Interpolation search

Introduction

The background to the interpolation search is described in **MOD007357 Coursework Brief 2021-22** (**Task a**), as search algorithm for sorted arrays to find the position of a search value. The task is to compare search performance for three strategies:

- 1. Pure interpolation search
- 2. Mixed interpolation search and binary search
- 3. Pure binary search

Implementation summary

The seearch algorithms have been implemented in the *python* programming language, and can be applied to an input array by initialising an ArraySearcher class with the array, and calling the class' search() method - providing a search strategy input.

ZIP archive

The code output of running the main() method can be found in the associated ZIP archive (interpolation-search.zip). Contents:

interpolation-search.zip

- Interpolation search.html
- Interpolation search.ipynb
- interpolation search.py
- README.md
- environment.yml

results/

- comparison_pandas.png
- testing df.csv

Dependencies

The conda environment required to run the interpolation_search.py code, and this notebook, can be recreated from the *environment.yml* file in the ZIP archive, and activated as follows:

```
conda env create -f environment.yml
conda activate interpolation-search
```

We can use a jupyter 'magic' command to generate the environment.yml file.

```
[n [ ]: !conda env export --from-history > environment.yml
```

Notebook

This notebook was created within the same python conda environment used to run the interpolation_search.py .

To generate the PDF version, the notebook was exported to HTML format and then printed to PDF format from Google Chrome.

Imports

```
In []: from matplotlib import pyplot as plt
import pandas as pd
import seaborn as sns

from interpolation_search import ArraySearcher
from interpolation_search import generate_random_array, run_tests

pd.set_option('display.max_rows', 10) # Keep table displays short
sns.set_context('talk') # Make seaborn plots have good font sizes, line widths, etc
```

Walkthrough examples

An example of each of our 3 searching algorithms can be seen below for a single input array.

Input parameters

```
In [ ]: MIN_VAL = 0

MAX_VAL = 1000

CARDINALITY = 50
```

Generate the array

```
In [ ]: array = generate_random_array(MIN_VAL, MAX_VAL, CARDINALITY)
    print(sorted(array))
[52, 63, 64, 93, 122, 140, 163, 184, 185, 197, 207, 252, 263, 299, 321, 360, 378, 378, 405, 4
```

17, 422, 444, 462, 476, 513, 519, 524, 537, 539, 546, 561, 568, 595, 615, 643, 668, 690, 692,

Make an ArraySorter

The ArraySearcher class will ensure any array passed to it is sorted before applying a search algorithm. Once initialised, it can also choose its own *query val*.

709, 715, 723, 750, 760, 824, 870, 881, 914, 918, 928, 941]

```
In [ ]: searcher = ArraySearcher(array)
   query_val = searcher.get_random_array_item()
   print(query_val)
690
```

Apply each search algorithm in turn

```
In [ ]: results = searcher.compare_methods(query_val, verbose=True)
```

```
Iteration: 0 | Starting info | Query value: 690 | Bottom index: 0 | Top index: 49 | Bottom value: 52 | Top value: 941

The query value (690) was found at index 36 (of the sorted array) after 2 iteration(s) With interpolation method it took 2 search step(s) to find the query value.

Iteration: 0 | Starting info | Query value: 690 | Bottom index: 0 | Top index: 49 | Bottom value: 52 | Top value: 941

The query value (690) was found at index 36 (of the sorted array) after 2 iteration(s) With mixed method it took 2 search step(s) to find the query value.

Iteration: 0 | Starting info | Query value: 690 | Bottom index: 0 | Top index: 49 | Bottom value: 52 | Top value: 941

The query value (690) was found at index 35 (of the sorted array) after 5 iteration(s) With binary method it took 5 search step(s) to find the query value.
```

Running the compare_methods() function returns a dictionary that summarises number of iterations used by each method to find the *query_val*.

```
In [ ]: results
Out[ ]: {'interpolation': 2, 'mixed': 2, 'binary': 5}
```

Testing increasing cardinalities

It is interesting to see how the different methods compare as we increase the length (cardinality) of the input array. To do this, we can make use of the run_tests() method from interpolation search.py.

```
In [ ]:
        def run_tests(start_cardinality, growth_mode, growth_factor, growth_steps, repeats,
                       min_array_val, max_val_factor):
            """Runs repeated tests on each provided cardinality, generating a random array
            each time and comparing each splitting method on that array.
            Args:
                start_cardinality (int): The cardinality to start with.
                 growth_mode (str): Whether to grow the cardinality arithmetically
                 ('arithmetic'), or geometrically ('geometric').
                growth_factor (int): The amount to grow the cardinality by on each step.
                 growth_steps (int): The number of times to grow the cardinality.
                repeats (int): Number of repeats.
                min array val (int): Minimum possible value in the array.
                 max_val_factor (int): Scaling factor to set the maximum possible
                                       value in the array by multiplying by the cardinality.
            Returns:
                list[dicts]: A list of dictionaries of length 'repeats'.
                              Keys are the method names, plus 'cardinality'.
                              Values are the method absolute subset differences,
                              plus cardinality
            0.00
            testing results = []
            for _ in range(repeats):
                 cardinality = start_cardinality
                for step in range(growth_steps):
                     if growth_mode == 'arithmetic':
                         cardinality = cardinality + (step * growth_factor)
                     elif growth mode == 'geometric':
                         cardinality = cardinality * (step + 1) * growth_factor
                     array = generate_random_array(min_val=min_array_val,
                                                   max_val=max_val_factor * cardinality,
                                                   cardinality=cardinality)
                     searcher = ArraySearcher(array)
                     query_val = searcher.get_random_array_item()
```

```
results = searcher.compare_methods(query_val)
    results['cardinality'] = cardinality
    testing_results.append(results)
return testing_results
```

This convenience function allows us to start with a cardinality value (*start_cardinality*), and grow this by *growth_factor* for *growth_steps* steps. Depending on *growth_mode*, the steps are either *arithmetic* or *geometric*.

The geometric steps give a real performance problem(!), so we will limit parameters accordingly.

Input parameters

```
In [ ]: MIN_ARRAY_VAL = 1
    START_CARDINALITY = 10
    MAX_VAL_FACTOR = 10
    GROWTH_STEPS = 6
    GROWTH_FACTOR = 2
    REPEATS = 100
```

Run the tests

```
In [ ]: testing_df = pd.DataFrame(testing_results)
   testing_df
```

Out[]:		interpolation	mixed	binary	cardinality
	0	2	3	3	20
	1	2	2	3	80
	2	3	4	8	480
	3	3	4	11	3840
	4	4	5	12	38400
	•••				
	595	3	3	4	80
	596	5	8	8	480
	597	3	4	11	3840
	598	3	4	12	38400
	599	4	7	18	460800

600 rows × 4 columns

Plotting with Seaborn

The above shape of the data is convenient for readability, but not so good for certain plotting approaches. The seaborn package makes plotting data (including data from a pandas.DataFrame) easy, but it is best to provide the data in a "tall, skinny" format.

Convert raw results into tall-skinny DataFrame

The raw data can be unpivotted using pandas.melt(). This gives us the tall, skinny version of the output data where each method appears not as a column, but as a value in a single 'method' column.

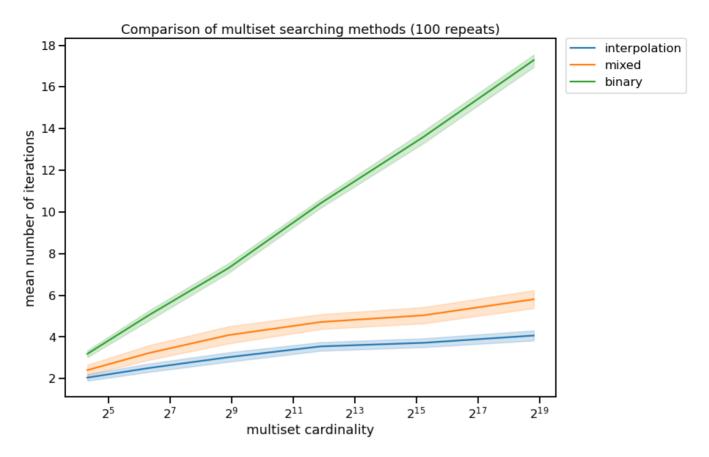
ut[]:		cardinality	method	iterations
	0	20	interpolation	2
	1	80	interpolation	2
	2	480	interpolation	3
	3	3840	interpolation	3
	4	38400	interpolation	4
	•••			
	1795	80	binary	4
	1796	480	binary	8
	1797	3840	binary	11
	1798	38400	binary	12
	1799	460800	binary	18

1800 rows × 3 columns

Plot

Seaborn's lineplot() method is an appropriate choice. This method will plot the mean of the data, and can automatically show variation from the mean in a number of ways. As we are probably most interested in how confident we are in the predicted mean values, 95% confidence intervals have been chosen (rather than standard deviations).

```
In []: fig, ax = plt.subplots(figsize=(12, 9), facecolor='white')
    ax = sns.lineplot(data=tall_df, x='cardinality', y='iterations', hue='method', ci=95)
    ax.set_xlabel('multiset cardinality')
    ax.set_xscale('log', base=2)
    ax.set_ylabel('mean number of iterations')
    # Put Legend outside plot
    ax.legend(bbox_to_anchor=(1.02, 1), loc='upper left', borderaxespad=0)
    # Underscore assignment to supress Text object output
    _ = ax.set_title(f'Comparison of multiset searching methods ({REPEATS} repeats)')
```



Both of the above steps can be conveniently run by calling the seaborn_plot() method from partition_problem.py. The inputs to this function are simply the raw result dictionary list, and the list of cardinality values used to generate the results:

fig = seaborn_plot(results=testing_results, repeats=REPEATS)