### APL SIMD Boolean Array Algorithms

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#### Abstract

Computation on large Boolean arrays is becoming more prevalent, due to applications such as cryptography, data compression, and image analysis and synthesis. The advent of bit-oriented vector extensions for microprocessors and of GPUS presents opportunities for significant performance improvements in such Boolean-dominated applications. Since APL is one of the few computer languages that supports dense (one bit per element, eight bits per byte), multi-dimensional Boolean arrays as first-class objects, it has naturally attracted research into optimizations for improved performance of Boolean array operations. This paper presents some of the Single Instruction, Multiple Data (SIMD) Boolean-related optimizations that have appeared in APL implementations, and suggests ways in which those optimizations might be exploited using contemporary hardware.

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- GPU and SIMD vector facilities can exploit them

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- Performance boosts: In a compiler, opportunity for other optimizations

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# Structural and Selection Verbs I

- catenate, laminate, rotate, reverse, rank, from,
- ▶ merge, take, drop...

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- Operates in SIMD mode(s) whenever possible
- Supports all type conversions

Operation on non-trailing array axes:

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- Operation on non-trailing array axes:
- ► SIMD copy entire subarrays at once, e.g. 1⊖2 3 4P124

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- Operation on non-trailing array axes:
- SIMD copy entire subarrays at once, e.g. 102 3 40124
- rbemove will copy 12 adjacent array elements at once

# Structural and Selection Verbs III

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# Structural and Selection Verbs III

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Bernecky, 1979: fast algorithms for φω & αφω

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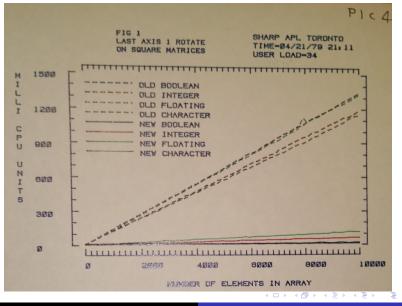
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- last-axis Boolean Φω did a byte at a time, w/table lookup... RevTab[uint8 ω]
- ▶ then byte-aligned the resulting vector, SIMD, a word at a time
- All non-last-axis operations copied entire cells at once, using rbemove

### Reverse and Rotate Performance on Booleans



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#### ► Reshape allows element reuse, e.g.: 8P1 0 0 1 0 0 1 0 0 1 0

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 Breed optimized Boolean reshape this way:

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801 0 0

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- Breed optimized Boolean reshape this way:
- Copy the argument to the result
- Catenate the partial result to itself, doubling its length, until its tail is byte-aligned.
- ► Do an overlapped move, or "smear" of the result to its tail

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► An unlikely candidate for SIMD, it would seem...

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## Transpose I

- An unlikely candidate for SIMD, it would seem...
- α\u03c0\u03c0 with unchanged trailing axes

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# Transpose I

- An unlikely candidate for SIMD, it would seem...
- $\alpha \otimes \omega$  with unchanged trailing axes
- ► T←2 2 2 3P124
  - 0 1 2
  - 3 4 5
  - 6 7 8 9 10 11

- 12 13 14
- 15 16 17
- 18 19 20
- 21 22 23

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Transpose with unchanged trailing axes

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# Transpose II

- Transpose with unchanged trailing axes
- SIMD copy six elements at once (rbemove)

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- Transpose with unchanged trailing axes
- SIMD copy six elements at once (rbemove)
- ▶ 1 0 2 3\T
  - 0 1 2
  - 3 4 5
  - 12 13 14 15 16 17

- 6 7 8 9 10 11
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Hacker's Delight: fast 8x8 Boolean matrix transpose

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- Hacker's Delight: fast 8x8 Boolean matrix transpose
- ► Kernel: 16 logical & shift operations on 64-bit ravel

A B K A B K

- Hacker's Delight: fast 8x8 Boolean matrix transpose
- Kernel: 16 logical & shift operations on 64-bit ravel
- ► Uses *perfect shuffle* (PDEP) on any power of two shape

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- ► Kernel generalizes to any power of two, *e.g.*, 16×16, 32×32

#### Bernecky, 1971: fast indexof and set membership

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• Algorithm used briefly for  $\vee/\omega$  and  $\wedge/\omega$ 

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 E.E. McDonnell, 1974: elegant APL models of Boolean scan and reduction for relationals

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- Result: linear-time, word-at-a-time, SIMD Boolean scan & reduce

#### ► John Heckman, 1970 or 1971: user-defined APL scan verbs

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- Recursive doubling:
  - r←nescanall y;s;biw
    - A Not-equal scan
    - r≁y
    - biw←「2⊗1「Py
    - :For s :In 2\*1biw A Heckman
      - r←r≠(-pr)↑(-s)↓r
    - :EndFor
- ► SIMD, word-at-a-time algorithm for Boolean ≠\ω and =\ω along last axis

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► So far, no X86 vectorization; perhaps we can do even better

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- ► redesigned Boolean inner product to use STAR algorithm

```
Z+X ipclassic Y;RX;CX;CY;I;J;K
RX \leftarrow (\rho X) [0]
CX \leftarrow (\rho X) [1]
CY \leftarrow (PY)[1]
Z \leftarrow (RX, CY) \rho 0.5
:For I :In 1RX
  :For J :In 1CY
   Z[I;J]←0
   :For K : Tn 1CX
     Z[I;J] \leftarrow Z[I;J] + X[I;K] \times Y[K;J]
    : EndFor
  : EndFor
: EndFor
```

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```
Z+X ipstar Y;RX;CX;CY;I;J;Xel
RX \leftarrow (\rho X) [0]
CX \leftarrow (\rho X) [1]
CY \leftarrow (PY)[1]
Z \leftarrow (RX, CY) \rho 0
 :For I :In 1RX
  :For J :In 1CX
   Xel + X[I;J]
   Z[I;] \leftarrow Z[I;] + Xel \times Y[J;]
  :EndFor
 :EndFor
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- Unfortunately, the APL primitive is still 30X faster than the APL model

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- Final result: Boolean inner products on SHARP APL/PC ran much faster than APL2 on huge mainframe

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- See also the Bernecky-Scholz PLDI2014 Arrays Workshop paper: Abstract Expressionism

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- Boolean sort can use Moore's SIMD +/Boolean in its first phase of execution
- Second phase can be performed in SIMD, e.g., by a single SAC data-parallel with-loop.

SIMD Boolean upgrade: ug ← { ((~ω)/ιρω), ω/ιρω }

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- SIMD Boolean upgrade: ug ← { ((~ω)/ιρω), ω/ιρω }
- Not stunningly SIMD, though.

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- A similar definition holds for word-oriented algorithms

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My sincere thanks to James A. Brown, Larry M. Breed, Walter Fil, Jay Foad, Roger K.W. Hui, Roger D. Moore, and Bob Smith, for their constructive suggestions and ideas regarding this paper. The APL programs in this paper were executed on Dyalog APL Version 15.0. Dyalog provides free downloads of their interpreters for educational use; they also offer a free download of their Raspberry Pi edition, at www.dyalog.com. The British APL Association (BAA) provided the author with financial assistance for attending this conference; their thoughtfulness and generosity is greatly appreciated.

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