To nest or not to nest, that is the question. An argument that NCList is unnecessary. BH 2019.08.09 rev.

**Just for discussion**; based on *Nested Containment List (NCList): a new algorithm for accelerating interval query of genome alignment and interval databases*, Alexander V. Alekseyenko and Christopher J. Lee, Vol. 23 no. 11 **2007**, pages 1386–1393. doi:10.1093/bioinformatics/btl647

These authors present a nested approach to finding overlapping intervals within a potentially very large set of intervals. They argue that in a situation such as given in their Fig. 2 (Figure 1), it is better to nest interval y into x than to leave it in the initially searched list.

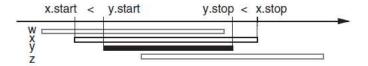


Fig. 2. Any contained interval breaks sortedness. If intervals (boxes) are sorted on their start coordinate, then any interval y (filled box) that breaks sortedness of the stop coordinate is properly contained in another interval x.

Figure 1. The original issue leading to the development of NCList.

The authors state the problem in this way (bold mine):

Interval query can be slow because the overlapping intervals for any given query may not be contiguous in standard indexing. Therefore, the database query cannot stop at the first non-overlapping interval, but must scan the rest of the database.

### And their solution:

We can easily solve this problem by realizing that it is caused solely by the intervals that are contained within other intervals, i.e. x. start < y. start y.stop < x.stop. (see Fig. 2). If a sorted list of intervals has both start and stop coordinates in ascending order, then the overlapping intervals for any query are guaranteed to be contiguous in the list.

The solution is clever. Create a list that is *monotonic* in both start and end for all intervals by packaging any nonconforming intervals (*y* in this case) into one of those conforming intervals (i.e., *x*). NCList's findOverlap(from,to) method involves a series of binary searches that are of sets potentially smaller than *N*, each followed by a linear contiguous scan. For example, consider using NCList, for all overlaps of interval *I*, as shown in (my) Figure 2. Intervals *w*, *x*, and *z* are monotonically increasing in both start and end positions, but *y* is not. We package *y* into *x*. A binary search for "the first end point after the start of interval *I*" finds *w*. A linear forward scan finds *x*, and *z*. A second binary search and linear scan through the subintervals of *x* finds *y*.

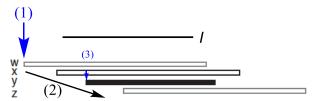


Figure 2: NCList looking for overlaps with *I* finds *w* first using a binary search for the closest end point after the start of *I* (1), then scans forward for intervals that have a starting point before the end of *I* (2), finding *x* and *z*. Interval *y* is found within *x* using a second binary search (3), followed again by a second forward search (not shown, because in this case *y* is the only subinterval of *x*).

#### Jalview's NCList implementation - IntervalStore

The implementation of NCList in Jalview, IntervalStore<SequenceFeature> is identical to the original specification, with the added efficiency that during initialization all intervals that have no subintervals are set aside in a separate list. While this adds complications when adding or removing intervals from the set, it is more efficient in terms of storage (just a simple ArrayList<SequenceFeature>). The Jalview implementation is further optimized by separating different *types* of features (plaf, variant, etc.) into separate FeatureStore objects, each with their own IntervalStore. This substantially reduces the number of levels of nesting as well as the number of intervals in any single binary search.

#### But do we have to nest?

There is another, considerably simpler, solution involving a linked list with similarly scalable performance. The problem addressed by NCList is that once the first binary search is over, we still don't know which intervals that start ahead of our specified interval overlap with it. Consider the situation in Figure 3. A set of eight intervals, a-h, are not ordered by start but not monotonic in end. NCList creates a set of subsets that are internally monotonic in both start and end: Set a contains b, g, and h; Set b contains c, d, and e; and Set e contains f. The key point here is that we are removing c, d, e, and f from the initial binary search.

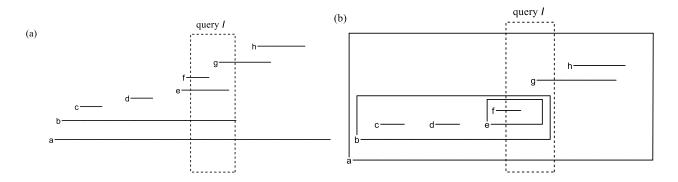


Figure 3. (a) Eight intervals a-h, in order of increasing start position is not monotonic in end position. The indicated query must return the set {a, b, e, f, g}. (b) NCList nests intervals that are not monotonically increasing in both start and end points, creating subset {b, g, h} within a, subset {c, d, e} within b, and subset {f} within e. Within each subset the intervals are monotonic in both start and end point.

NCList processing finds the "first interval having an end point not before the query interval." In this case, it finds a. Then, within a's subsets it finds b, within b it finds e, and within e it finds f. Interval g is found by scanning the subset {b, g, h} in sequential order until the start of an interval (h in this case) is past the end of the query interval. This works, but might it be overly complicated?

Now consider the simple alternative shown in Figure 4, where we allow for a linked list of pointers from an interval to the *nearest prior interval that contains its starting point*.

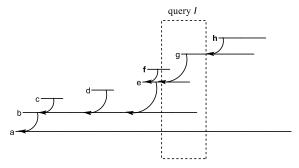


Figure 4. An alternative to NCList that utilizes a reference for each interval pointing to the nearest prior interval that contains its starting point. Note that intervals f and h flank this interval but are not included in it.

A dual binary search for the two nearest starting points flanking this query gives us f and h. The set of all intervals starting between these ( $\{g\}$  in this case) is added to the results. Interval f along with all intervals that are linked from it ( $f \rightarrow e \rightarrow b \rightarrow a$ ), are added only if their end is not before the start of the query interval. Note that intervals c and d are never checked, since they are not in the linked list starting from f.

The implementation at <a href="https://github.com/BobHanson/IntervalStoreJ">https://github.com/BobHanson/IntervalStoreJ</a> uses a simple int[] array to manage the linked list, where the elements of the array are relative index offsets rather than actual object references. Thus, for the set {a b c d e f g h} the offsets would be [\*, 1, 1, 2, 3, 1, 2, 1], where \* is a reserved number (Integer.MIN\_VALUE, as implemented) meaning "not contained", which stops all link processing when it is found.

As described, the algorithm is not scalable for sets where there is substantial overlap. The problem is that this sort of set produces long strings of pointers. An improvement involves indicating an offset with a negative number when the pointer is to an interval that has a higher end point than any that comes before it (Figure 5). If that interval is checked and found to have an end point prior to the start of the query interval, it is guaranteed that no further checking along the chain will lead to a result. We can stop scanning.

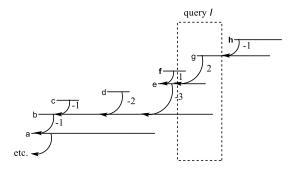


Figure 5: Offsets indicated with negative numbers to indicate a link to the interval with the highest end point of any previous intervals. The process (f -> e -> b -> a -> etc.) can be terminated after checking interval a, since its offset from b is -1, meaning that we know that no interval before Interval a extends beyond Interval a (into query I).

## **Advantages and Disadvantages**

The advantage of NCList is that it pre-partitions the binary search of N objects into a set of n binary searches of  $m_i \le N$  objects, where SUM(i){ $m_i$ } = N. Depending upon the extent of nesting, this could be significant. With minimally nested sets, however, it is unlikely that this advantage would be noticeable.

The primary advantages of the linked list approach include:

- (1) It processes queries very efficiently, with a single binary search, followed by a single link-based check.
- (2) It requires minimal initialization, with very little allocated memory (just simple arrays Intervall[N] and int[N]). The simple linked list avoids the necessity for all the nesting structure that comes with NCNodes and NCList, as well as all initialization that goes with those objects.
- (3) It allows for "lazy" initialization. That is, we can do all the loading of the list, including minor addition/removal with the option to not sort the actual list or build the links until it is absolutely necessary (the first findOverlap() call, generally). Rebuilding after addition or removal is simply a recalculation of the offsets array.
- (4) The return list is in the same order (albeit reversed, for performance reasons) as the original sorted list. In contrast, IntervalStoreJ's implementation of NCList uses of separate nested and unnested lists, which are processed sequentially. It thus returns a list that might or might not be ordered. In some situations, this could be an advantage.

## Timing results: query

Timing results for querying (Table 1 and Figure 6) suggest that using a linked list is from two to three times faster than NCList alone or the IntervalStoreJ/NCList implementation. In fact, compared to NCList, the linked list alternative will return 100,000 queries from a set containing 464K intervals in the time it takes NCList to return 100,000 queries from a list that contains only 2K intervals. All three method times are linear in log N for large N and governed by other factors at low N.

logN	N	IntNC	NCList	IntLink
3.33	2154	160.4	222.9	88.2
3.67	4641	165.1	232.8	90.6
4.00	10000	173.5	241.8	95.4
4.33	21544	183.9	262.6	99.4
4.67	46415	201.2	306.9	105.7
5.00	100000	244.8	365.3	121.5
5.33	215443	289.7	406.5	144.3
5.67	464158	327.3	435.1	169.6
6.00	1000000	363.7	467.8	189.1

Table 1. Timing results (in ms) for returning 100,000 queries from a "sequence" of length N \* 10 that contains N intervals of pseudorandom length 1 to 50 for IntervalStore/NCList ("IntNC"), Nclist alone, and IntervalStore/linked list ("IntLink"). The start positions of the intervals within the sequence are in the range 1 to N \* 10 - 50. Each query has a length of 1000 and a pseudorandom position within the sequence.

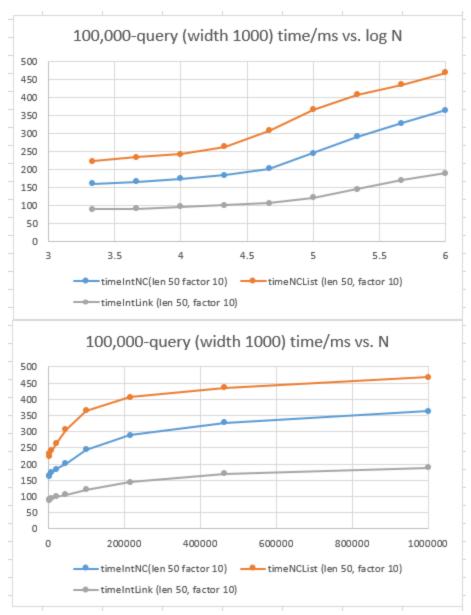


Figure 6. Query time vs. logN and N for the linked-list variation of IntervalStore (lowest set), along with the IntervalStore implementation of NCList (middle data set), and NCList without IntervalStore (highest set).

# Full run:

[RemoteTestNG] detected TestNG version 6.14.2

Java version: 1.8.0\_191

amd64 Windows 10 10.0 cores:4 14 Aug 2019 21:09:15 GMT

					time			
Test	size N	tests	time/ms	rate/(N/ms)	stderr	rate stderr		
# Query IntStoreNCList store interval size 50 store sequence factor 10 query width -1000 query count 100000								
IntStoreNCList query	2154	10	160.4	13.5	2.93	0.24		
IntStoreNCList query	4641	10	165.1	28.1	0.54	0.09		
IntStoreNCList query	10000	10	173.5	57.6	0.7	0.23		
IntStoreNCList query	21544	10	183.9	117.2	0.48	0.3		
IntStoreNCList query	46415	10	201.2	230.7	0.79	0.89		
IntStoreNCList query	100000	10	244.8	408.4	0.54	0.9		
IntStoreNCList query	215443	10	289.7	743.7	0.47	1.2		
IntStoreNCList query	464158	10	327.3	1418.4	1.16	4.95		
IntStoreNCList query	1000000	10	363.7	2749.4	0.57	4.33		
# dimensions [7 1000000]								
# Query IntStoreNCList0 store interv	al size 50 stor	e sequer	nce factor 10	query width -1	000 query cou	nt 100000		
IntStoreNCList0 query	2154	10	183.3	11.8	3.89	0.22		
IntStoreNCList0 query	4641	10	190.5	24.4	1.6	0.19		
IntStoreNCList0 query	10000	10	197.1	50.7	0.56	0.14		
IntStoreNCList0 query	21544	10	207	104.1	0.46	0.23		
IntStoreNCList0 query	46415	10	225.2	206.1	0.43	0.39		
IntStoreNCList0 query	100000	10	286.5	351.2	8.01	8.63		
IntStoreNCList0 query	215443	10	315.6	683	2.03	4.21		
IntStoreNCList0 query	464158	10	354.2	1310.4	0.59	2.19		
IntStoreNCList0 query	1000000	10	389.6	2567	0.99	6.6		
# dimensions [7 0]								
# Query IntStoreLink store interval s	ize 50 store se	quence	factor 10 que	ery width -1000	query count 1	100000		
IntStoreLink query	2154	10	88.2	24.5	2.24	0.52		
IntStoreLink query	4641	10	90.6	51.2	0.62	0.34		
IntStoreLink query	10000	10	95.4	104.8	0.41	0.45		
IntStoreLink query	21544	10	99.4	216.8	0.24	0.53		
IntStoreLink query	46415	10	105.7	439.3	0.34	1.43		
IntStoreLink query	100000	10	121.5	823.4	0.3	2		
IntStoreLink query	215443	10	144.3	1493.6	0.58	5.89		
IntStoreLink query	464158	10	169.6	2748.3	3.96	54.12		
IntStoreLink query	1000000	10	189.1	5287.4	0.29	8.23		
# dimensions [126 416364]								
# Query IntStoreLink0 store interval size 50 store sequence factor 10 query width -1000 query count 100000								
IntStoreLink0 query	2154	10	83.1	25.9	0.18	0.06		
IntStoreLinkO query	4641	10	87	53.4	0.81	0.47		
IntStoreLinkO query	10000	10	89.7	111.5	0.28	0.34		
IntStoreLinkO query	21544	10	93.6	230.2	0.31	0.76		
IntStoreLinkO query	46415	10	99.7	465.7	0.31	1.46		
IntStoreLinkO query	100000	10	120	833	0.29	2.01		
4001	_00000		0	000	0.23	2.01		

IntStoreLinkO query	215443	10	146	1476.1	0.14	1.45		
IntStoreLinkO query	464158	10	170.2	2726.5	0.42	6.71		
IntStoreLinkO query	1000000	10	193.3	5173.2	0.19	5.14		
# dimensions [126 416364]								
# Query NCList store interval size 50 store sequence factor 10 query width -1000 query count 100000								
NCList query	2154	10	222.9	9.7	0.71	0.03		
NCList query	4641	10	232.8	19.9	0.41	0.03		
NCList query	10000	10	241.8	41.3	0.51	0.09		
NCList query	21544	10	262.6	82.8	9.44	2.31		
NCList query	46415	10	306.9	151.2	0.4	0.2		
NCList query	100000	10	365.3	273.7	0.49	0.37		
NCList query	215443	10	406.5	530	0.19	0.25		
NCList query	464158	10	435.1	1066.9	0.21	0.52		
NCList query	1000000	10	467.8	2142.7	8.04	32.38		
# dimensions [7 528974]								
# Query NCList0 store interval size 50 store sequence factor 10 query width -1000 query count 100000								
NCList0 query	2154	10	230.3	9.4	2.1	0.08		
NCList0 query	4641	10	239.7	19.4	0.86	0.07		
NCList0 query	10000	10	246.3	40.6	0.27	0.04		
NCList0 query	21544	10	260.3	82.8	0.85	0.27		
NCList0 query	46415	10	310.2	149.6	0.37	0.18		
NCList0 query	100000	10	365	274	0.64	0.48		
NCList0 query	215443	10	404.9	532.1	0.33	0.44		
NCList0 query	464158	10	434.4	1068.6	0.66	1.62		
NCList0 query	1000000	10	462.4	2163.4	3.36	14.88		
# dimensions [7 ?]								

# resultcounts [0, 0, 0, 0, 0, 0, 0, 0, 0, 100, 113, 96, 112, 100, 93, 98, 112, 88]

PASSED: testLoadTimeBulk

PASSED: testLoadTimeIncrementalAllowDulicates PASSED: testLoadTimeIncrementalNoDuplicates

PASSED: testQueryTime PASSED: testRemoveTime

Default test

Tests run: 5, Failures: 0, Skips: 0

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Default suite

Total tests run: 5, Failures: 0, Skips: 0

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```
/**
 * factor to multiply first parameter of generateIntervals(sequenceWidth,
 * count, length) by to set store sequence width; higher number reduces number
 * of overlaps
private static final int QUERY_STORE_SEQUENCE_SIZE_FACTOR = 10;// 10;
 * interval size for the store; absolute(negative) or maximum(positive);
private static final int QUERY_STORE_INTERVAL_SIZE = 50;// -1 for SNPs;
 * width of query intervals; negative for absolute, positive for max value
*/
// private static final int QUERY_WINDOW = -1;// overview single-pixel overlap
private static final int QUERY_WINDOW = -1000;// -1000 standard view
 * number of queries to generate (independently of the size of the sequence
private static final int QUERY_COUNT = 100000;
  int sequenceWidth = count * QUERY_STORE_SEQUENCE_SIZE_FACTOR;
      List<Range> ranges = generateIntervals(sequenceWidth, count,
              QUERY_STORE_INTERVAL_SIZE);
      List<Range> queries = generateIntervals(sequenceWidth, QUERY_COUNT,
              QUERY_WINDOW);
/**
 * Generates a list of <code>count</code> intervals of length [1,length] in
 * the range [1, sequenceWidth]
* @param sequenceWidth
            scale of the sequence, based on the number of intervals present,
           not the number of queries
 * @param count
           the number of intervals to generate
           maximum (positive) or absolute(negative) number of intervals to
           generate
 * @return list of intervals
private synchronized List<Range> generateIntervals(int sequenceWidth,
        int count, int length)
  int maxPos = sequenceWidth - Math.abs(length);
  List<Range> ranges = new ArrayList<>();
  for (int j = 0; j < count; j++)</pre>
    int from = 1 + rand.nextInt(maxPos);
    int to = from + (length < 0 ? -length - 1 : rand.nextInt(length));</pre>
    ranges.add(new Range(from, to));
  }
  return ranges;
}
```