```
1.import random
```

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# Pre-defined evidence (modify as needed)
test_results = {"WBC Count": 15000, "Imaging": "Possible appendicitis"}
intraoperative_findings = "Bleeding encountered during initial incision"
expert_consultation = "Proceed cautiously, consider potential for atypical presentation"
# Decision point and answer options
decision prompt = "Based on the available evidence, what is the best course of action?"
answer_options = [
   "Continue with the laparoscopic appendectomy as planned.",
   "Convert to an open procedure to gain better visibility.",
   "Consult with another surgical specialist for further guidance."
1
def simulate experiment():
  # Present background information
  print("Simulating an appendectomy...")
  print("Background: Patient presents with classic appendicitis symptoms.")
  # Present evidence
  print("Pre-operative Test Results:", test_results)
  print("Intra-operative Findings:", intraoperative_findings)
  print("Expert Consultation:", expert_consultation)
  # Present decision point
  print(decision_prompt)
  for i, option in enumerate(answer_options):
     print(f"{i+1}. {option}")
  # Simulate user input (replace with actual user input mechanism)
  user_choice = input("Select an option (1, 2, 3): ") # Modified to accept user input
  while user choice not in ['1', '2', '3']:
     print("Invalid choice. Please select 1, 2, or 3.")
     user choice = input("Select an option (1, 2, 3): ")
  user choice = int(user choice)
  print("Selected option:", user choice)
  # Prompt for justification
  justification = input("Explain your reasoning for this choice: ")
  print("Justification:", justification)
  evaluate justification(justification, user choice)
def evaluate_justification(justification, user_choice):
  # Implement your evaluation logic here
  # For simplicity, let's provide basic feedback based on the chosen option
  if user choice == 1:
     print("Your choice: Continue with the laparoscopic appendectomy.")
     print("Feedback: Proceeding cautiously with the planned procedure.")
   elif user choice == 2:
     print("Your choice: Convert to an open procedure.")
     print("Feedback: Opting for better visibility due to encountered bleeding.")
   elif user choice == 3:
     print("Your choice: Consult with another surgical specialist.")
     print("Feedback: Seeking further guidance due to uncertain circumstances.")
# Run the simulation
simulate experiment()
```

```
2.import pandas as pd
from scipy import stats
# Data
fertilizer_conc = ["Control", "Low", "Medium", "High"]
plant_height = [10, 12, 15, 18] # Average height for each group
# Create dataframe
data = pd.DataFrame({"Fertilizer": fertilizer_conc, "Height": plant_height})
# Descriptive statistics for each group
print(data.groupby('Fertilizer')['Height'].describe())
# Hypothesis test (ANOVA)
anova_results = stats.f_oneway(
  data[data['Fertilizer'] == "Control"]["Height"],
  data[data['Fertilizer'] == "Low"]["Height"],
  data[data['Fertilizer'] == "Medium"]["Height"],
  data[data['Fertilizer'] == "High"]["Height"]
# Print ANOVA results
print("ANOVA p-value:", anova_results.pvalue)
# Interpretation based on results
if anova_results.pvalue < 0.05: # Adjust significance level as needed
  print("There is a statistically significant difference in plant height between groups. Further analysis is needed to identify
specific relationships between fertilizer concentration and plant growth.")
  print("There is no statistically significant evidence that fertilizer concentration affects plant height in this data set.")
3.import random
# Define events
event_A = "It will rain today"
event_B = "The bus will be late"
# Assign probabilities to events (P(A) and P(B))
probability_A = 0.6
probability_B = 0.4
# Function to calculate the probability of an event (P(X))
def get_probability(event):
  if event == event_A:
     return probability A
  elif event == event_B:
     return probability_B
  else:
     print("Invalid event")
     return None
# Simulate 1000 trials
trials = 1000
rain count = 0
late_bus_count = 0
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both_count = 0
for _ in range(trials):
  # Simulate events happening
  if random.random() < probability_A:
     rain_count += 1
  if random.random() < probability_B:
     late bus count += 1
  if rain_count and late_bus_count:
     both_count += 1
# Estimated probabilities based on simulations
estimated_probability_A = rain_count / trials
estimated_probability_B = late_bus_count / trials
estimated_intersection_probability = both_count / trials
estimated union probability = estimated probability A + estimated probability B - estimated intersection probability
# Print results
print("Event A:", event_A)
print("Estimated P(A):", estimated_probability_A)
print("Event B:", event_B)
print("Estimated P(B):", estimated_probability_B)
print("Estimated P(A and B):", estimated_intersection_probability)
print("Estimated P(A or B):", estimated_union_probability)
4.def analyze_believability(claim, source, sensationalism=0):
  This function analyzes the believability of a claim based on source and sensationalism.
  Args:
     claim (str): The textual claim to be analyzed.
     source (str): The source of the claim (e.g., news website name).
     sensationalism (int, optional): A score indicating the level of sensational language (default 0).
  Returns:
    str: A basic believability classification (e.g., "Likely Believable", "Needs Verification").
  source credibility = {"Reputable News": 1, "Social Media": 0.5, "Anonymous": 0.2}
  credibility_score = source_credibility.get(source, 0.5) # Default 0.5 for unknown sources
  # Adjust score based on sensationalism (more exclamation points = less believable)
  credibility score -= sensationalism * 0.1
  if credibility score > 0.7:
     return "Likely Believable"
  elif credibility_score > 0.4:
     return "Needs Verification"
  else:
     return "Likely Unbelievable"
# Example usage
claim = "Giant lizards discovered living in the Amazon rainforest!"
source = "Unknown Social Media Post"
believability = analyze believability(claim, source, sensationalism=2) # High sensationalism score (2)
print(f"Claim: {claim}")
print(f"Source: {source}")
print(f"Believability: {believability}")
```

```
5.import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score
# Load the dataset
data = pd.read csv("iris.csv") # Replace with your dataset path
# Preprocess data (handle missing values, encode categorical features)
# For simplicity, let's assume no missing values and no categorical features to encode
# You may need to perform preprocessing steps here
# Separate features (X) and target variable (y)
X = data.drop("target", axis=1)
y = data["target"]
# Split data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Create and train the decision tree model
model = DecisionTreeClassifier()
model.fit(X_train, y_train)
# Make predictions on the testing set
y_pred = model.predict(X_test)
# Evaluate the model's performance
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy:", accuracy)
# Placeholder for iterative refinement process
# Analyze model's performance, gather user feedback, and refine rules
# Repeat iteration until desired performance is achieved
# You may want to visualize the decision tree and analyze its structure
# Retrain the model with refined rules (if applicable)
# For simplicity, let's skip this step in the placeholder
# Evaluate the performance of the refined model (if applicable)
# For simplicity, let's skip this step in the placeholder
6.import pandas as pd
from sklearn.preprocessing import StandardScaler
from sklearn.cluster import KMeans
import matplotlib.pyplot as plt
# Load and preprocess data
data = pd.read_csv('your_data.csv') # Replace 'your_data.csv' with your actual file
scaler = StandardScaler()
scaled data = scaler.fit transform(data)
# Choose number of clusters
n clusters = 4 # Adjust as needed
```

Create and train KMeans model

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kmeans = KMeans(n_clusters=n_clusters, random_state=42)
kmeans.fit(scaled_data)
# Assign cluster labels to data points
data['cluster'] = kmeans.labels_
# Analyze clusters: Print cluster centers
cluster centers = kmeans.cluster centers
print("Cluster centers:", cluster_centers)
# Visualize clusters
plt.scatter(scaled_data[:, 0], scaled_data[:, 1], c=data['cluster'], cmap='viridis')
plt.title('Yeast Gene Expression Clusters')
plt.xlabel('Gene 1')
plt.ylabel('Gene 2')
plt.show()
# Further analysis: Compute summary statistics of cluster means
clustered_data = data.groupby('cluster')
print(clustered_data.mean().describe())
7.from owlready2 import *
# Define the ontology
plant_phenotype_ontology = Ontology("http://plantphenotype.org/ontology.owl")
# Define classes
class Leaf(Thing):
  pass
class Shape(Thing):
  pass
class Size(DataRange):
# Define relationships (object properties & data properties)
has_shape = ObjectProperty(domain=Leaf, range=Shape)
has_size = DataProperty(domain=Leaf, range=Size)
# Define data types for attributes
float_datatype = DatatypeFactory.getBuiltInDatatype(xmlschema="float")
# Example instance
oak_leaf = Leaf(name="Oak Leaf")
oak_leaf.has_shape = "elliptical"
oak_leaf.has_size = 10.5 # using float data type
# Save the ontology
plant_phenotype_ontology.save(file="plant_phenotype.owl", format="owl")
print("Ontology created and saved successfully!")
```