Algorithms and Datastructures Cache Efficiency, Divide and Conquer

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Structure

Cache Efficiency Introduction Cache Organization

Divide and Conquer Introduction

Introduction

Background:

- Up to now we always counted number of operations
- Assuming this is a good measure for the runtime of a algorithm/tool
- ▶ Today we will see examples where this is not suitable

Introduction

Example:

- ▶ We sum up all elements of a field a of size n in . . .
 - natural order:

$$sum(a) = a[1] + a[2] + \cdots + a[n]$$

random order:

$$sum(a) = a[21] + a[5] + \cdots + a[8]$$

Linear Order - Python

Python:

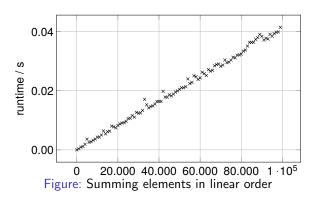
```
def init(size):
    # use system time as seed
    random.seed(None)
    # set linear order as accessor
    order = [a for a in range(0, size)]
    # init array with random data
    data = [random.random() for a in order]
    return (order, data)
```

Linear Order - Python

Python:

```
def run(param):
    # unpack data
    (order, data) = param
    # init the sum value
    s = 0
    for index in order:
        s += data[index]
    return s
```

Linear Order

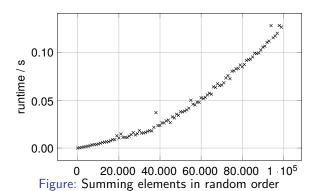


Random Order - Python

Python:

```
def init(size):
    # use system time as seed
    random.seed(None)
    # set random order as accessor
    order = [a for a in range(0, size)]
    random.shuffle(order)
    # init array with random data
    data = [random.random() for a in order]
    return (order, data)
```

Random Order

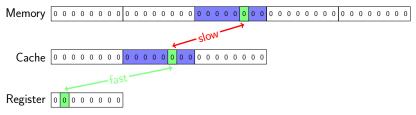


Algorithm Comparision

Conclusion:

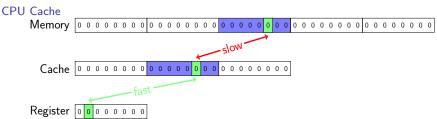
- ▶ The number of operations are identical for both algorithms
- Accessing elements in random order takes a lot longer (Factor 10) Why?
- ▶ The costs in terms of memory access are very different

CPU Cache



Principle / organization:

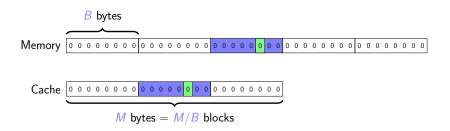
- lacktriangle Accessing one byte of the main memory takes $pprox 100\,\mathrm{ns}$
- ▶ Accessing one byte of (L1-)cache takes $\approx 1 \, \text{ns}$
- ▶ Accessing one or more byte/s of main memory loads a whole block $\approx 100\,\text{B}$ into the cache
- As long as this block is in the cache, it is not neccessary to access the memory for bytes of this block



Cache organization:

- ► The (L1-)cache can hold multiple memory blocks (cache lines)
 - ▶ $\approx 100 \, \text{kB}$
- If the capacity is reached unused blocks are discarded
 - ► Least recently used (LRU)
 - ► Least frequently used (LFU)
 - ► First in first out (FIFO)
- Details of discarding are not the topic for today

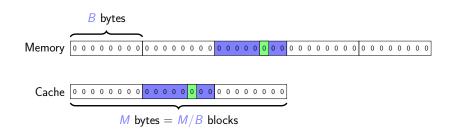
Block Operations



Terminology:

- ▶ The system consists of slow and fast memory
- ► The slow memory is divided in blocks of size *B*
- ▶ The fast cache has size M an can store M/B blocks
- ▶ If data is not in fast memory, the corresponding block is loaded into the cache

Block Operations



Terminology:

- The program defines which blocks are held in the cache
- We use the number of block operations as runtime estimation
- ▶ We ignore runtime costs of cache accesses / management

Block Operations

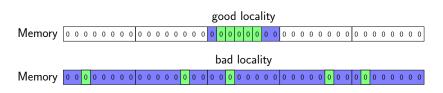


Figure: Comparison good / bad locality

Accessing the cache B times:

- ▶ Best case: 1 block operation → good locality
- Worst case: B block operations → bad locality

Block Operations

Additional factors:

- ► The following settings change only a small constant factor in number of block operations
 - ▶ The partionining of the slow memory into blocks
 - ▶ If the block is 1 Bytes or 4 Bytes or 8 Bytes

Note:

- ▶ If the input size is smaller than *M* we load the complete data chunk directly into the cache
- ► Cache handling is only interesting when the input size is greater than *M*

Block Operations

Typical values: (Intel@ i7-4770 Haswell, WD@ Blue 2TB)

- ► CPU L1 Cache: $B = 64 \, \text{B}$, $M = 4 \times (32 \, \text{kB} + 32 \, \text{kB})$
- ► CPU L2 Cache: $B = 64 \, \text{B}$, $M = 4 \times 256 \, \text{kB}$
- ► CPU L3 Cache: $B = 64 \, \text{B}$, $M = 8 \, \text{MB}$
- ▶ Disk Cache: B = 64 kB, M = 64 MB
 - Many operating systems use free system memory as disk cache

Block Operations

Terminology:

- ▶ Block loads on CPU-cache are called cache misses
- Block operations on disk-cache are called IOs (input / output operations)
- ► These also fall under the term cache efficiency or IO efficiency

Block Operations - Linear Order

Example 1 - Linear order:

▶ We sum up all elements in natural order

$$sum(a) = a[1] + a[2] + \cdots + a[n]$$

▶ The number of block operations is $\operatorname{ceil}\left(\frac{n}{B}\right)$

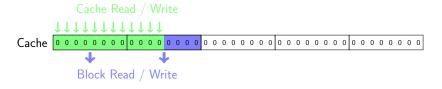


Figure: Good locality of sum operation

Block Operations - Random Order

Example 2 - Random order:

We sum up all elements in random order

$$sum(a) = a[21] + a[5] + \cdots + a[8]$$

- ▶ The number of block operations is *n* in the worst case
- ▶ This leads to a runtime factor difference of *B*

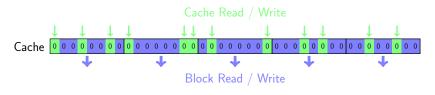


Figure: Bad locality of sum operation

Block Operations

Generally the factor is substantially < B

- Even with a random or5bder we access per element 4 (int) / neighboring bytes at once
- If not n ≫ M the next element might already with a high probability loaded in cache

Block Operations - QuickSort

QuickSort:

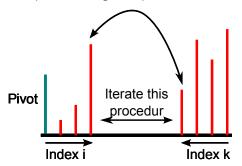
- Strategy: Divide and conquer
- ▶ Divide the data into two parts where the "left" part contains all values ≤ those in the right part
- ► Choose one element (e.g the first one) as "pivot"-element
- Ideally both parts are the same size
- Both parts are sorted recursively

р	list							
lower list	р	upper list						

Figure: QuickSort with pivot-element

Idea of Quicksort

- ▶ at start: pivot in first position, first re-arrange list such that left part contains small, right part larger elements
- do required changes in place



▶ end point: k is left to left-most element greater than pivot swap position 0 (pivot) with k (smaller than pivot)

Block Operations - QuickSort - Python

Python:

```
def quicksort(1, start, end):
   if (end - start) < 1:
      return

i = start
   k = end
   piv = 1[0]</pre>
```

```
Block Operations - QuickSort - Python
   def quicksort(l, start, end):
     while k > i:
       while l[i] <= piv and i <= end and k > i:
         i += 1
       while l[k] > piv and k >= start and k >= i:
         k = 1
       if k > i: # swap elements
         (1[i], 1[k]) = (1[k], 1[i])
     (1[start], 1[k]) = (1[k], 1[start])
     quicksort(l, start, k - 1)
     quicksort(1, k + 1, end)
```

Block Operations - QuickSort

Number of operations for Quicksort:

- Let T(n) be the runtime for the input size n
- Assumptions:
 - Fields are always separated perfectly in the middle
 - ▶ *n* is a power of two and recursion depth is $k = \log_2 n$

Block Operations - QuickSort

$$T(n) \leq \underbrace{A \cdot n}_{\text{splitting in two parts}} + \underbrace{2 \cdot T\left(\frac{n}{2}\right)}_{\text{recursive sort}}$$

$$\leq A \cdot n + 2\left(A \cdot \frac{n}{2} + 2 \cdot T\left(\frac{n}{4}\right)\right)$$

$$= 2A \cdot n + 4 \cdot T\left(\frac{n}{4}\right)$$

$$\leq 3A \cdot n + 8 \cdot T\left(\frac{n}{8}\right)$$

$$\leq \cdots$$

$$\leq k \cdot A \cdot n + 2^k \cdot T(1)$$

$$= \log_2 n \cdot A \cdot n + n \cdot T(1)$$

$$\leq \log_2 n \cdot A \cdot n + n \cdot A \in \mathcal{O}(n \log_2 n)$$

Block Operations - QuickSort

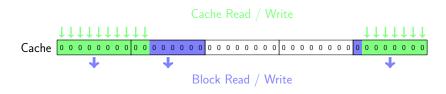


Figure: Locality of quicksort

- Let IO(n) be the number of block operations for input size n
- Assumptions as before but recursion depth is $k = \log_2 \frac{n}{B}$ Why?

Block Operations - QuickSort

 $IO(n) \leq A \cdot n/B + 2 \cdot IO(n/2)$

splitting in two parts recursive sort

$$\leq A \cdot n/B + 2(A \cdot n/2B + 2 \cdot IO(n/4))$$

$$\leq 2 \cdot A \cdot n/B + 4 \cdot IO(n/4)$$

$$\leq 3 \cdot A \cdot n/B + 8 \cdot IO(n/8)$$

$$\leq \cdots$$

$$\leq k \cdot A \cdot n/B + 2^k \cdot IO(n/2^k)$$

$$= \log_2(n/B) \cdot A \cdot (n/B) + n/B \cdot IO(B)$$

$$\leq \log_2(n/B) \cdot A \cdot (n/B) + A \cdot n/B \in O(\frac{n}{B} \cdot \log_2(\frac{n}{B}))$$

Introduction

Concept:

- ▶ Divide the problem into smaller subproblems
- Conquer the subproblems through recursive solving. If subproblems are small enough solve them directly
- Connect all solutions of the subproblems to a solution of the full problem
- Recursive application of the algorithm to ever smaller subproblems
- Direct solving of sufficently small subproblems

Introduction - Python

► Function solve for solving a problem of size *n*

```
def solve(problem):
    if n < threshold:</pre>
        # solve directly
        return solution
    else:
        # divide problem into subproblems
        # P1, P2, ..., Pk with k \ge 2
        S1 = solve(P1)
        S2 = solve(P2)
        Sk = solve(Pk)
        # combine solutions
        return S1 + S2 + ... + Sk
```

Features

- Can help with conceptual hard problems
 - Solution of the trivial problems has to be known
 - Dividing in subproblems has to be possible
 - Combination of solutions has to be possible
- Realization of efficient solutions
 - ▶ If trivial solution is $\in O(1)$
 - ▶ And separation / combination of subproblems is $\in O(n)$
 - And the number of subproblems is limited
 - ▶ The runtime is $\in O(n \cdot \log n)$
- Suitable for parallel processing
 - Subproblems are independent of each other
 - Only needed input for each subproblem has to be known

Implementation

Definition of the trivial case:

- Smaller subproblems are elegant and simple
- Otherwise the efficiency will be improved if relative big subproblems can be solved directly
- Recursion depth should not get too big (stack / memory overhead)

Implementation

Division in subproblems:

 Choosing the number of subproblems and the concrete allocation can be demanding

Combination of solutions:

Typically conceptional demanding

Example - Maximum Subtotal

Example - Maximum Subtotal Input:

▶ Progression *X* of *n* integers

Output:

Maximum sum of related subsequence and its index boundary

Index	0	1	2	3	4	5	6	7	8	9
Value	31	-41	59	26	-53	58	97	-93	-23	84

Output: Sum: 187, Start: 2, End: 6

Example - Maximum Subtotal

Application:

Maximum profit of buying and selling shares



Example - Maximum Subtotal - Python

Naive solution (brute force)

```
def maxSubArray(X):
    # Store sum, start, end
    result = (X[0], 0, 0)
    for i in range(0, len(X)):
        for j in range(i, len(X)):
             subSum = 0
            for k in range(i, j + 1):
                 subSum += X[k]
            if result[0] < subSum:</pre>
                 result = (subSum, i, j)
    return result
```

Example - Maximum Subtotal - Python

Runtime - Upper bound

```
def maxSubArray(X):
    result = (X[0], 0, 0)
    # n loops \rightarrow O(n)
    for i in range(0, len(X)):
         # max n loops \rightarrow O(n)
         for j in range(i, len(X)):
              # max n loops \rightarrow O(n)
              subSum = sum(X[i:j+1])
              if result [0] < subSum: # 0(1)
                  result = (subSum, i, j)
    return result
```

Example - Maximum Subtotal

Upper bound:

- Three interleaved loops
- ► Each loop with runtime *O*(*n*)
- ► Algorithm runtime of $O(n^3)$

Example - Maximum Subtotal - Runtime

Lower bound:

Table: Operations $i \quad | \text{Additions} | \quad j$

 $\frac{n}{3} \in O(n) \mid \frac{n}{3} \in O(n) \mid \frac{n}{3} \in O(n)$

• We iterate at least
$$\frac{n}{3}$$
 values for i

- ► For each *i* we iterate at least $\frac{n}{3}$ values for *j*
- ► For each j we have at least $\frac{n}{3}$ additions
- ▶ We need at least $T(n) = (\frac{n}{3})^3 \in \Omega(n^3)$ steps

Example - Maximum Subtotal - Runtime

Runtime:

▶ With $T(n) \in O(n^3)$ and $T(n) \in \Omega(n^3)$ we know:

$$T(n) \in \Theta(n^3)$$

▶ It is hard to solve the problem in a worse way . . .

Example - Maximum Subtotal - Runtime

Current approach:

► Calculating the sum for range from *i* to *j* with loop

$$S_{i,j} = X[i] + X[i+1] + \cdots + X[j]$$

Better approach:

Incremental sum instead of loop

$$S_{i,j+1} = X[i] + X[i+1] + \dots + X[j] + X[j+1]$$

 $S_{i,j+1} = S_{i,j} + X[j+1] \in O(1)$ instead of $\in O(n)$

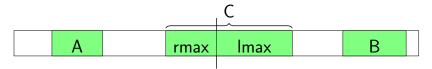
Example - Maximum Subtotal - Python

Better solution:

```
def maxSubArray(X):
    result = (X[0], 0, 0)
    # n loops \rightarrow O(n)
    for i in range(0, len(X)):
         subSum = 0
         # max n loops \rightarrow O(n)
         for j in range(i, len(X)):
              subSum += X[j] # O(1)
             if result [0] < subSum: # 0(1)
                  result = (subSum, i, j)
    return result
 ▶ Runtime \in O(n^2)
```

Example - Maximum Subtotal

Divide and Conquer:



Divide and Conquer Idea to solve:

- split the sequence in the middle
- ► Solve the left half of the problem
- Solve right half and combine both solutions into a total solution
- ► OK if maximum is located in left half (A) or right half (B)
- ▶ Problem: Maximum can overlap split
- ▶ To solve this case we have to calculate *rmax* and *lmax*
- ▶ The overall solution is the maximum of A, B and C

Example - Maximum Subtotal

Principle - Divide and Conquer:

- ▶ Small problems are solved directly: $n = 1 \Rightarrow \max = X[0]$
- ▶ Bigger problems are partitioned into two subproblems and recursivly solved. Subsolutions A and B are returned.
- ► To determine subsolution C, rmax and Imax for the subproblems are computed.
- ▶ The overall solution is the maximum of A, B and C

Example - Maximum Subtotal - Python

Divide and conquer solution

```
def maxSubArray(X, i, j):
    if i == j: #trivial case
        return (X[i], i, i)
    m = (i + j) / 2
        #recursive Subsolutions for A,B
    A = \max SubArray(X, i, m)
    B = \max SubArray(X, m + 1, j)
    #rmax and lmax for bordercase C
    C1 = rmax(X, i, m)
    C2 = lmax(X, m + 1, j)
    C = (C1[0] + C2[0], C1[1], C2[1])
    #Solution results from A,B,C
    return max([A, B, C], \
        key=lambda item: item[0])
```

Further Literature

General

- [CRL01] Thomas H. Cormen, Ronald L. Rivest, and Charles E. Leiserson. Introduction to Algorithms. MIT Press, Cambridge, Mass, 2001.
- [MS08] Kurt Mehlhorn and Peter Sanders.
 Algorithms and data structures, 2008.
 https://people.mpi-inf.mpg.de/~mehlhorn/
 ftp/Mehlhorn-Sanders-Toolbox.pdf.

Further Literature

Caching

```
[Wik] Cache https://en.wikipedia.org/wiki/Cache
```