for T={T}s, dt={dt}s, and v0={v0}m/s') plt.title(f'Total Energy for T={T}s, dt={dt}s, and v0={v0}m/s')
	plt.ylabel('Total Energy (J)') # Plot average kinetic energy as a function of time for each species plt.subplot(1, 2, 2) plt.plot(time, avg_kinetic_energy_A, label='Species A') plt.plot(time, avg_kinetic_energy_B, label='Species B') plt.title(f'Average Kinetic Energy for T={T}s, dt={dt}s, and v0={v0}m/s') plt.xlabel('Time (s)') plt.ylabel('Average Kinetic Energy (J)') plt.lgend()
	<pre>plt.tight_layout() plt.show() equilibrium_reached = False for i in range(len(avg_kinetic_energy_A)): if np.abs(avg_kinetic_energy_A[i] - avg_kinetic_energy_B[i] < tolerance: print(f'Equilibrium reached at T_therm = {i^dt_s}s for T={T}s, dt={dt}s, and v0={v0}m/s') equilibrium_reached = True T_threm = i^dt_s # Found thermalization time break</pre>
	if not equilibrium_reached: print(f'Equilibrium not reached for T={T}s, dt={dt}s, and v0={v0}m/s.') T_threm = 100 # Default value Considerations The kinetic and potential energy plots provide crucial insights into the dynamics of the system we are simulating. Ideally, the plot of total energy should remain constant, as per the law of conservation of energy, which states that energy cannot be created or destroyed, but only changed.
	from one form to another. Any fluctuations we observe could be a result of numerical approximations used in the simulation. The effect of these approximations should diminish as we reduce the simulation time step of . The second plot, which demonstrates the average kinetic energy of each species over time, sheds light on the energy distribution within the system. As the system approaches thermal equilibrium, we expect the kinetic energies of the two species to become equal, signifying an equal distribution of energy. However, it's important to note that despite various attempts at altering the initial velocities and simulation time, equilibrium wasn't consistently achieved. This indicates that the two species aren't consistently reaching a state of equal energy distribution. This lack of equilibrium could be due to a variety of factors, including the inherent properties of the species (such as mass or particle count), initial conditions, or other factors not considered within the simulation. Identifying the point at which equilibrium is achieved, if at all, is key to understanding the timescales at which the system evolves. Thus, the inability to consistently reach equilibrium presents an interesting area for further investigation. Particles Animation
	This code will generate di animation of the particles A and B inside the box. # Set initial conditions simulation = ParticleSimulation(N_A, N_B, M_A, M_B, R, v_0, seed) T = 200 # Time interval (in seconds) dt = 0.004 # Simulation steps (in seconds)
	dt_s = 8.84 # Sampling step (in seconds) positions_samples, velocities_samples, avg_kinetic_energy_B, avg_potential_energy_A, avg_potential_energy_B = simulation.evolve(T, dt, dt_s) # Animate the motion of particles interval = 59 HTML(simulation.animate(positions_samples, interval,-1)) Velocity and Height Distributions
T	The provided code implements the calculation and visualization of velocity and height distributions for two species of particles in a simulation. The simulation is run for an extended period of time, including a thermalization phase and additional 500 seconds. After reaching equilibrium (if it's reached), the velocities of the particles are used to compute the squared velocity values, which are then used to populate histograms for each particle species. Similarly, the positions of the particles are used to calculate histograms for the height distributions. The code utilizes the numpy.histogram function to compute the histograms and matplotlib.pyplot.hist to visualize the distributions. The resulting histograms provide insights into the velocity and height distributions, which are expected to resemble the Maxwell-Boltzmann distributions observed in gas particles. The value of kT, representing the average kinetic energy, is used in the calculation of the exponential probabilities for the velocity and height distributions. The code allows for exploration of different initial parameters, such as v ₀ , to observe the effects on the distributions.
In []:	# Changing the time for the simulation T_tot = 180 + 500 # Total time positions_samples, velocities_samples, avg_kinetic_energy_A, avg_kinetic_energy_B, avg_potential_energy_A, avg_potential_energy_B = simulation.evolve(T_tot, dt, dt_s) # Calculate kT kT = 0.5 * avg_kinetic_energy_A # Compute velocity histograms hist A, bins_A = np. histogram(velocities_samples[:, 0, :], bins='auto', density=True) bist_B bins_B = np. histogram(velocities_samples[:, 1, :], bins='auto', density=True) bist_B bins_B = np. histogram(velocities_samples[:, 1, :], bins='auto', density=True)
	hist_B, bins_B = np.histogram(velocities_samples[:, 1, :], bins='auto', density=True) # Compute height histograms hist_height_A = np.histogram(positions_samples[:, 0, 1], bins='auto', density=True) hist_height_B, bins_height_B = np.histogram(positions_samples[:, 1, 1], bins='auto', density=True) # Create a figure and a 1x2 array of subplots fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(16, 6)) # Plot velocity histograms on the first subplot
	axi.hist(bins,A[:-1], bins,A, weights=hist,A, alpha=0.5, label='Type A') axi.hist(bins,B[:-1], bins,B, weights=hist,B, alpha=0.5, label='Type B') axi.set_xlabel('Velocity Squared') axi.set_tlite('Velocity Distribution') axi.set_tlite('Velocity Distribution') axi.legend() # Plot height histograms on the second subplot ax2.hist(bins,height,A[:-1], bins_height,A, weights=hist_height,B, alpha=0.5, label='Type A') ax2.hist(bins,height,B[:-1], bins_height,B, weights=hist_height,B, alpha=0.5, label='Type B')
	ax2.set_xlabel('Height') ax2.set_ylabel('Probability Density') ax2.set_title('Height Distribution') ax2.slegend() # Show the plots plt.tight_layout() # Adjusts subplot params so that subplots are nicely fit in the figure. plt.show() Considerations
	Considerations The plots for velocity and height distributions should give us important information about the state of our simulated system and how the particles of the two species are behaving. The velocity distribution plot is expected to be a reflection of the Maxwell-Boltzmann distribution. At thermal equilibrium, we would expect the distribution of velocities for each species to form a bell curve shape, centered around a peak velocity that reflects the most probable speed of particles in the system. This distribution arises from the principle that in a gas at thermal equilibrium, the velocities of particles are distributed according to their kinetic energies. Similarly, the height distribution should provide us with an insight into the spatial distribution of heights. If the system is not at equilibrium, then the particles might be congregated in specific regions, leading to a non-uniform distribution of heights.
	It's important to note that these are ideal expectations. Depending on the specifics of the simulation, such as the properties of the two species and the initial conditions, the actual distributions may differ. This type of analysis can provide valuable insights into the timescales over which equilibrium is reached, and the factors that might speed up or slow down the process.