Sum of Two Terms

Mac Radigan

Are The Individual Terms Of A Two-Term Sum Present In An Array?

Overview

Given an array of integers, \mathbb{X} , determine whether or not there exist two elements, m and n, in the array (at different positions) whose sum is equal to some target value, Σ .

Algorithm #1

Background

Algorithm 1 makes use of the fact that:

$$n = (n - k) + k$$

holds true for any integers n and k.

Therefor, given knowledge of the elements present in the sequence \mathbb{X} between 0 and Σ , we can apply the above formula as a test for existence.

This algorithm makes an assumption about the maximal sum that will be encountered, M, and uses this size in allocating a histogram.

Implementation

Initially size a histogram of unit bin size based on the expected maximum sum to be supported (say, upper bound M).

Build a histogram of unit bin size from the input sequence.

$$h_k = |\{k : k \in \mathbb{X}\}|$$

Scan the histogram up to half the number of bins, applying the formula:

$$n = (n - k) + k$$

If the above equation holds for any element encountered, then two terms have been found that add to the given sum.

Note that in the implementation we are using the C++ behavior that of integral type having a truth value of T if and only if their register value is non-zero.

```
Algorithm 1 Has Two Sum Terms given set of terms X, goal sum \Sigma
```

```
for each x \in \mathbb{X} do h_x \leftarrow h_x + 1 \alpha build a histogram end for each for k \leftarrow 0 \cdots |\mathbb{X}| do if \Sigma = h_{\Sigma - k} + h_k then return \top \alpha check sum of terms holds end if if 2 \mid \Sigma \wedge h_{\lceil \frac{k}{2} \rceil} \geq 2 then return \top \alpha check two terms in histogram center end if end for return \bot
```

Performance

For a sequence xs, having N elements, we have:

Measure	Performance
average time complexity worst case time complexity	$\frac{\mathrm{O(N)}}{\mathrm{N}+1/2\;\mathrm{M}}$
constant space complexity	M

Note that time complexity assumptions for the average case are not strictly valid without knowledge of the underlying statistical distribution of the input data.

Note that for large M, the space complexity of this algorithm may be substantial. Algorithm 2 provides better performance when M is large.

```
//
   //
         returns true if there exists elements m and n in sequence xs such that
   //
                      sum = m + n, where xs and sum are given
   //
   //
         inputs:
   //
          xs : vector<T> - a sequence of terms to consider
11
   //
                            - the specified target sum for testing terms
          sum : T
   //
  //
         template paramters:
14
  //
15
   //
          T : class
                            - the data type of the terms and target sum
   //
   //
          M: size_t
                            - an upper bound on the expected sum
   //
19
   //
        returns:
20
  //
21
  //
           has_terms : bool - true if the input sequence contains two elements
22
  //
                                     equal to a given sum
23
                              false otherwise
24
25
26
   template<class T, std::size_t M>
   class SequenceCheck
28
29
    public:
30
     inline bool has_two_sum_terms(const std::vector<T> &xs, const T sum)
31
32
        // build a histogram
33
       hist_.fill(0);
34
       for(auto x : xs) if(x<=sum) hist_[x]++;</pre>
35
       // each element up to half of the histogram
36
       for(auto k=0; k<std::floor(sum/2.0); ++k)</pre>
37
         // test if n = (n-k) + k
39
         if( hist_[sum-k] && hist_[k] ) return true;
40
        } //  k = 0...sum/2 
41
        // special case at ceil(N/2) when N odd, must have at least two terms
       if( !(sum%2) && hist_[std::ceil(sum/2.0)]>1 ) return true;
43
       return false; // otherwise no such two terms
     } // has_two_terms
45
    private:
     std::array<T,M> hist_{};
   }; // SequenceCheck
```

Algorithm #2

Note that Algorithm 2 currently does not enforce the selection of two terms in the sequence to be from two distinct different positions.

Background

Algorithm 2 makes use of the algebraic group property that every number has an inverse, and thus we may rewrite:

$$\Sigma = m + n \leftrightarrow n = \Sigma - m$$

Thus for each element m encountered in \mathbb{X} , we know uniquely of a corresponding n in \mathbb{X} that we seek.

Therefor, we may scan xs once to identify its compliment with respect to Σ .

Now, with a set of compliments, say $\overline{\mathbb{X}}$, we may scan \mathbb{X} again to determine if any element exists in $\overline{\mathbb{X}}$.

If we find that an element in X is found in the set of compliments, then we know the sum can be produced from two terms that exist in the sequence.

Since it is also required that the terms in the sum are at distictly different positions in the sequence, use an unordered map for the set of compliments, and use the position to track the position of the original term. When checking the compliment, verify that the position is also distinct from the original term.

Implementation

Initially an empty, unordered map (uses a hash map implementation). Call this the compliment map $\overline{\mathbb{X}}$.

Scan the input sequence, \mathbb{X} , for each x in \mathbb{X} . For each x, compute the compliment of the sum, Σ , and x, say: $\bar{c} = \Sigma - x$, and insert the compliment \bar{c} and the ordinal position of x (say k) into $\overline{\mathbb{X}}$.

Scan the input sequence, \mathbb{X} , for each x in \mathbb{X} again, checking the compliment map $\overline{\mathbb{X}}$ for x. If x exists in $\overline{\mathbb{X}}$, and the position is distinct from the map's value of k, then we have found two terms that produce the sum.

If the end of the sequence is reached without finding a matching x in the compliment map, $\overline{\mathbb{X}}$, then there are no two terms in $\operatorname{mathbb} X$ that will produce the sum.

Performance

For a sequence X, having N elements, we have:

Algorithm 2 Has Two Sum Terms

```
given set of terms \mathbb{X}, goal sum \Sigma for k \leftarrow 0 \cdots |\mathbb{X}| do \overline{x} \leftarrow \Sigma - \mathbb{X}_k \mathbb{F}_{\overline{x}} \leftarrow k \Theta map of compliments to position end for for k \leftarrow 0 \cdots |\mathbb{X}| do if \mathbb{X}_k \in \overline{\mathbb{X}} \wedge \mathbb{F}_k \neq k then return \top \Theta check are compliments present and distinct end if end for return \bot
```

Measure	Performance
average time complexity	O(N)
best case time complexity	N + 1
worst case time complexity	2 * N
average case space complexity	O(N)
best case space complexity	1
worst case space complexity	N

Note that the space complexity is dependent only on the number of unique elements in the input sequence (X).

Note that for small M, algorithm 1 can have better worst case time complexity as well as substantially better space complexity. These assumptions can be further bounded and refined given statistical knowledge of the input data.

```
// -----
  // has_two_sum_terms (Algorithm 2)
  //
      returns true if there exists elements m and n in sequence xs such that
                 sum = m + n, where xs and sum are given
      inputs:
        xs : vector<T> - a sequence of terms to consider
        sum : T
                     - the specified target sum for testing terms
      template paramters:
                     - the data type of the terms and target sum
  //
        T : class
16
      returns:
                                                      5
        has_terms : bool - true if the input sequence contains two elements
                            equal to a given sum
                       false otherwise
22
23
  bool has_two_sum_terms(const std::vector<T> &xs, const T sum)
```

```
30
       const T x = xs[k];
31
       const auto diff = sum - x; // caution: T must be a signed datatype
32
       compliments[diff] = k;
33
     } // foreach index k of x in xs
     // scan the input sequence again to identify if any compliments are present
35
     for(auto k=0; k<xs.size(); ++k)</pre>
37
       const T x = xs[k];
       const auto it = compliments.find(x);
39
       // check that the compliment is at a distict position from the original term
40
       if( (it!= compliments.end()) && (k != it->second) ) return true;
41
     } // foreach x in xs
     return false; // otherwise no such two terms
   } // has_two_terms
```

Source Code

```
// sum-two-terms.cc
  // Mac Radigan
    #include <array>
    #include <assert.h>
    #include <cmath>
    #include <cstdlib>
    #include <iomanip>
    #include <iostream>
10
    #include <map>
11
    #include <sys/types.h>
12
13
    #include <unordered_map>
    #include <vector>
14
15
16
    // -----
17
    // has_two_sum_terms (Algorithm 1)
18
    // -----
19
    //
20
    //
        returns true if there exists elements m and n in sequence xs such that
21
    //
                   sum = m + n, where xs and sum are given
22
    //
23
    //
        inputs:
25
```

```
xs : vector<T> - a sequence of terms to consider
26
     //
27
     //
                              - the specified target sum for testing terms
             sum : T
     //
29
     //
           template paramters:
     //
31
     //
             T : class
                              - the data type of the terms and target sum
     //
33
     //
                              - an upper bound on the expected sum
            M: size_t
     //
35
     //
          returns:
36
     //
37
     //
            has_terms : bool - true if the input sequence contains two elements
                                       equal to a given sum
     //
39
     //
                                false otherwise
40
     //
41
     //
42
     //
43
     //
44
     // Background:
45
     //
46
     //
          Algorithm 1 makes use of the fact that:
47
     //
            n = (n-k) + k
48
     //
          holds true for any integers n and k.
     //
50
     //
           Therefor, given knowledge of the elements present in the sequence xs
51
     //
            between 0 and sum, we can apply the above formula as a test for
52
     //
             existence.
53
     //
54
     //
          This algorithm makes an assumption about the maximal sum that will
             be encountered, M, and uses this size in allocating a histogram.
     //
56
     //
     //
58
     // Implementation:
59
     //
60
     //
           Initially size a histogram of unit bin size based on the expected
61
     //
             maximum sum to be supported (say, upper bound M).
     //
63
     //
          Build a histogram of unit bin size from the input sequence.
     //
65
          Scan the histogram up to half the number of bins, applying the formula:
     //
     //
             n = (n-k) + k
67
     //
     //
           If the above equation holds for any element encountered, then two terms
69
             have been found that add to the given sum.
     //
```

```
//
71
      //
            There is one additional case to consider, that is, when considering the
72
      11
              center element of the histogram when the sum is odd. In this case,
      //
              since is it required that the terms in the sum are at different
74
      //
             positions, we must check that there are two terms in the sequence,
75
      //
              in other words, that the histogram count at this position is greater
76
      //
              than one.
      //
78
      //
           Note that in the implementation we are using the C++ behavior that of
      //
              integral type having a truth value of T if and only if their register
80
      //
              value is non-zero.
81
      //
82
      //
      // Performance:
84
      //
85
      //
           For a sequence xs, having N elements, we have:
86
      //
87
                                               O(N)
      //
              average time complexity:
88
      //
              worst case time complexity:
                                               N + 1/2 M
89
      //
      //
              constant space complexity:
91
      //
92
      //
93
      //
           Note that time complexity assumptions for the average case are not strictly
      11
             valid without knowledge of the underlying statistical distribution of the
95
      //
              input data.
      //
97
      //
           Note that for large M, the space complexity of this algorithm may be
aa
      //
              substantial. Algorithm 2 provides better performance when M is large.
      //
101
      //
102
      namespace demo::algo1 {
103
        template < class T, std::size t M>
104
        class SequenceCheck
105
106
         public:
          inline bool has_two_sum_terms(const std::vector<T> &xs, const T sum)
108
          {
             // build a histogram
110
            hist_.fill(0);
111
            for(auto x : xs) if(x<=sum) hist_[x]++;</pre>
112
             // each element up to half of the histogram
113
            for(auto k=0; k<std::floor(sum/2.0); ++k)</pre>
114
             {
115
```

```
// test if n = (n-k) + k
116
             if( hist_[sum-k] && hist_[k] ) return true;
117
             // k = 0...sum/2 
           // special case at ceil(N/2) when N odd, must have at least two terms
119
           if( !(sum%2) && hist_[std::ceil(sum/2.0)]>1 ) return true;
           return false; // otherwise no such two terms
121
         } // has_two_terms
        private:
123
         std::array<T,M> hist_{};
124
       }; // SequenceCheck
125
     } // demo::algo1
126
127
      129
     // has_two_sum_terms (Algorithm 2)
130
     131
      //
132
     //
          returns true if there exists elements m and n in sequence xs such that
133
     //
                      sum = m + n, where xs and sum are given
134
     //
135
      //
          inputs:
136
      //
137
     //
            xs : vector<T> - a sequence of terms to consider
138
      //
139
      //
            sum : T
                            - the specified target sum for testing terms
140
     //
     //
          template paramters:
142
      //
      //
            T : class
                            - the data type of the terms and target sum
144
     //
          returns:
     //
146
      //
     //
            has_terms : bool - true if the input sequence contains two elements
148
                                   equal to a given sum
     //
149
     //
                              false otherwise
150
151
152
     //
153
     //
154
     // Background:
155
     //
     //
          Algorithm 2 makes use of the algebraic group property that every number
157
      //
            has an inverse, and thus we may rewrite:
     //
159
      //
160
            s = m + n as n = s - m
```

```
//
161
      //
            Thus for each element m encountered in xs, we know uniquely of a
162
      //
              corresponding n in xs that we seek.
      //
164
      //
           Therefor, we may scan as once to identify its compliment with respect
165
      //
166
      //
      //
           Now, with a set of compliments, say xs', we may scan xs again to
168
      //
              determine if any element exists in xs'.
169
      //
170
      //
           If we find that an element in xs is found in the set of compliments,
171
      //
              then we know the sum can be produced from two terms that exist in
172
      //
              the sequence.
173
      //
174
      //
           Since it is also required that the terms in the sum are at distictly
      //
              different positions in the sequence, use an unordered map for the
176
      //
             set of compliments, and use the position to track the position of
177
              the original term. When checking the compliment, verify that
      //
178
      //
              the position is also distinct from the original term.
179
      //
180
      // Implementation:
181
      //
      //
           Initially an empty, unordered map (uses a hash map implementation).
183
      //
             Call this the compliment map cs.
      //
185
      //
           Scan the input sequence, xs, for each x in xs. For each x, compute
      //
              the compliment of the sum, s, and x, say: c = s - x, and insert
187
      //
              the compliment c and the ordinal position of x (say k) into cs.
      //
180
      //
           Scan the input sequence, xs, for each x in xs again, checking the
      //
              compliment map cs for x. If x exists in cs, and the position is
191
      //
              distinct from the map's value of k, then we have found two terms
192
      //
              that produce the sum.
193
      //
194
      //
           If the end of the sequence is reached without finding a matching x in
195
      //
              the compliment map, cs, then there are no two terms in xs that will
196
      //
             produce the sum.
197
      //
198
      //
199
      // Performance:
200
      //
      11
           For a sequence xs, having N elements, we have:
202
      //
              average time complexity:
      //
                                                O(N)
204
              best case time complexity:
                                                N + 1
```

```
2 * N
              worst case time complexity:
206
207
      //
              average case space complexity:
                                                  O(N)
      //
              best case space complexity:
                                                  1
209
      //
              worst case space complexity:
                                                  N
210
       //
211
      //
            Note that the space complexity is dependent only on the number of
      //
213
      //
              unique elements in the input sequence (xs).
214
      //
215
      //
216
      11
            Note that for small M, algorithm 1 can have better worst case time complexity
217
      //
              as well as substantially better space complexity. These assumptions can
      //
              be further bounded and refined given statistical knowledge of the input data.
219
      //
220
      //
221
      namespace demo::algo2 {
222
         template<class T>
223
         bool has_two_sum_terms(const std::vector<T> &xs, const T sum)
224
225
           // hash map of compliments: CS := \{ c : sum-x=c \text{ forall } x \text{ in } xs \}
226
           std::unordered_map<T,std::size_t> compliments;
           for(auto k=0; k<xs.size(); ++k)</pre>
228
             const T x = xs[k];
230
             const auto diff = sum - x; // caution: T must be a signed datatype
231
             compliments[diff] = k;
232
           } // foreach index k of x in xs
233
           // scan the input sequence again to identify if any compliments are present
234
235
           for(auto k=0; k<xs.size(); ++k)</pre>
236
             const T x = xs[k];
237
             const auto it = compliments.find(x);
238
             // check that the compliment is at a distinct position from the original term
239
             if( (it!= compliments.end()) && (k != it->second) ) return true;
240
           } // foreach x in xs
241
           return false; // otherwise no such two terms
         } // has_two_terms
243
      } // demo::algo2
244
245
246
      //
247
      // main test driver
249
250
      int main(int argc, char *argv[])
```

```
251
252
         // default element type (domain)
        typedef int64_t element_t;
254
         /*
256
257
            Basic Fobonacci tests
258
259
          */
260
261
        // An array containing the first 20 terms of the Fibonacci sequence
262
         //
263
        11
              F[n] := F[n-1] + F[n-2], with F[1] := 1 and F[2] := 2 for n = 0...20
264
         //
265
        //
              Note that this subsequence (xs) contains the following pairs:
266
         //
267
                (3, 5) in xs, and 8 = 3 + 5
         //
268
         //
269
         //
                (13,21) in xs, and 34 = 13 + 21
         //
271
         //
         //
              And also that this subsequence (xs) does not contain pairs satisfying the followin
273
         //
         //
                19 = m + n for any m and n in xs
275
         //
276
        //
                41 = m + n for any m and n in xs
277
         //
278
         //
279
         std::vector<element_t> xs = {
280
              1,
                    2,
                           3,
                                 5,
                                         8,
281
                   21,
                          34,
                                55,
                                        89,
             13,
282
                        377,
            144,
                 233,
                              610,
                                       987,
283
           1597, 2584, 4181, 6765, 10946
284
        }; // fibonnaci sequence xs
286
         // unit test for both algorithms
        auto my_assert = [](const std::vector<element_t> &xs, element_t x, bool expect) -> bool
288
           // assumptions about the maximum expected target sum
           constexpr std::size_t M = 1024;
290
           // test algorithm 1
291
           demo::algo1::SequenceCheck<element_t, M> check;
292
           auto result_1 = check.has_two_sum_terms(xs, x);
293
           assert(result_1 == expect);
294
           // test algorithm 2
295
```

```
auto result_2 = demo::algo2::has_two_sum_terms<element_t>(xs, x);
296
           assert(result_2 == expect);
297
           std::cout << "test case for sum "</pre>
                      << std::setw(3) << x
299
                      << " passed"
                      << std::endl << std::flush;
301
         }; // my_assert
302
303
         // list of test cases with expected results
304
         std::map<element_t, bool> test_cases = {
305
           { 8, true},
306
           {34, true},
307
           {19, false},
308
           {41, false}
309
         }; // test_cases
310
311
         // run all tests
312
         for(auto &test : test_cases) my_assert(xs, test.first, test.second);
313
314
         /*
315
316
             Some additional stress tests to handle special cases
318
          */
319
320
         std::vector<element_t> xs_2 = {
321
             1, 1,
322
             1, 1,
323
             4,
                4,
324
             9,
325
            12
326
         }; // stress sequence x2_s
327
328
         // list of test cases with expected results
329
         std::map<element_t, bool> test_cases_2 = {
330
           { 2, true},
331
           { 8, true},
           { 9, false},
333
           {12, false}
         }; // test_cases_2
335
336
         // run all tests
337
         for(auto &test : test_cases_2) my_assert(xs_2, test.first, test.second);
338
339
         return EXIT_SUCCESS;
340
```

```
341 } // main
342
343 // *EOF
```

Test Data Generation

```
#!/usr/bin/env stack
-- Fibonacci.hs
-- Mac Radigan

fib :: Integer -> Integer
fib 0 = 1
fib 1 = 1
fib 1 = 1
fib n = fib (n-1) + fib (n-2)

main :: IO ()
main = print $ map fib [1..20]

-- *EOF*
```

Unit Test Results

```
test case for sum 8 passed
test case for sum 19 passed
test case for sum 34 passed
test case for sum 41 passed
test case for sum 2 passed
test case for sum 8 passed
test case for sum 9 passed
test case for sum 12 passed
```