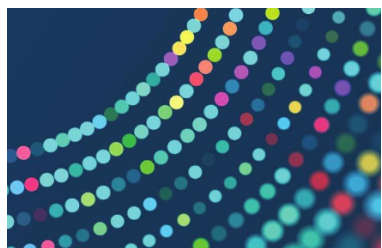


# Fast Interpolation of Grid Data at a Non-Grid Point

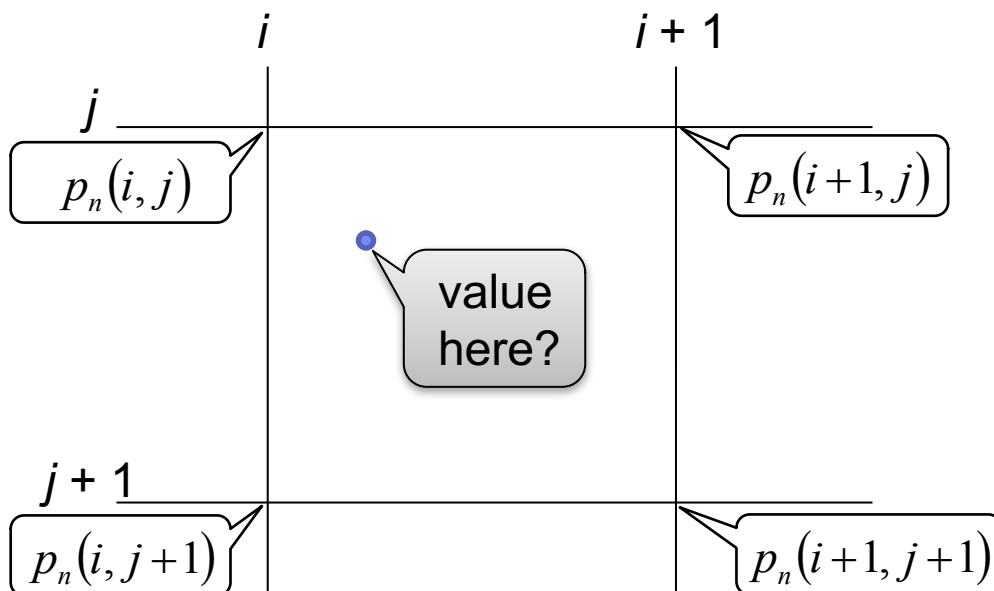


Hiroshi Inoue  
IBM Research – Tokyo

## Interpolation from Grid Data

Goal: to make compute-intensive interpolation operation faster

- Input: values at grid points
- Output: estimated (interpolated) value at a non-grid point



Target workloads include:

- medical imaging
  - CT reconstruction
  - registration etc
- stencil applications
  - particle simulation etc

## Contributions

- Developed an fast method to interpolate values from grid data at a non-grid point
- Evaluated with 3D Computed Tomography (CT) reconstruction benchmark (RabbitCT)
  - this technique itself can be applicable for other imaging and non-imaging applications
  - although we explain the technique using bi-linear interpolation in this talk, it is applicable for more accurate interpolation algorithms (See paper for detail)

# CT Reconstruction Overview

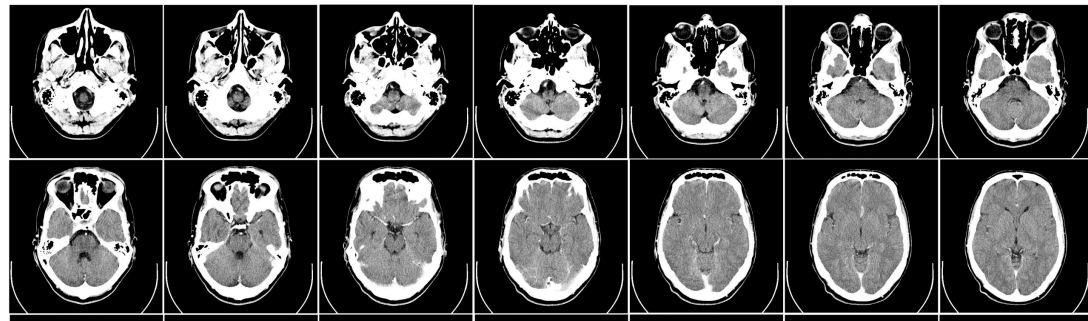
- Input: a set of 2D projection images obtained from different angles (and geometry information for each image)
- Output: density values for voxels in a 3D volume

## Example of a (C-arm) CT system



Source: <http://www.sharpmedical.com/refurbished-c-arms/ziehm-c-arms/ziehm-exposcop-7000-c-arm/>

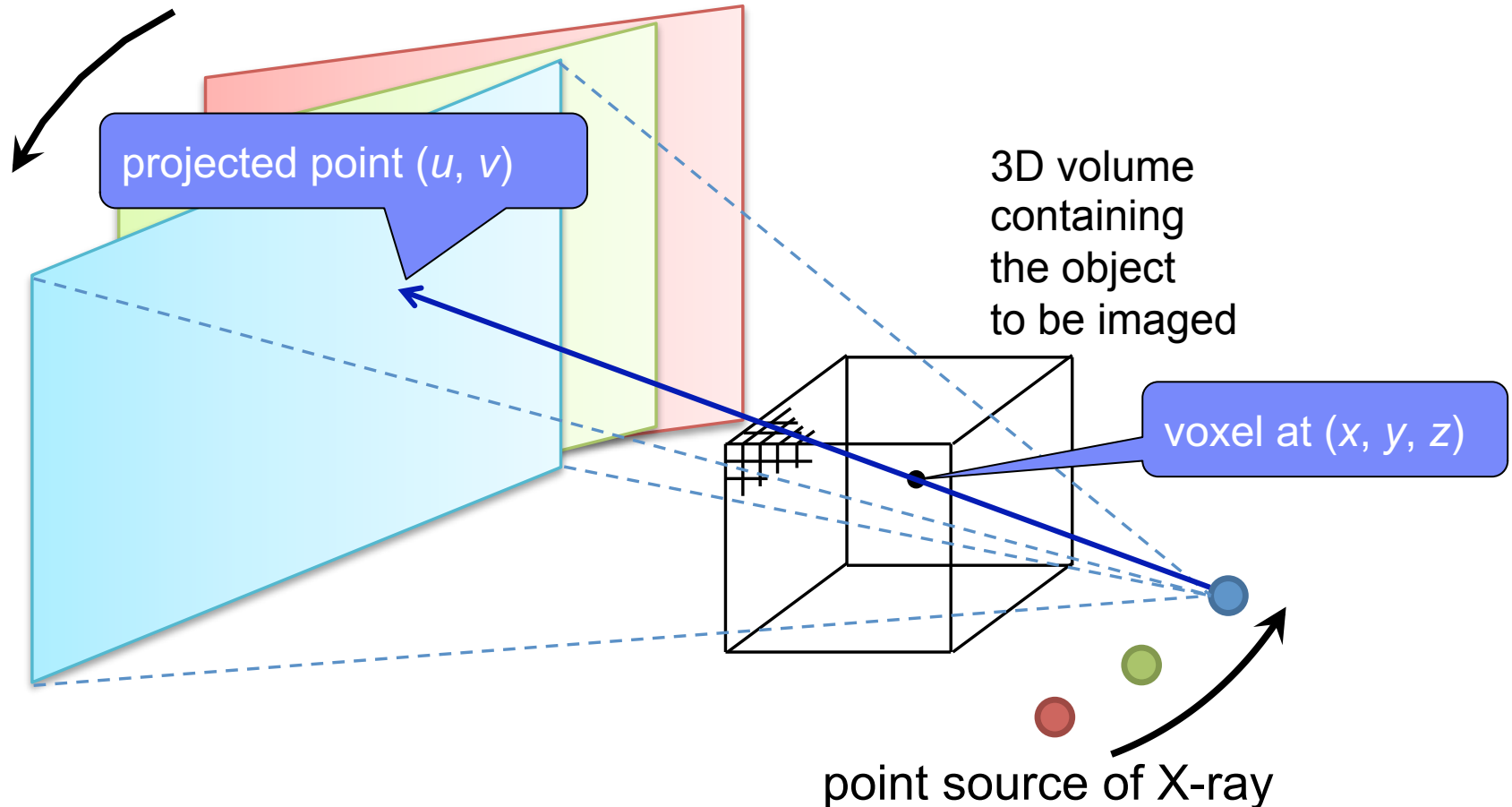
## Example of output from a CT system



Source: [https://en.wikipedia.org/wiki/CT\\_scan](https://en.wikipedia.org/wiki/CT_scan)

## Projection in CT Reconstruction

Flat-panel detector (capturing 2D projection images)



## Baseline Reconstruction Algorithm Overview [2]

```
for each projection image  $I_n$ 
  // reconstruction (projection)
  for  $z = 0$  to  $L-1$ 
    for  $y = 0$  to  $L-1$ 
      for  $x = 0$  to  $L-1$ 
        1) project voxel  $(x,y,z)$  onto  $I_n$ 
        2) read values from surrounding four grid points
        3) estimate value at projected point by interpolation
        4) update density value of voxel  $(x,y,z)$ 
      end
    end
  end
end
end
```

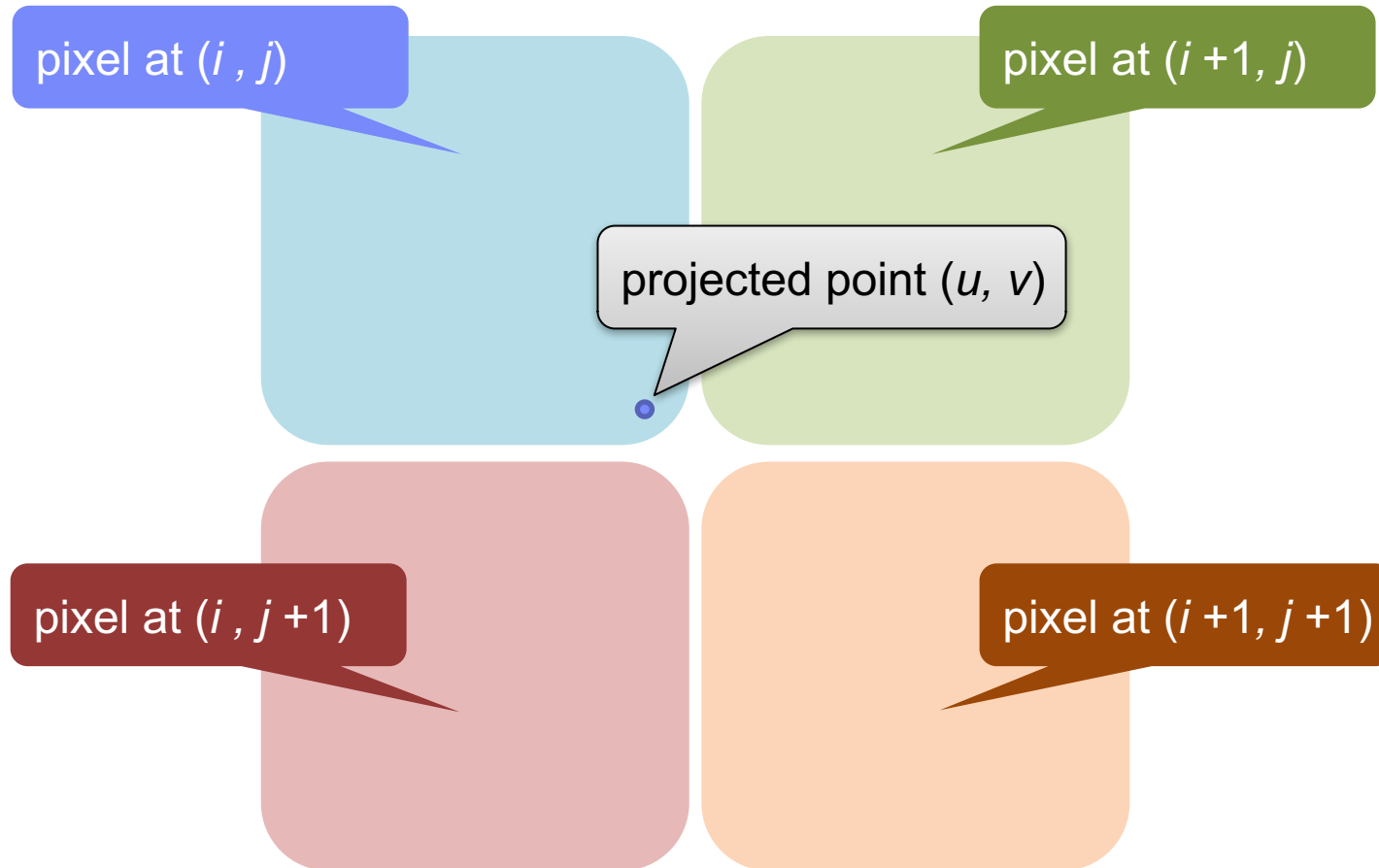
} for each voxel in 3D volume

## Baseline Reconstruction Algorithm Overview [2]

```
for each projection image  $I_n$ 
  // reconstruction (projection)
  for  $z = 0$  to  $L-1$ 
    for  $y = 0$  to  $L-1$ 
      for  $x = 0$  to  $L-1$ 
        1) project voxel  $(x,y,z)$  onto  $I_n$ 
        2) read values from surrounding four grid points
        3) estimate value at projected point by interpolation
        4) update density value of voxel  $(x,y,z)$ 
      end
    end
  end
end
```

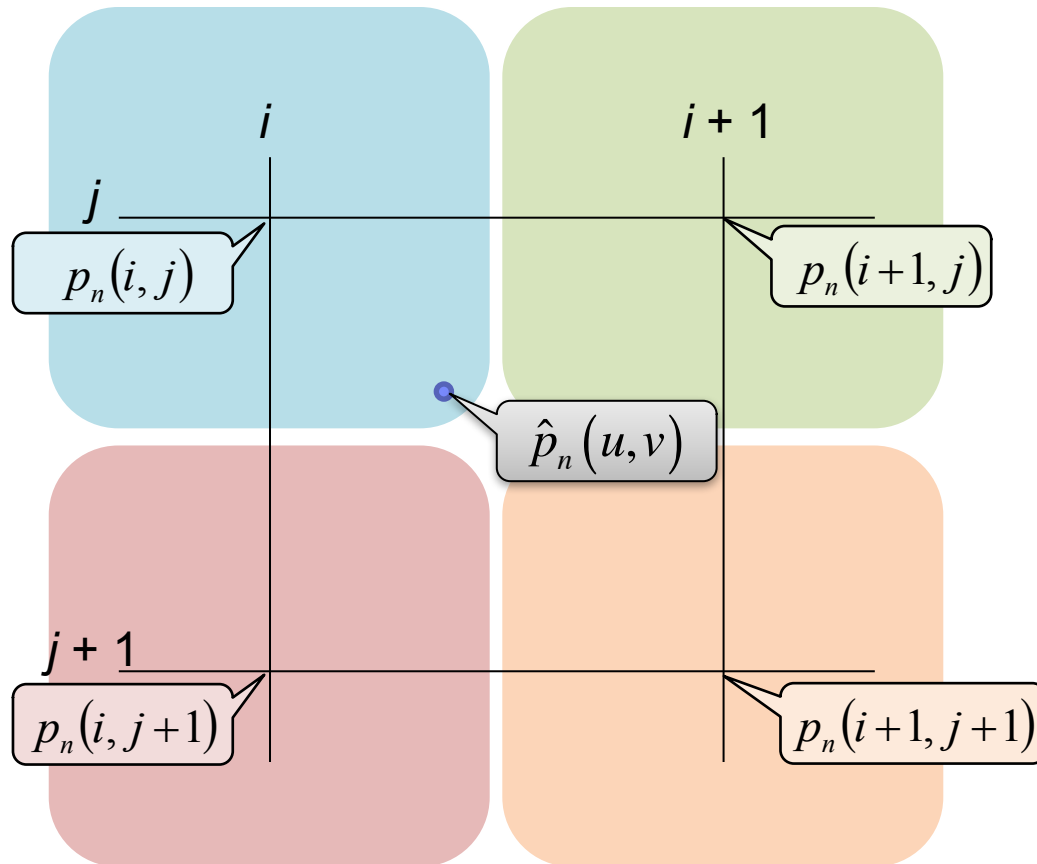
} for each voxel in 3D volume

# Interpolation from Grid Data at Non-Grid Point

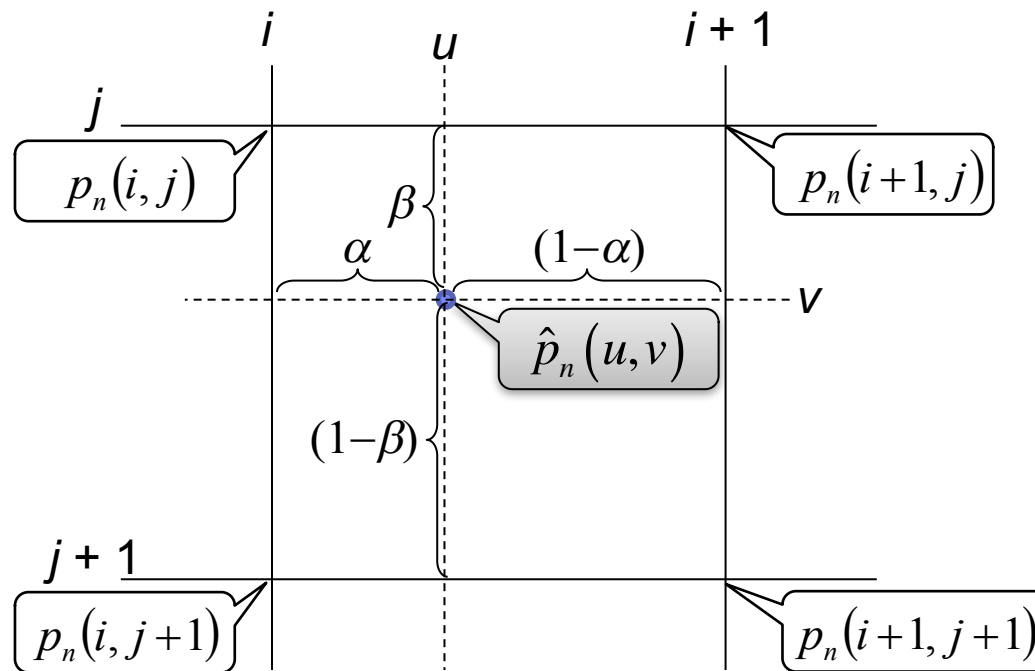




# Interpolation from Grid Data at Non-Grid Point



# Interpolation from Grid Data at Non-Grid Point



$$\alpha = u - i = u - \lfloor u \rfloor$$

$$\beta = v - j = v - \lfloor v \rfloor$$

## Bilinear interpolation

$$\hat{p}_n(u, v) = (1 - \alpha)(1 - \beta)p_n(i, j) + \alpha(1 - \beta)p_n(i + 1, j) + (1 - \alpha)\beta p_n(i, j + 1) + \alpha\beta p_n(i + 1, j + 1)$$

## Do we have any redundancy in this formula?

$$\hat{p}_n(u, v) = (1 - \alpha)(1 - \beta)p_n(i, j) + \alpha(1 - \beta)p_n(i + 1, j) + (1 - \alpha)\beta p_n(i, j + 1) + \alpha\beta p_n(i + 1, j + 1)$$

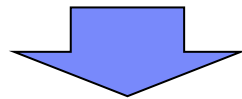
😊 Yes, we can simplify it if many interpolation operations are done in one grid

- In CT reconstruction, hundreds to thousands of interpolations are executed in one grid on average

➔ Now, think  $p_n(i, j)$ ,  $p_n(i + 1, j)$ ,  $p_n(i, j + 1)$ ,  $p_n(i + 1, j + 1)$  as constant values

## Do we have any redundancy in this formula?

$$\begin{aligned}
 \hat{p}_n(u, v) &= (1 - \alpha)(1 - \beta)p_n(i, j) + \alpha(1 - \beta)p_n(i + 1, j) + (1 - \alpha)\beta p_n(i, j + 1) + \alpha\beta p_n(i + 1, j + 1) \\
 &= (1 - \alpha - \beta + \alpha\beta) \cdot p_n(i, j) \\
 &\quad + (\alpha - \alpha\beta) \cdot p_n(i + 1, j) \\
 &\quad + (\beta - \alpha\beta) \cdot p_n(i, j + 1) \\
 &\quad + \alpha\beta \cdot p_n(i + 1, j + 1)
 \end{aligned}$$



Group terms by the number of  $\alpha$  and  $\beta$

$$= \alpha\beta \cdot C_0(i, j) + \alpha \cdot C_1(i, j) + \beta \cdot C_2(i, j) + C_3(i, j)$$

$$C_0(i, j) = p_n(i, j) - p_n(i + 1, j) - p_n(i, j + 1) + p_n(i + 1, j + 1)$$

## Our Efficient Interpolation Method

In the paper, we group terms based on  $u$  and  $v$  instead of  $\alpha$  and  $\beta$  for further performance boost

$$\alpha = u - i = u - \lfloor u \rfloor \quad \beta = v - j = v - \lfloor v \rfloor$$

$$\begin{aligned} \hat{p}_n(u, v) &= (1 - \alpha)(1 - \beta)p_n(i, j) + \alpha(1 - \beta)p_n(i + 1, j) + (1 - \alpha)\beta p_n(i, j + 1) + \alpha\beta p_n(i + 1, j + 1) \\ &= \alpha(\beta \cdot C_0(i, j) + C_1(i, j)) + (\beta \cdot C_2(i, j) + C_3(i, j)) \end{aligned}$$

➔ If we have these four coefficients  $C_0 - C_3$ , we can compute this formula with only **three multiply-and-add instructions!**

$C_0 - C_3$  are independent from  $\alpha$  and  $\beta$  (and hence  $x, y, z$ )

➔ We can pre-compute them at run time before iterating voxels and store in memory

original (without pre-computation)

for each projection image

end

projection

with pre-computation

for each projection image

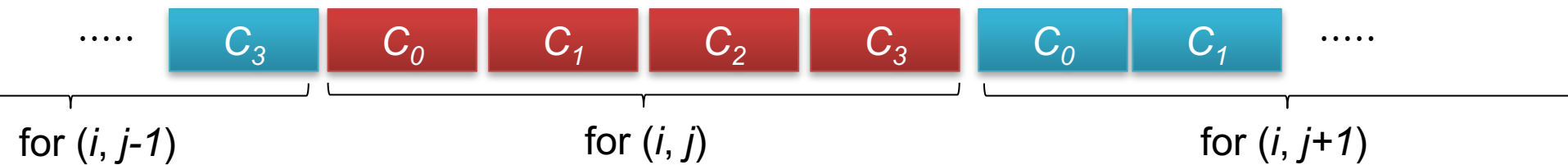
end

pre-computation

projection

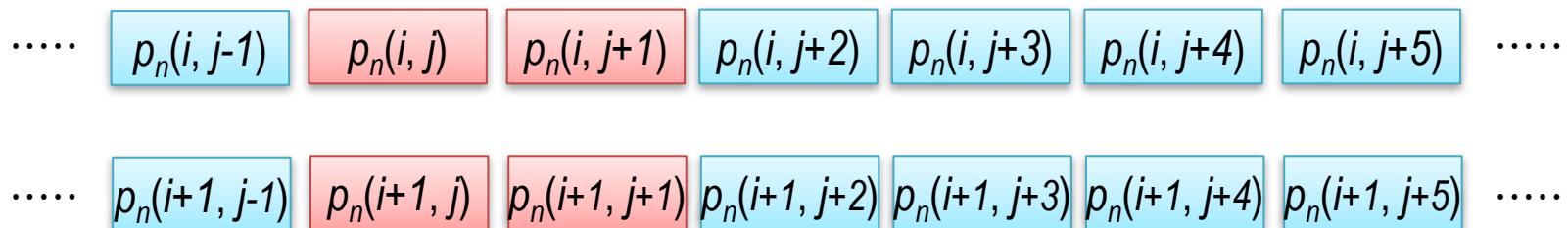
# In-memory Pre-computed Table

## Pre-computed Table



- ➔ 😞 total size of pre-computed table is 4x larger than original data
- ➔ 😊 need to read from only one cache line (w/ one aligned vector load)

Original data (Projection image) ➔ 😞 need to read from two cache lines



## Overall Algorithm with Pre-Computation

```

for each projection image  $I_n$ 
  // pre-computation
  for  $i = 0$  to  $S_x-1$ 
    for  $j = 0$  to  $S_y-1$ 
      } for each pixel in 2D projection image
      calculate and store coefficients  $C_0$  to  $C_3$  for pixel ( $i, j$ )
    end
  end
  // reconstruction (projection)
  for  $x = 0$  to  $L-1$ 
    for  $y = 0$  to  $L-1$ 
      for  $z = 0$  to  $L-1$ 
        } for each voxel in 3D volume
        1) project voxel ( $x, y, z$ ) onto  $I_n$ 
        2) read coefficients  $C_0$  to  $C_3$  from pre-computed table
        3) estimate value at projected point by interpolation
        4) update density value of voxel ( $x, y, z$ )
      end
    end
  end
end

```

## Performance Evaluation with RabbitCT

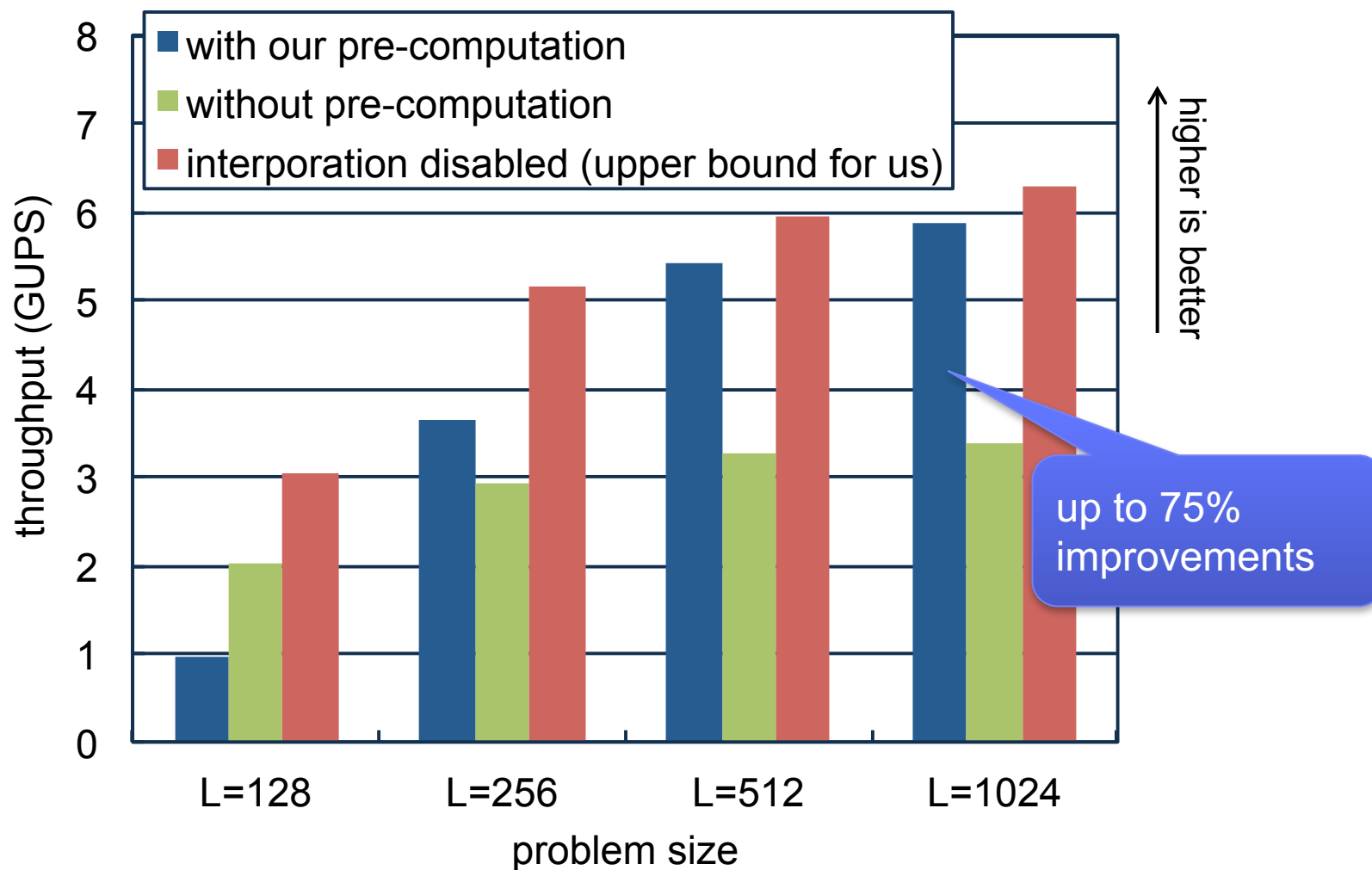
- RabbitCT is a framework for evaluating 3D CT reconstruction on performance and accuracy
- It includes:
  - benchmark driver
  - reference implementations of the backprojection algorithm
  - input data (a C-arm CT dataset of a rabbit)
    - 496 projection images of 1248x960 pixels associated with transformation matrixes
- Output data is 3-D images of  $256^3$  mm<sup>3</sup> space,  $L^3 = 128^3, 256^3, 512^3, 1024^3$  voxels, 12-bit value per voxel



## System used for evaluations

- 2-socket POWER8 3.69 GHz
  - 20 cores in total (5 cores / NUMA node)
  - 8 SMT threads / core
- 256 GB system memory
- Ubuntu Linux 14.10 for Little Endian POWER
- IBM XL C compiler 13
  - all algorithms are implemented with VSX (128-bit SIMD instructions) using intrinsics

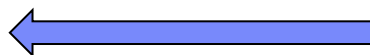
# Throughput with and without pre-computation



# Root Mean Squared Error

(lower is better)

Problem size	With pre-computation	Without pre-computation	Interpolation disabled
$L=128$	0.534	0.513	12.088
$L=256$	0.538	0.517	12.108
$L=512$	0.538	0.518	12.118
$L=1024$	0.545	0.526	12.120



😊 negligible degradation  
in image quality



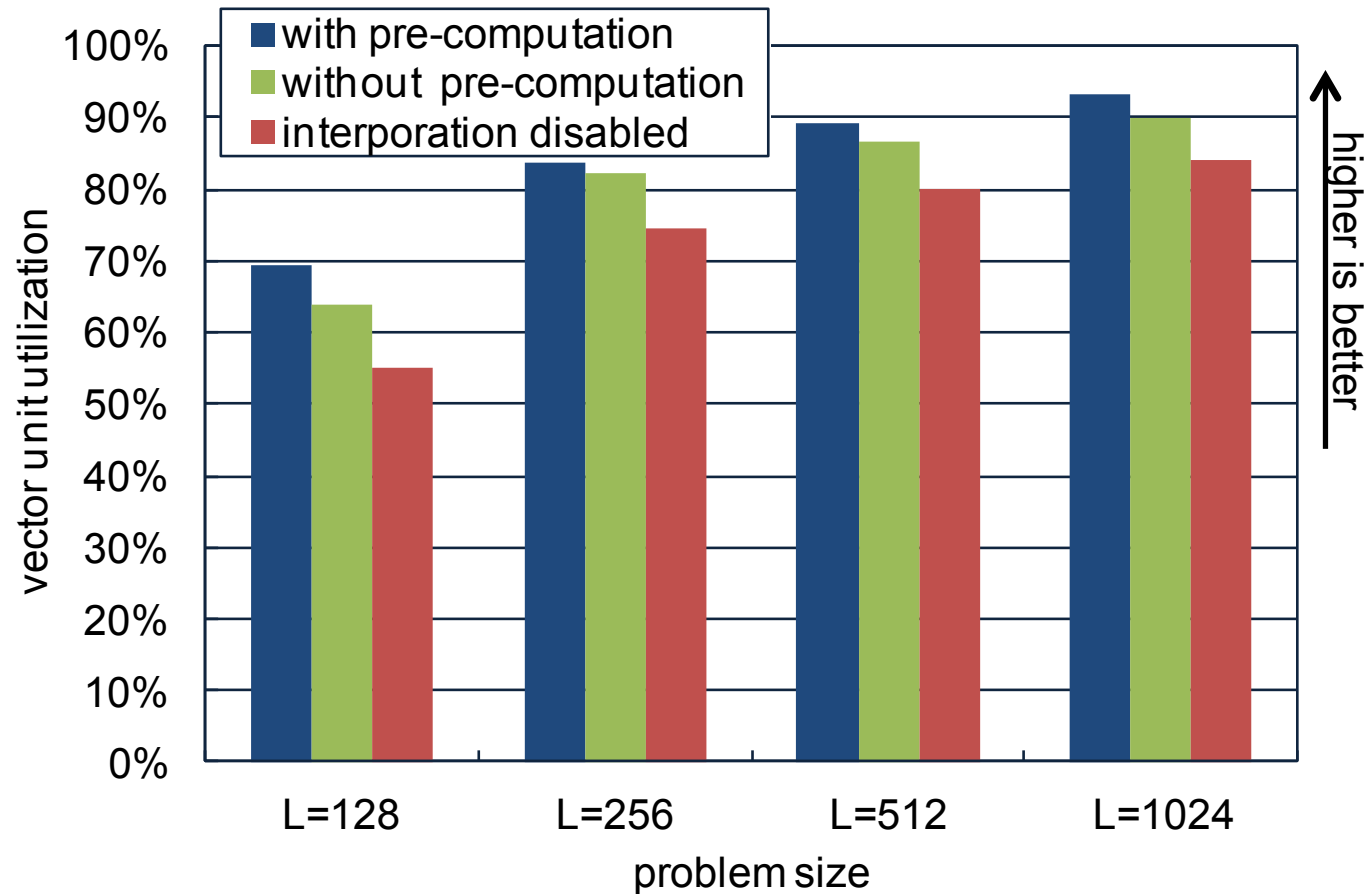
😞 significant degradation  
in image quality

## Overhead of Pre-Computation

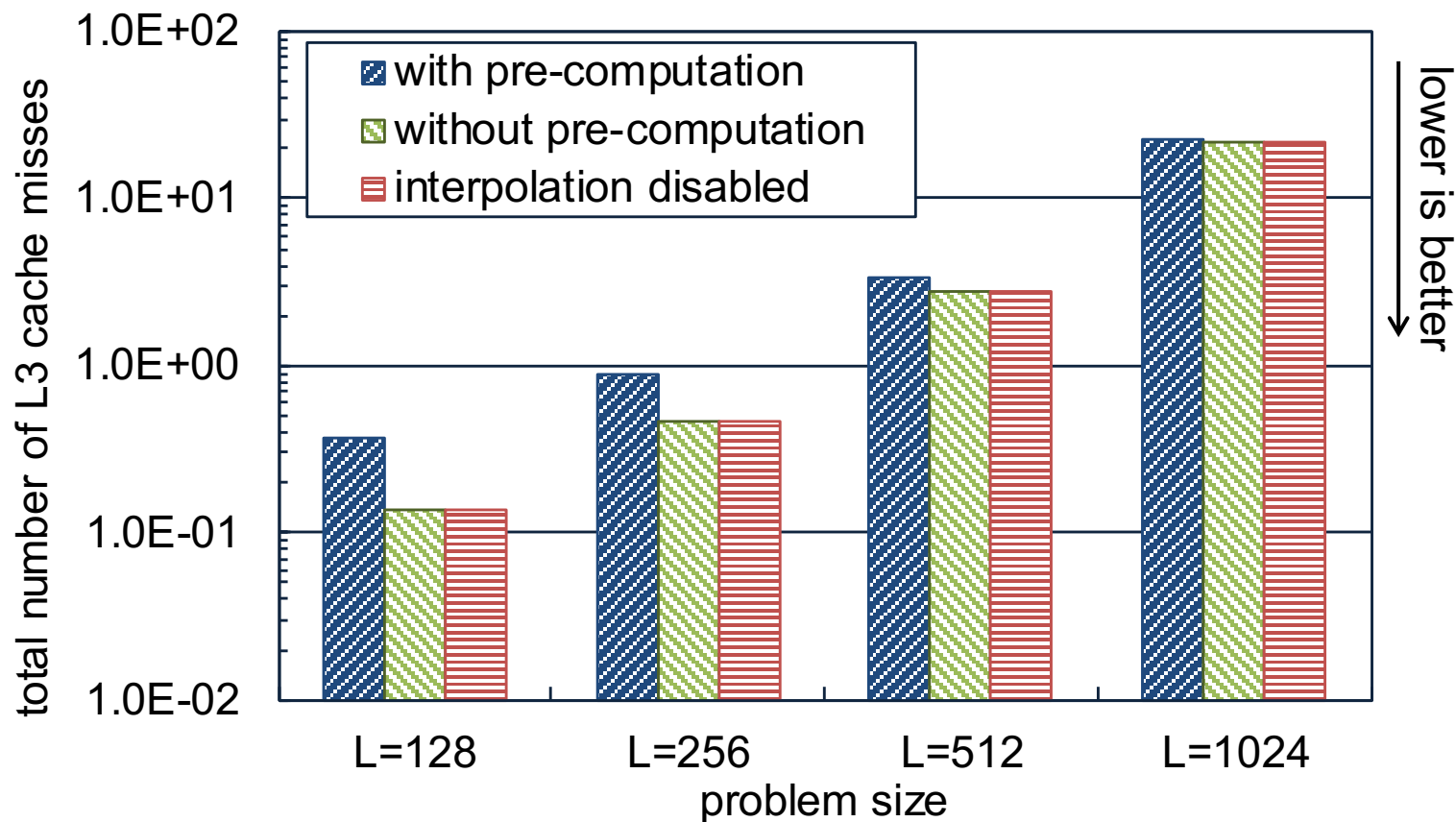
Problem size	With pre-computation			Without pre-computation
	precomputation	reconstruction	total	
$L=128$	0.93 msec (47%)	1.11 msec	2.01 msec	0.94 msec
$L=256$	0.94 msec (22%)	3.36 msec	4.19 msec	5.21 msec
$L=512$	0.94 msec (4.1%)	22.20 msec	22.59 msec	37.90 msec
$L=1024$	0.93 msec (0.6%)	169.30 msec	167.02 msec	293.87 msec

- The numbers show the execution time per projection image.
- The percentages shown in parenthesis show the ratios to the total execution time.
- Average numbers of interpolations (i.e # voxles) per pixel
  - $L=128 \rightarrow 1.75$
  - $L=1024 \rightarrow 896$

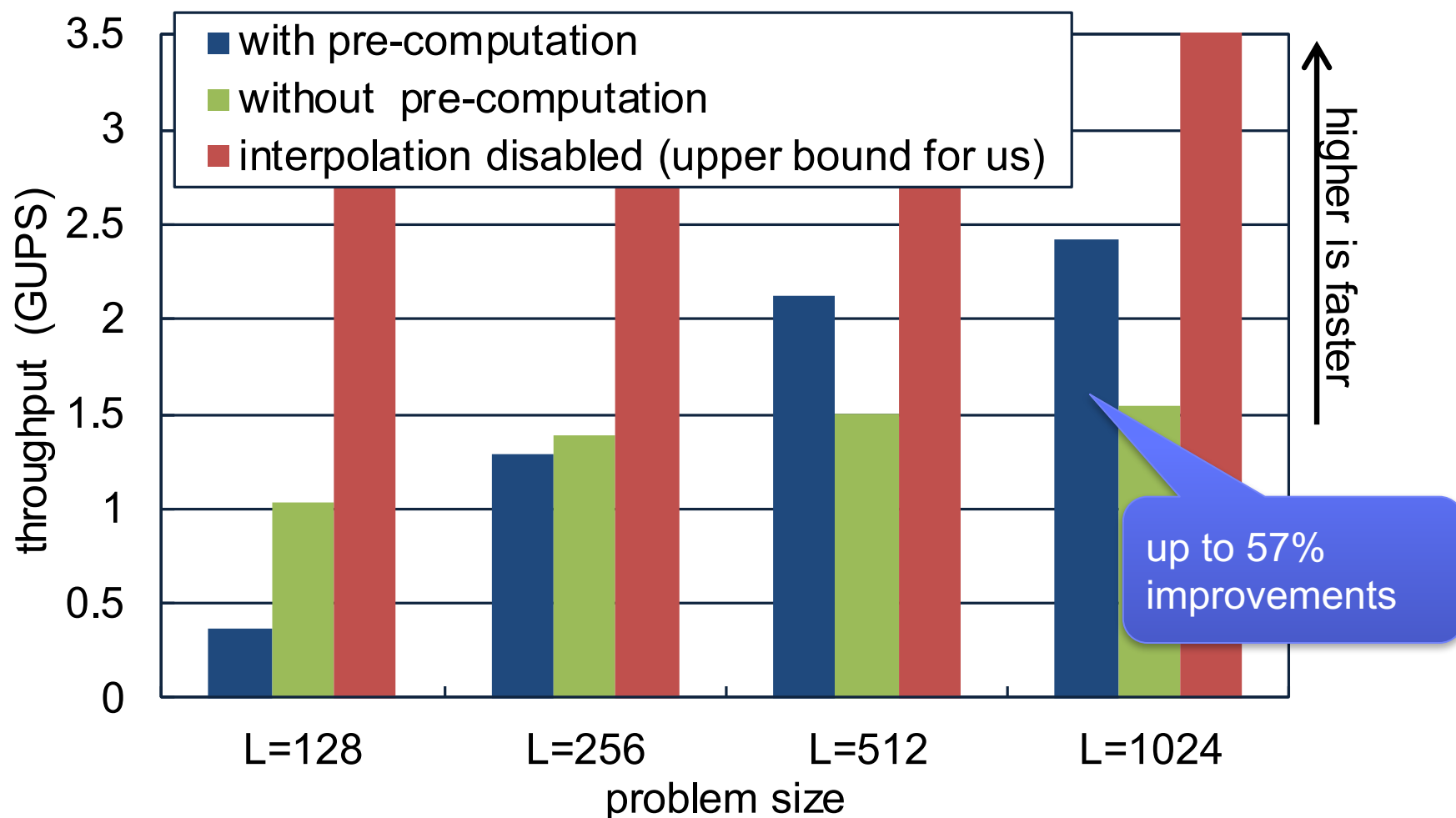
# Vector Unit Utilization



# System memory bandwidth requirements



# Throughput with and without pre-computation using 3rd degree Lagrange interpolation



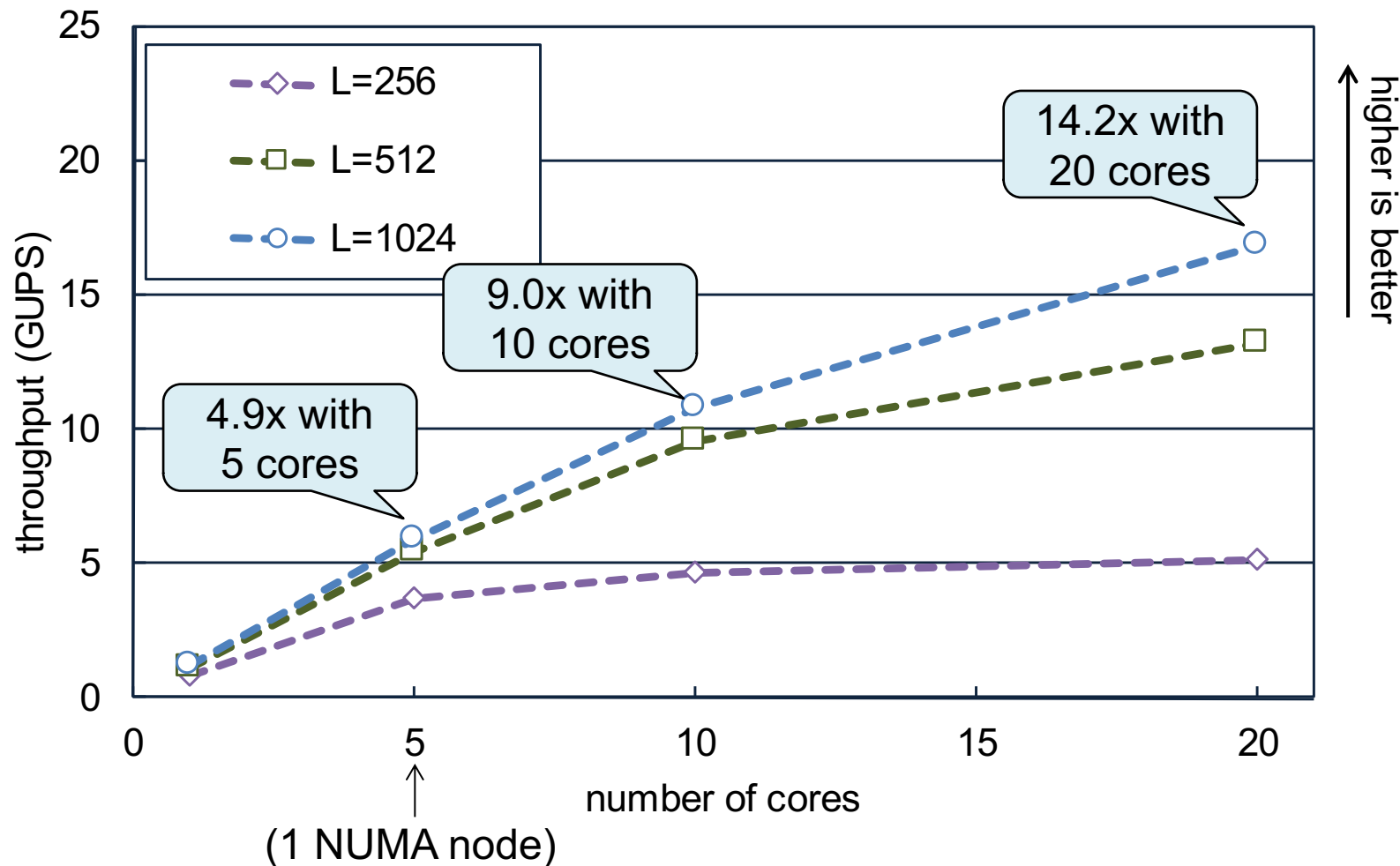
## Summary

- We developed an efficient way of interpolation from grid data at a non-grid point
  - Our pre-computation simplifies the computation drastically
  - The cost of pre-computation is not significant for realistic data size
- This technique is not specialized for CT reconstruction and applicable for other applications
- Refer the paper for more detail including:
  - Results for a more accurate interpolation algorithm
  - Performance modeling    — Handling of floating point errors
  - NUMA optimization



backup

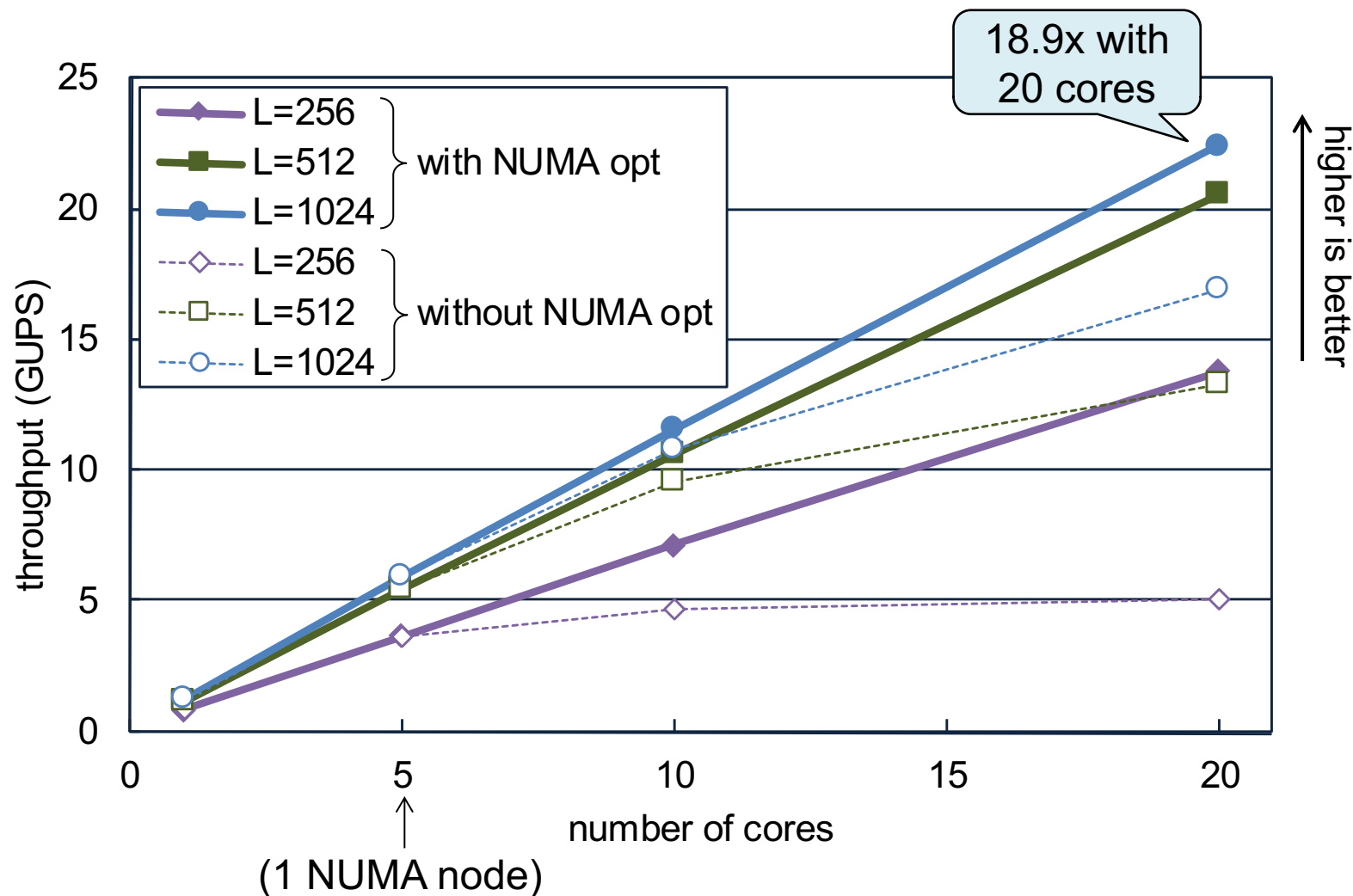
# Scalability with Multiple NUMA Nodes



## Memory Optimization for NUMA Machine

- Each NUMA node processes a projection image independently from other NUMA nodes to avoid remote memory accesses
  - Within a NUMA node, all threads process one projection image by dividing voxels into small blocks
- We gather the partial results from each NUMA nodes after processing all projection images to sum up them

# Scalability with Multiple NUMA Nodes



## Comparing to Previous RabbitCT Scores (L=512)

Processor	# Core / # Boards	Year	Source	GUPS
POWER8 3.69 GHz	20 cores (2 sockets)	2015	Ours	20.5
POWER8 3.69 GHz	10 cores (1 socket)	2015	Ours	10.6
IvyBridge-EP 2.2 GHz	20 cores (2 sockets)	2014	Paper [4]	about 7.0
Westmere-EX 2.4 GHz	40 cores (4 sockets)	2011	Official ranking	8.3
Xeon Phi 5110P	1 board	2014	Paper [4]	about 8.5
nVidia GTX 670	2 boards	2014	Official ranking	152.9

- Today's GPUs support bilinear interpolation in hardware!
- Our method will be beneficial even for GPUs when a higher-order interpolation algorithm is used