

- Task: compute "similarity" of documents (think Google)
 - One of the fundamental tasks in information retrieval (IR)
- Example: search engine / database of scientific papers needs to suggest similar papers for a given one
- Assumption: all documents are over a given, fixed vocabulary V consisting of N words (e.g., all English words)
 - Consequence: set of words, V , occurring in the docs is known & fixed
- Assumption: don't consider word order \rightarrow bag of words model
 - Consequence: "*John is quicker than Mary*" = "*Mary is quicker than John*"

- Representation of a document D :

- For each word $w \in V$: determine $f(w)$ = frequency of word w in D

- Example:

	Anthony & Cleopatra	Julius Caesar	The Tempest	Hamlet	Othello	Macbeth
ANTHONY	157	73	0	0	0	1
BRUTUS	4	157	0	2	0	0
CAESAR	232	227	0	2	1	0
CALPURNIA	0	10	0	0	0	0
CLEOPATRA	57	0	0	0	0	0
MERCY	2	0	3	8	5	8
WORSER	2	0	1	1	1	5
...

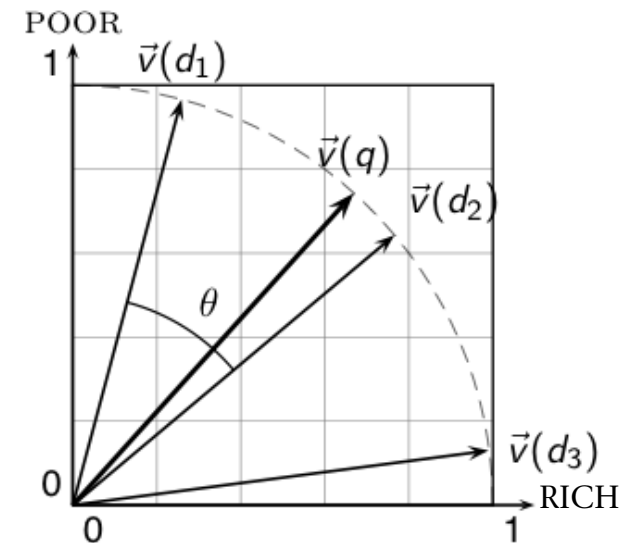
- Fix a word order in $V = (v_1, v_2, v_3, \dots, v_N)$ (in principle, any order will do)
- Represent D as a vector in \mathbf{R}^N :

$$D = (f(v_1), f(v_2), f(v_3), \dots, f(v_N))$$

- Note: our vector space is HUGE ($N \sim 100,000 - 10,000,000$)
 - For each word w , there is one axis in our vector space!

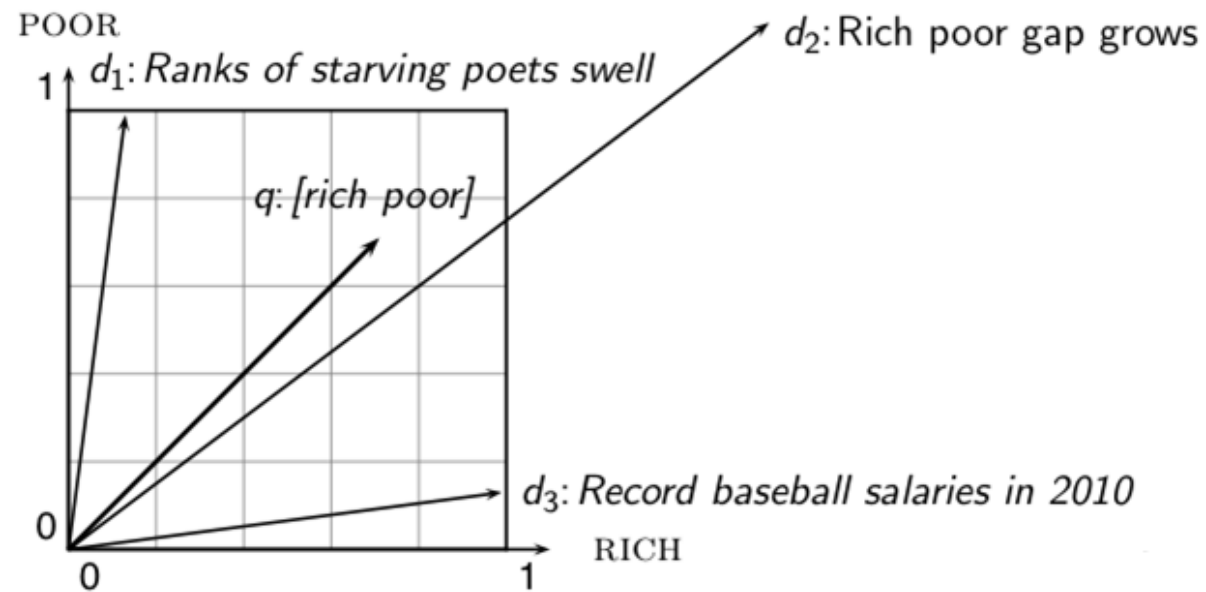
- Define similarity s between documents D_1 and D_2 as

$$s(D_1, D_2) = \frac{D_1 \cdot D_2}{\|D_1\| \cdot \|D_2\|} = \cos(D_1, D_2)$$



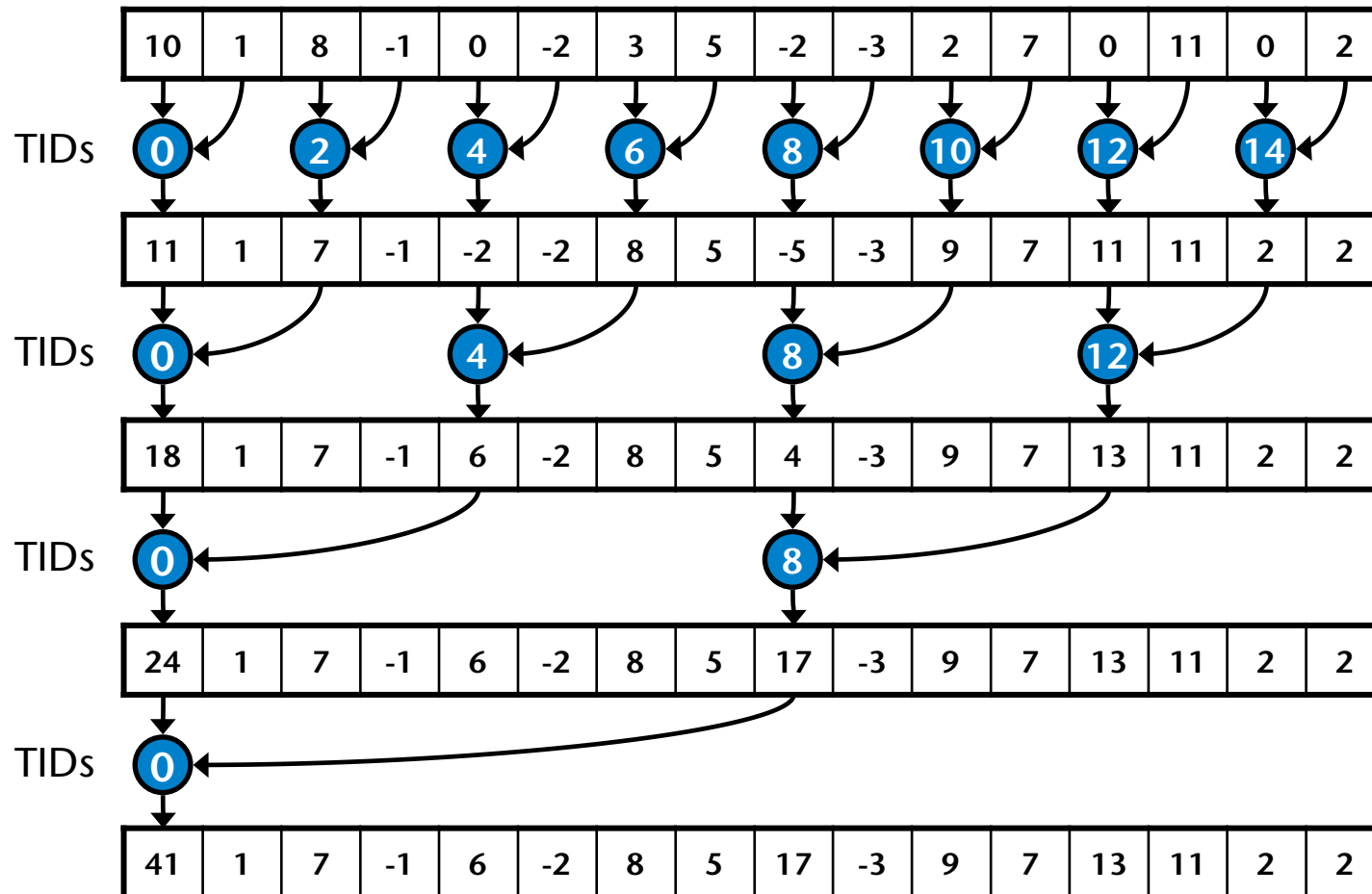
- This similarity measure is called "**vector space model**"
 - One of the most frequently used similarity measures in IR
- Note: our definition is a slightly simplified version of the commonly used one (we omitted the tf-idf weighting)

- Why not the Euclidean distance $\|D_1 - D_2\|$?
 - Otherwise: documents D , and D concatenated to itself would be very *dissimilar!*



- Why do we need the normalization by $\frac{1}{\|D_1\| \|D_2\|}$?
 - Same reason ...

- Why didn't we do the reduction this way?

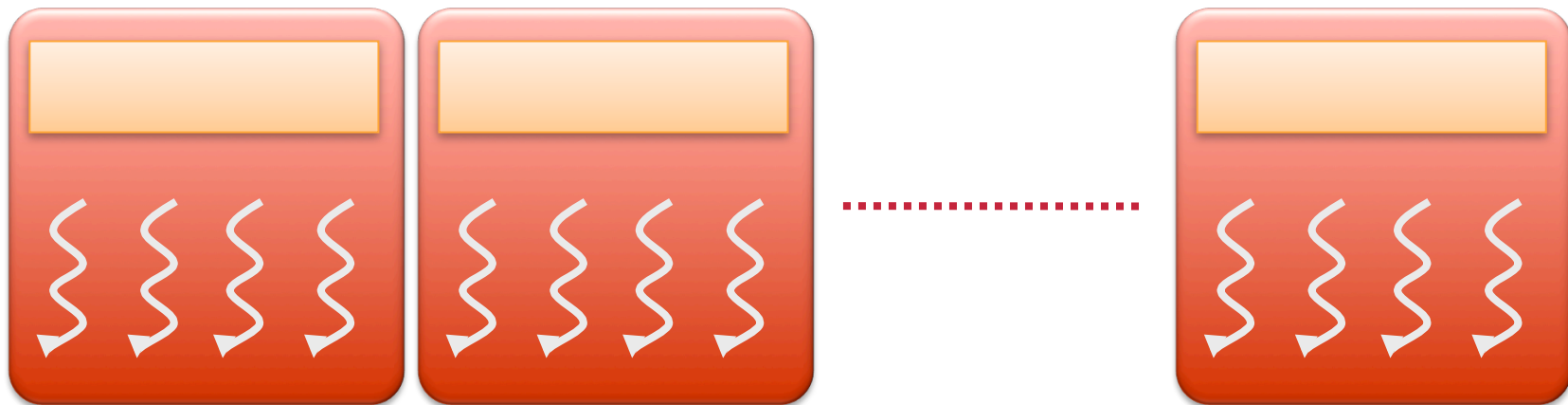


- The kernel for this algorithm:

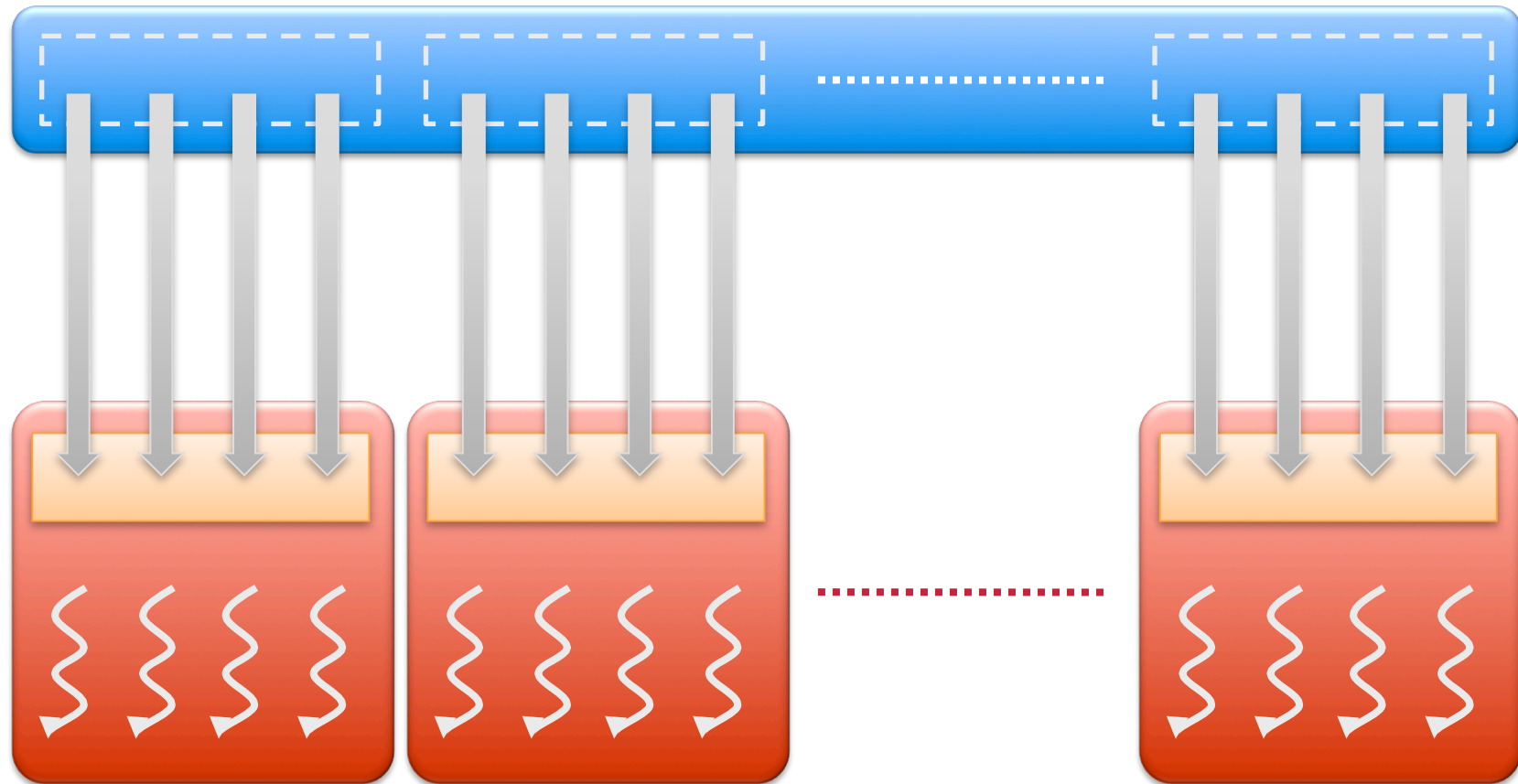
```
// do reduction in shared mem
__syncthreads();
for ( int i = 1; i < blockDim.x; i *= 2 )
{
    if ( threadIdx.x % (2*i) == 0 )
        cache[threadIdx.x] += cache[threadIdx.x + i];
    __syncthreads();
}
```

Problem:
highly
divergent
warps are
very inefficient

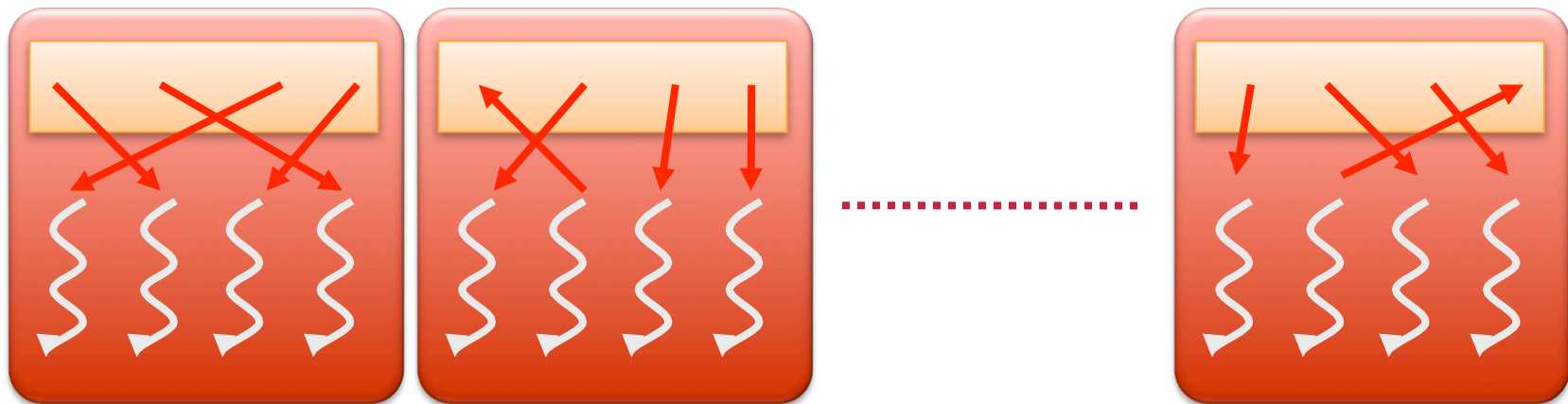
A Common, Massively Parallel Programming Pattern



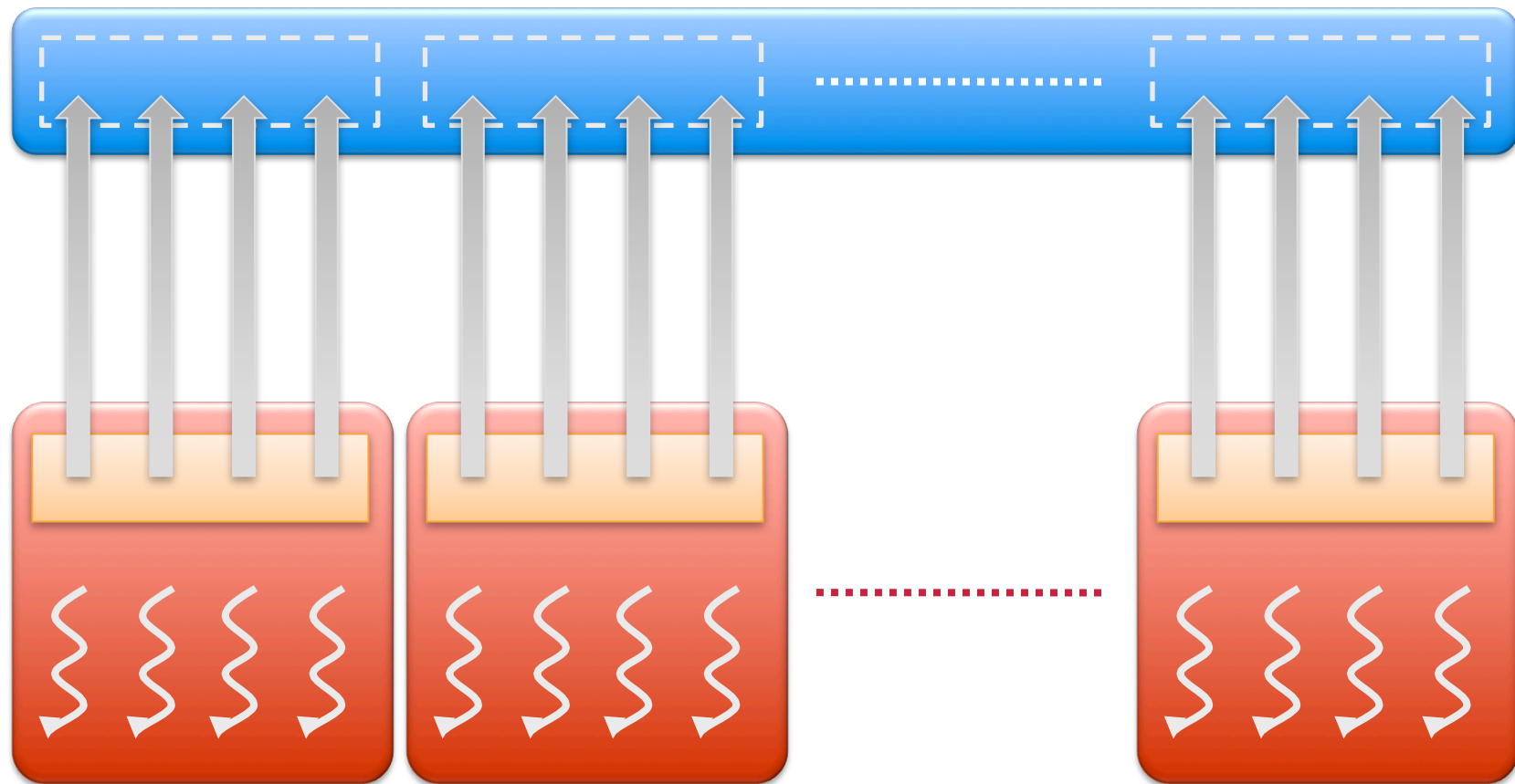
- Partition your domain such that each subset fits into shared memory; handle each data subset with one thread block



- Load the subset from global memory to **shared memory**; exploit memory-level parallelism by loading one piece per thread; don't forget to **synchronize** all threads before continuing!



- Perform the computation on the subset in **shared memory**



- Copy the result from **shared memory** back to global memory



Remarks on Memory (Applies to GPUs and CPUs)



- In our dot product kernel, we could have done everything in global memory, but ...
- Global memory bandwidth is loooow:

Ideal



Reality



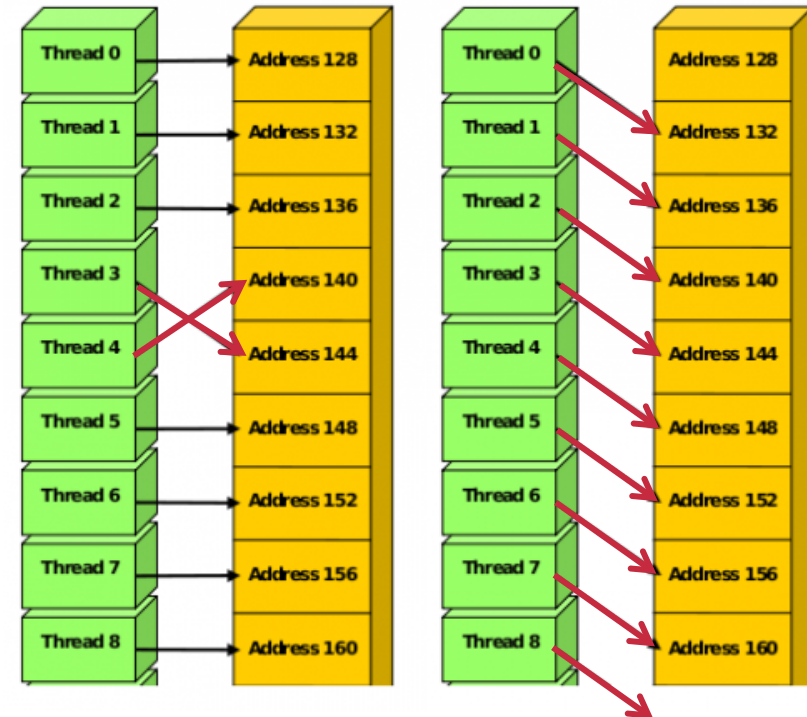
Coalesced Memory Access

- One of the most important optimization techniques for massively parallel algorithm design (on GPUs and – to some degree – CPUs!)

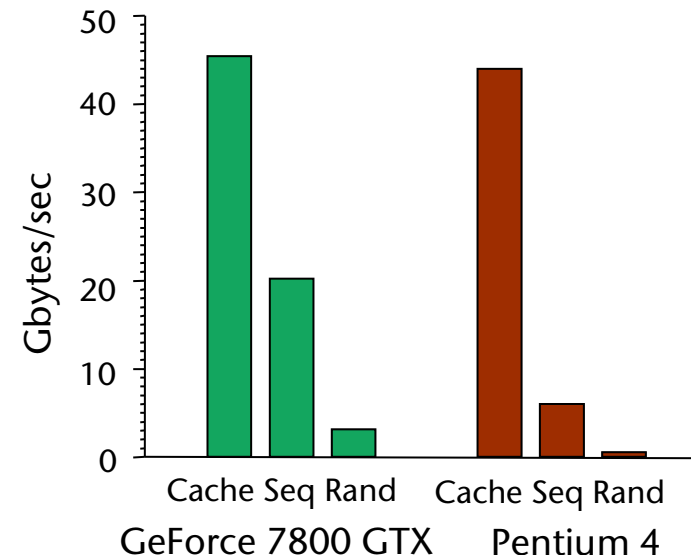
Coalesced memory accesses



Uncoalesced memory accesses

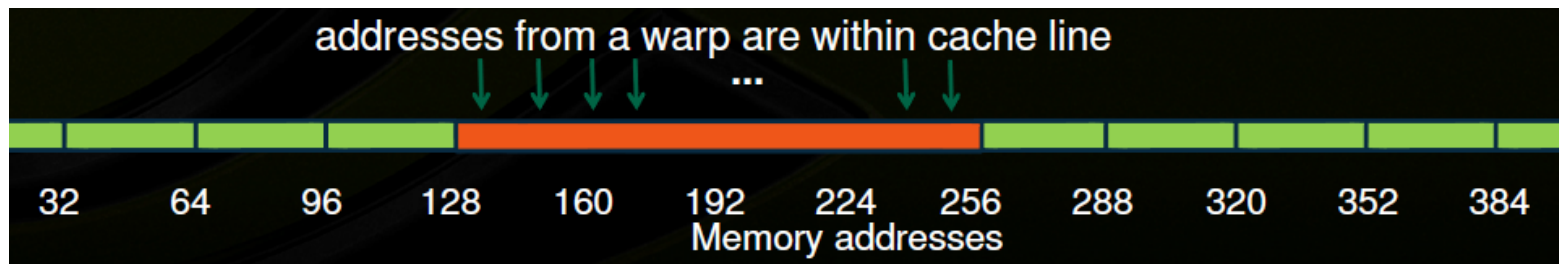


- When does the GPU win over the CPU?
- **Arithmetic intensity** of an algorithm := $\frac{\text{number of arithmetic operations}}{\text{amount of transferred bytes}}$
 - Sometimes also called **computational intensity**
- Unfortunately, many (most?) algorithms have a low arithmetic intensity → they are **bandwidth limited**
- GPU wins if memory access is "streamed" = coalesced
 - Hence, "stream programming architecture"



How to Achieve Coalesced Access

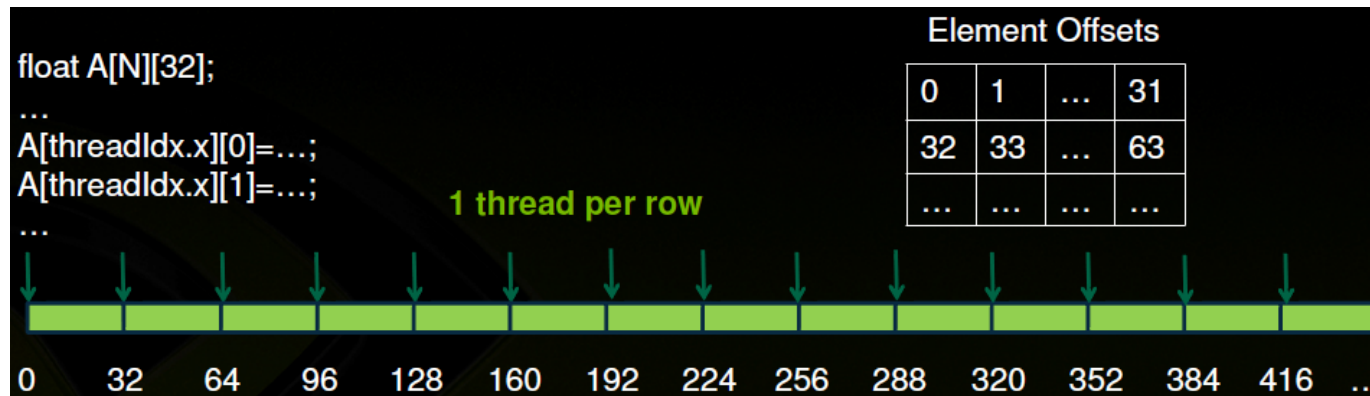
- Addresses from a warp (“thread-vector”) are converted into line requests
 - line sizes: 32B and 128B
 - Goal is to maximally utilize the bytes in these lines



2D Array Access Pattern (row major)

- Consider the following piece in a kernel (e.g., matrix \times vector):

```
for ( int j = 0; j < 32; j ++ ) {
    float x = A[threadIdx.x][j];
    ... do something with it ...
}
```



➤ Uncoalesced access pattern:

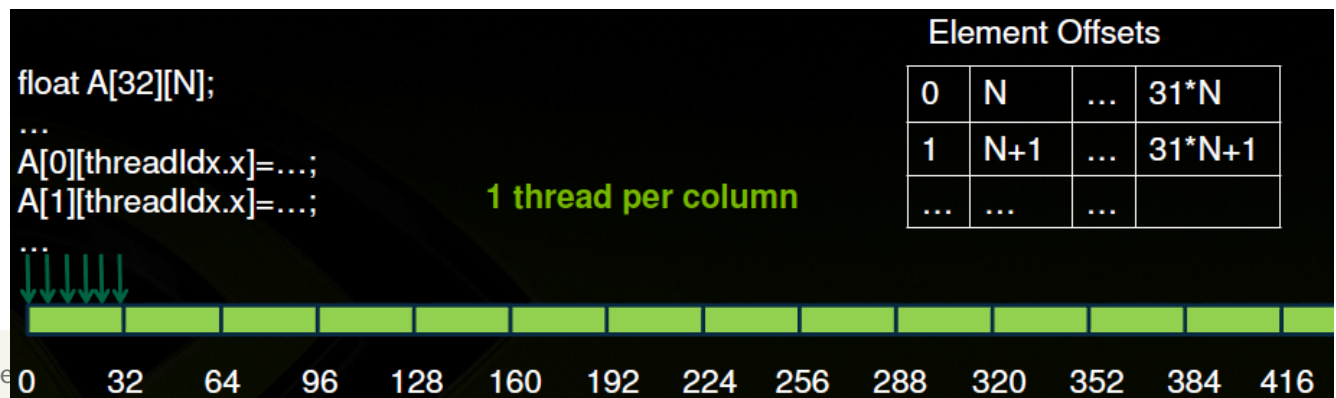
- Elements read on 1st SIMT access: 0, 32, 64, ...
- Elements read on 2nd SIMT access: 1, 33, 65, ...
- Also, extra data will be transferred in order to fill the cache line size
- Generally, most natural access pattern for direct port of a C/C++ code!

Transposed 2D Array Access Pattern

- The "natural" way to store matrices is called **row major order**
- **Column major** := store a logical row in a physical column
 - I.e., $A_{00} \rightarrow A[0][0]$, $A_{01} \rightarrow A[1][0]$, $A_{02} \rightarrow A[2][0]$, ...
 $A_{10} \rightarrow A[0][1]$, $A_{11} \rightarrow A[1][1]$, $A_{12} \rightarrow A[2][1]$, ...
 $A_{20} \rightarrow A[0][2]$, ...
- Transform the code piece (e.g., row×column) to column major:

```
for ( int j = 0; j < 32; j ++ ){
    float x = A[j][treadIdx.x];
    ... do something with it ...
}
```

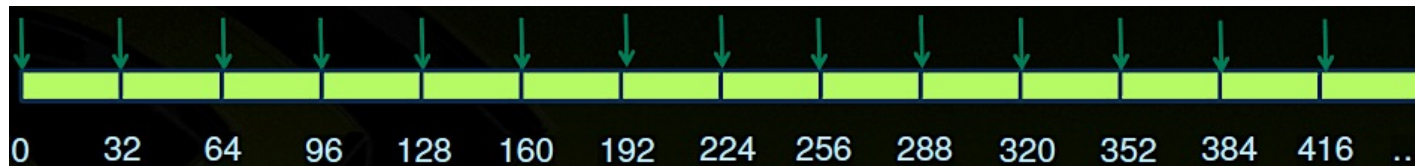
- Now, we have coalesced accesses:
 - Elements read on 1st SIMT access: 0, 1, 2, ..., 31
 - Elements read on 2nd SIMT access: 32, 33, ..., 63



Array of Structure or Structure of Array?

- An **array of structures (AoS)** behaves like row major accesses:

```
struct Point {
    float x; float y; float z;
};
Point PointList[N];
...
PointList[threadIdx.x].x = ...
```

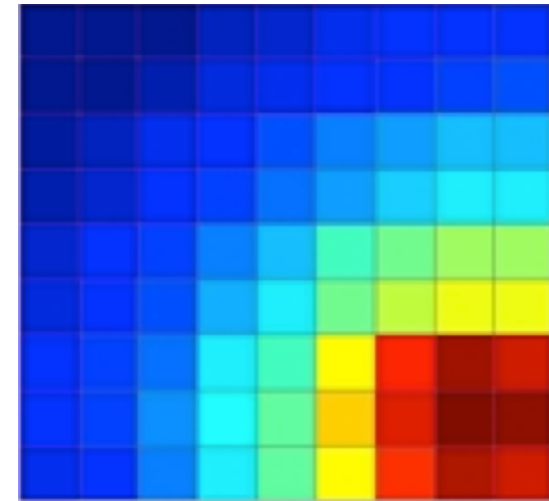


- A **structure of arrays (SoA)** behaves like column major access:

```
struct PointList {
    float x[N];
    float y[N];
    float z[N];
};
...
PointList.x[threadIdx.x] = ...
```



- Assumptions:
 - For sake of illustration, our domain is 2D
 - Discrete domain → 2D grid
(common approach in simulation)
 - A few designated cells are "heat sources"
→ cells with constant temperature
- Simulation model (simplistic):



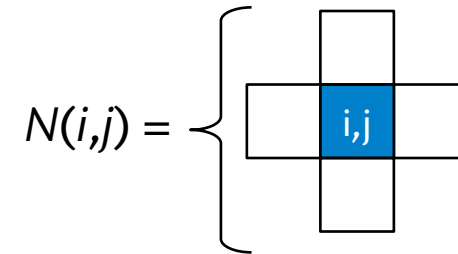
$$T_{i,j}^{n+1} = T_{i,j}^n + \sum_{(k,l) \in N(i,j)} \mu (T_{k,l}^n - T_{i,j}^n)$$

$$\Leftrightarrow T_{i,j}^{n+1} = (1 - N\mu) T_{i,j}^n + \mu \sum_{(k,l) \in N(i,j)} T_{k,l}^n \quad (1)$$

N = number of cells in the neighborhood

- Iterate this (e.g., until convergence to steady-state)

- Do we achieve energy conservation?
- For sake of simplicity, assume

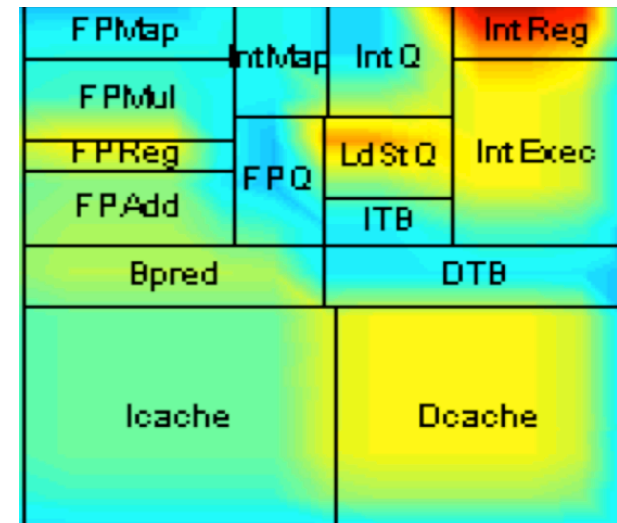
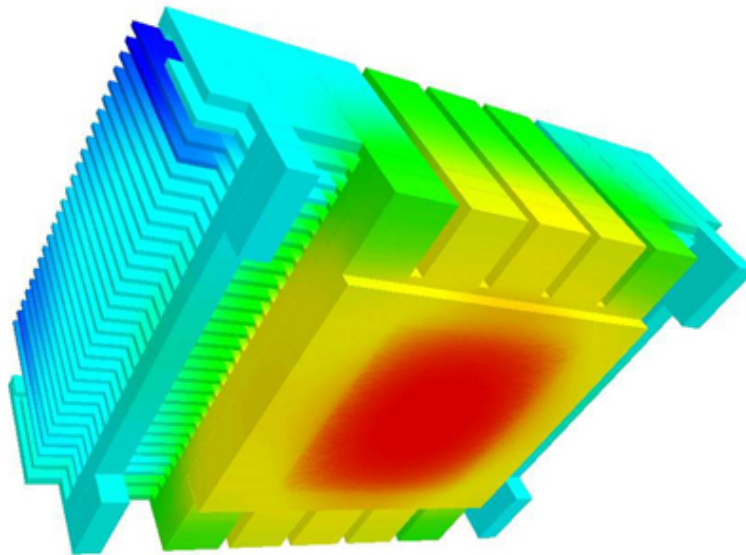


- Energy consumption iff $\sum_{i,j} T_{i,j}^{n+1} \stackrel{!}{=} \sum_{i,j} T_{i,j}^n$
- Plugging (1) into (2) yields

$$\underbrace{(1 - N\mu) \sum_{i,j} T_{i,j}^n + \mu \sum_{i,j} \sum_{(k,l) \in N(i,j)} T_{k,l}^n}_{=0} \stackrel{!}{=} \sum_{i,j} T_{i,j}^n \tag{2}$$

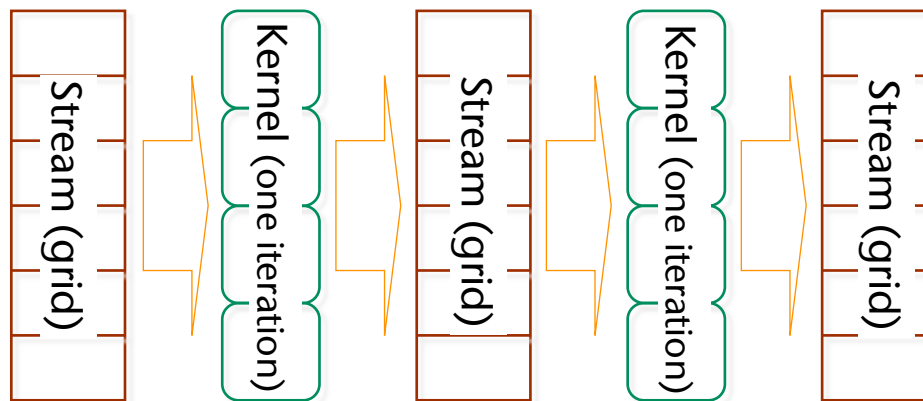
- Therefore, μ is indeed a free material parameter (= "heat flow speed")

- Example: heat simulation of ICs and cooling elements

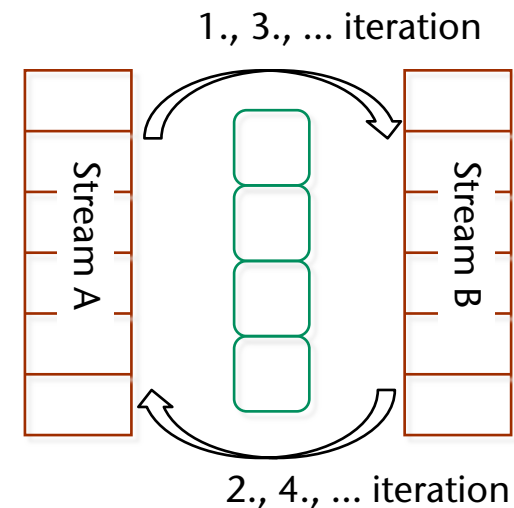


Pattern: Double Buffering

- Observations:
 - Each cell's next state can be computed completely independently
- We can arrange our computations like this:



- General parallel programming pattern: **double buffering** ("ping pong")



Algorithm

- One thread per cell

1. Kernel for re-setting heat sources:

```
if ( cell is heat cell ) :  
    read temperature from constant "heating stencil"
```

2. Kernel for one transfer step:

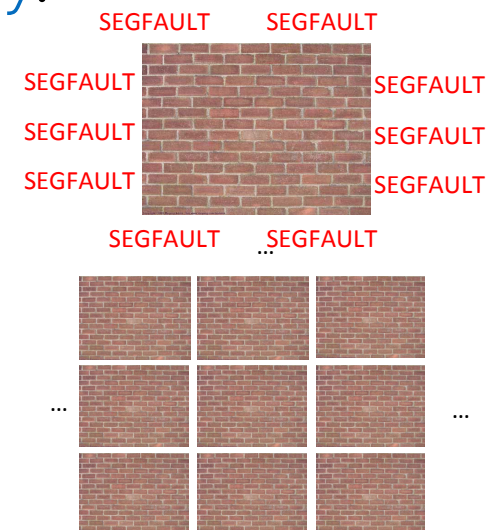
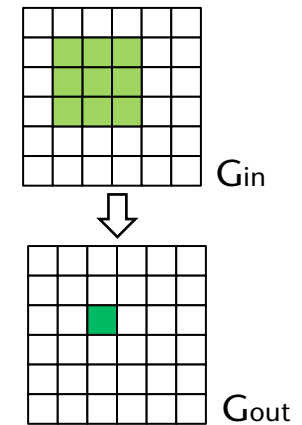
```
Read all neighbor cells  input_grid[tid.x+-1][tid.y+-1]  
Accumulate them  
Write new temperature in output_grid[tid.x][tid.y]
```

3. Swap pointers to input & output grid (done on host)

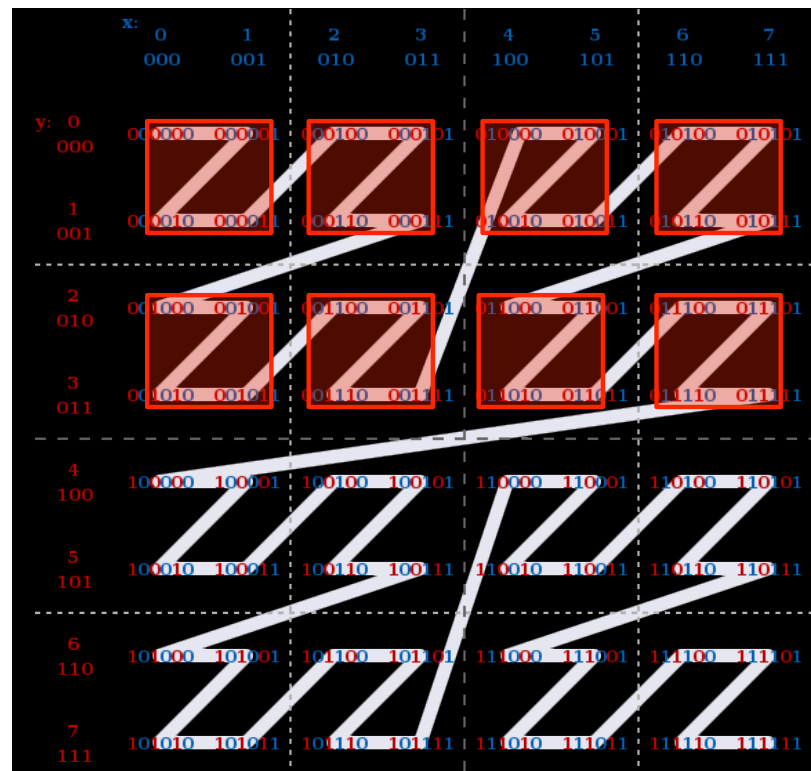
- Challenge: border cells! (very frequent problem in sim. codes)
 - Use if-then-else in above kernel?
 - Use extra kernel that is run only for border cells?
 - Introduce padding around domain? Arrange domain as torus?

- Many computations have the following characteristics:
 - They *iterate* a simple function many times
 - They work on a 2D/3D *grid*
 - We can run *one thread per grid cell*
 - Each thread only needs to look at *neighbor cells*
 - Each iteration transforms an input grid into an output grid

- For this kind of algorithms, there is **texture memory**:
 - Special cache with optimization for **spatial locality**
 - **Access to neighbor cells is very fast**
 - Important: can handle out-of-border accesses automatically by clamping or wrap-around!
 - For the technical details: see "Cuda by Example",
Nvidia's "CUDA C Programming Guide",

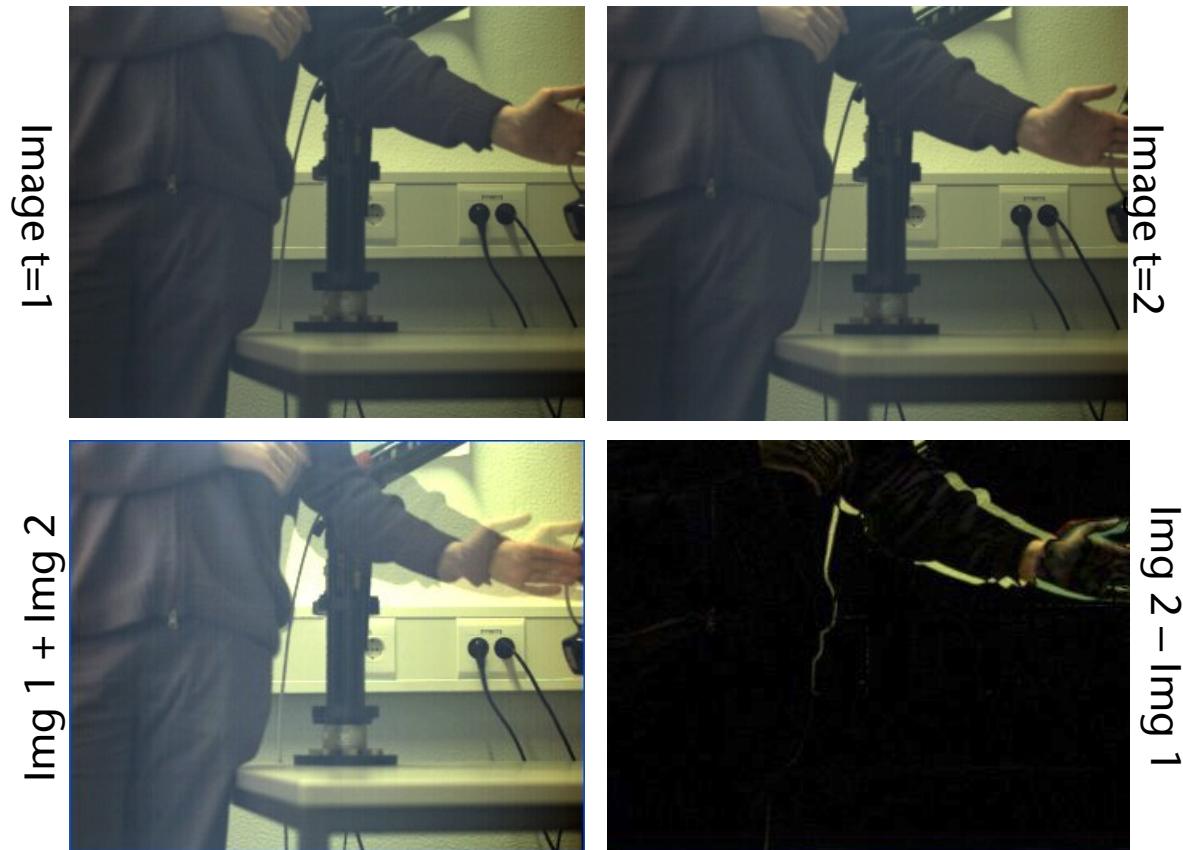


- The locality-preserving cache is probably achieved by arranging data via a **space-filling curve**:



Other Applications of Texture Memory

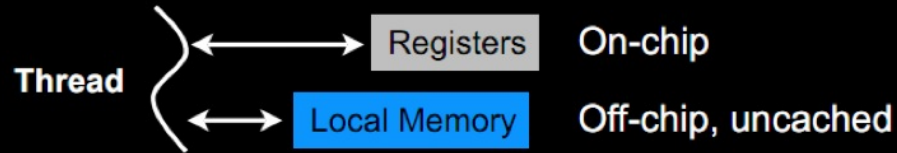
- Most image processing algorithms exhibit this kind of locality
- Trivial example: image addition / subtraction → neighboring threads access neighboring pixels



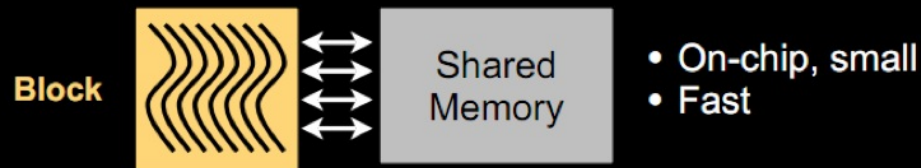
Kernel Memory A

Review

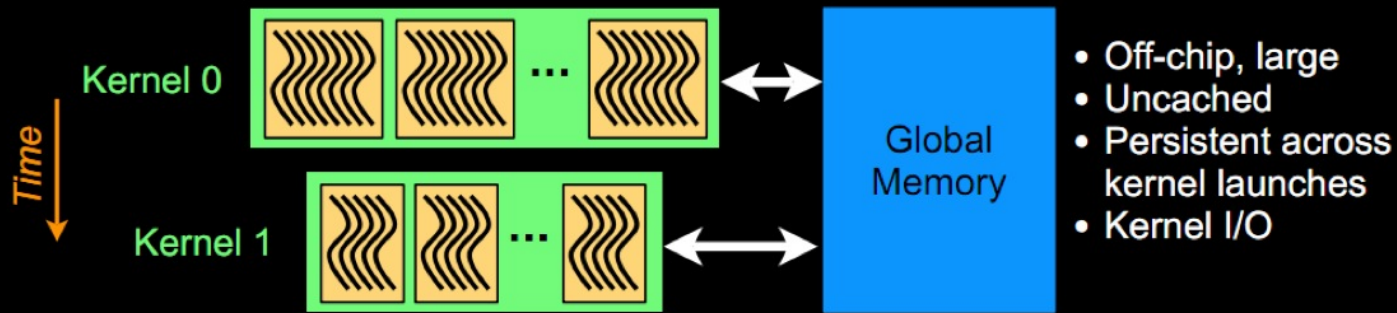
● Per-thread

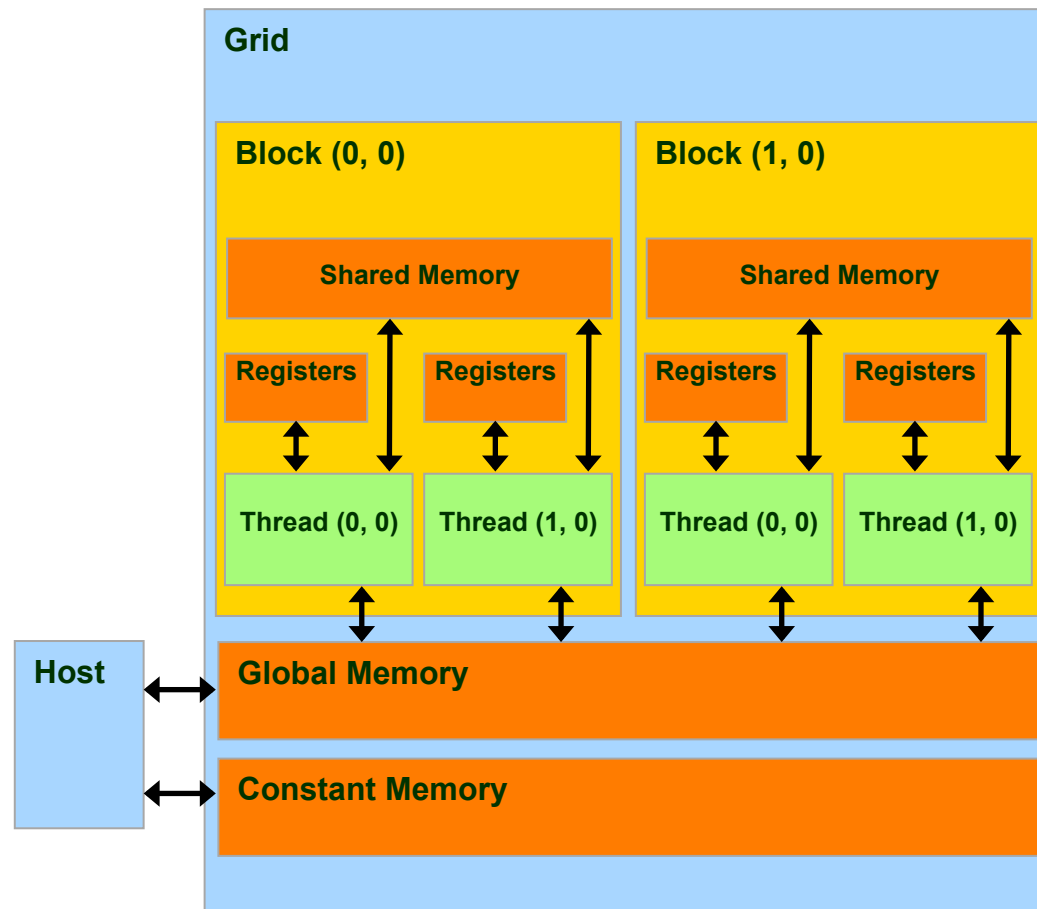


● Per-block



● Per-device





Variable declaration		Memory	Access	Lifetime
<code>__device__ __local__</code>	<code>int LocalVar;</code>	local	thread	thread
<code>__device__ __shared__</code>	<code>int SharedVar;</code>	shared	block	block
<code>__device__</code>	<code>int GlobalVar;</code>	global	grid	application
<code>__device__ __constant__</code>	<code>int ConstantVar;</code>	constant	grid	application

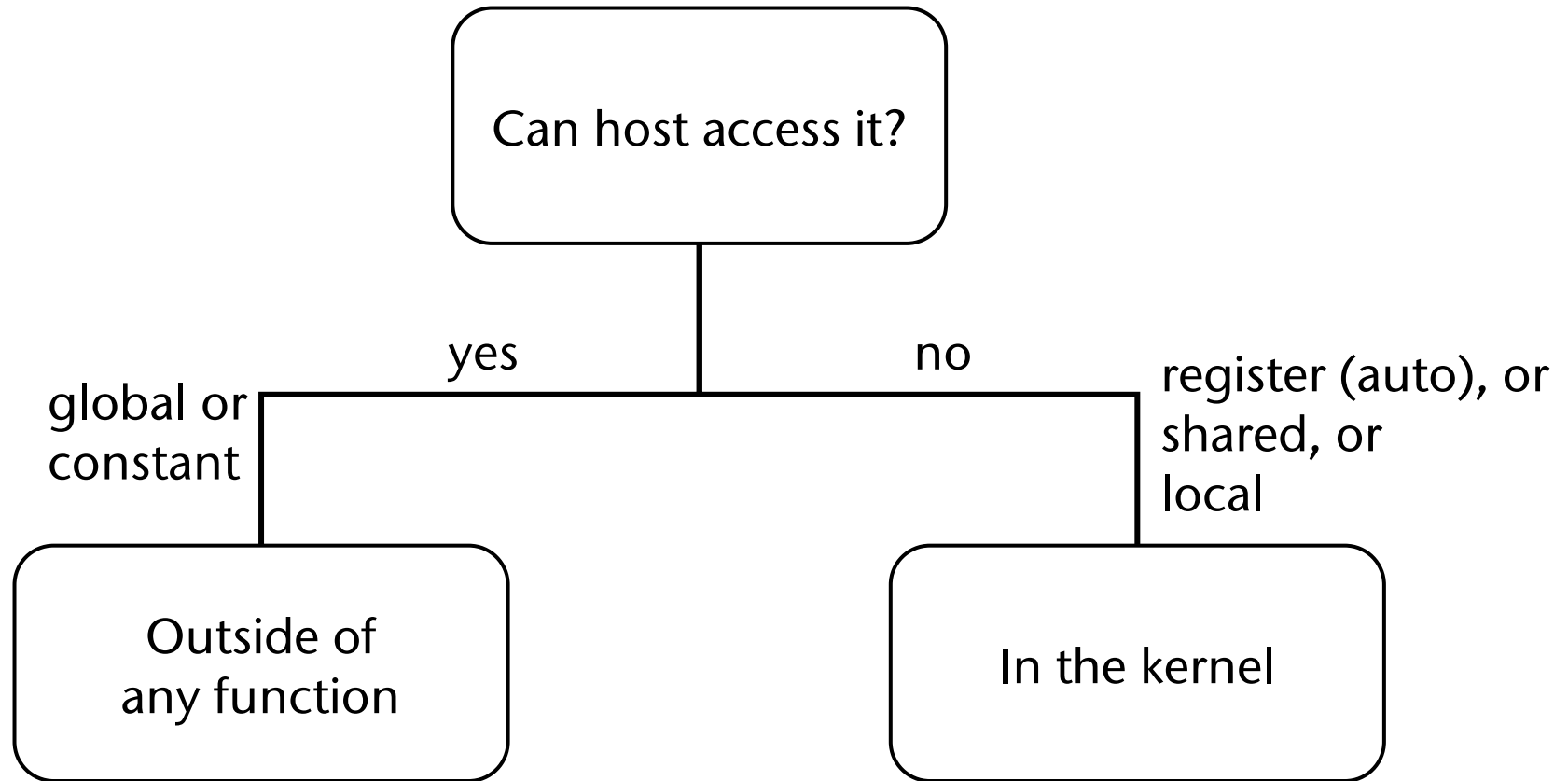
- Remarks:

- `__device__` is optional when used with `__local__`, `__shared__`, or `__constant__`
- **Automatic variables** without any qualifier reside in a **register**
 - Except arrays, which reside in local memory (slow)

	Variable declaration	Memory	Penalty
	<code>int var;</code>	register	1x
	<code>int array_var[10];</code>	local	100x
<code>__shared__</code>	<code>int shared_var;</code>	shared	1x
<code>__device__</code>	<code>int global_var;</code>	global	100x
<code>__constant__</code>	<code>int constant_var;</code>	constant	1x

- scalar variables reside in fast, on-chip registers
- shared variables reside in fast, on-chip memories
- thread-local arrays & global variables reside in uncached off-chip memory
- constant variables reside in cached off-chip memory

Where to Declare Variables?



Massively Parallel Histogramm Computation

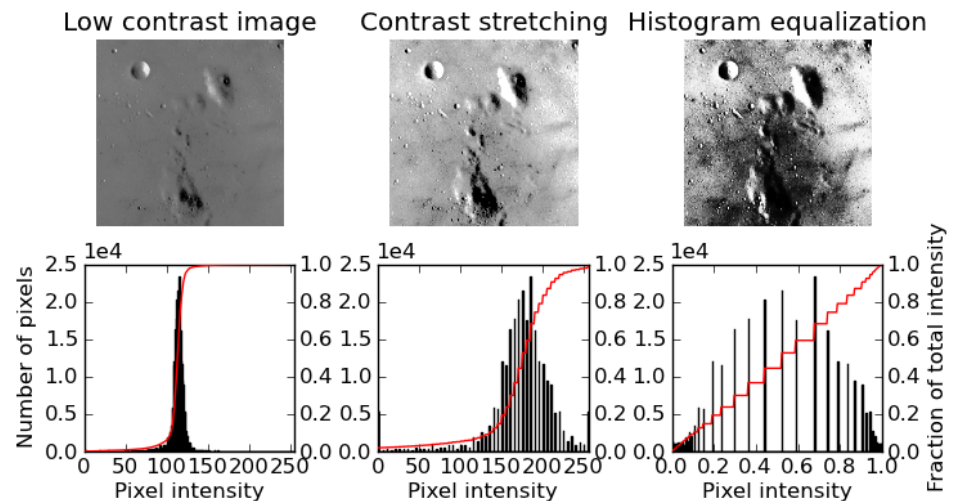
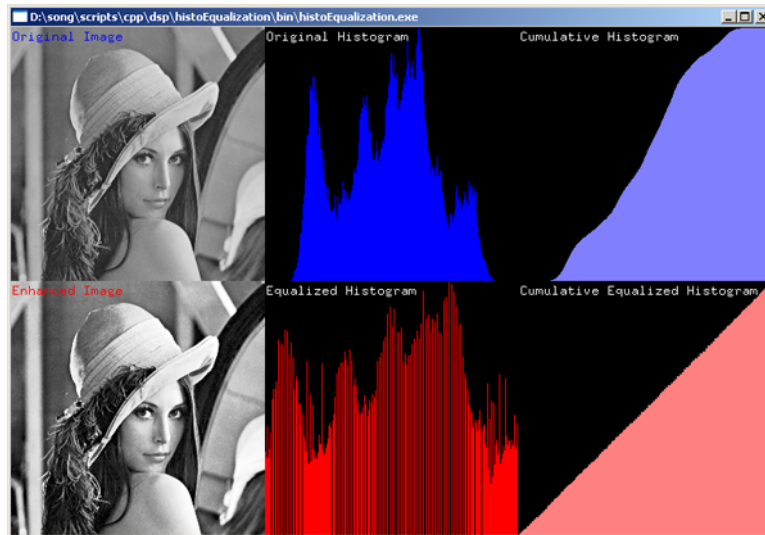
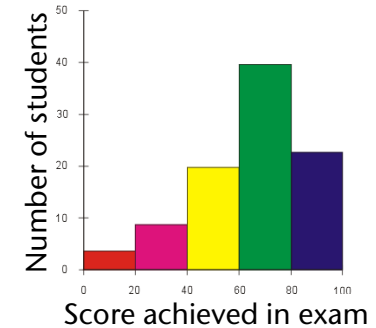
- Definition (for images):

$$h(x) = \# \text{ pixels with level } x$$

$$x \in 0, \dots, L - 1 \quad L = \# \text{ levels}$$

- Applications: many!

- Huffman Compression (see Info 2)
- Image Equalization (see Advanced Computer Graphics)



- The sequential algorithm:

```
unsigned char input[MAX_INP_SIZE]; // e.g. image
int input_size; // # valid chars in input
unsigned int histogram[256]; // 256 ASCII chars

// clear histogram
for (int i = 0; i < 256; i ++ )
    histogram[i] = 0;
for (int i = 0; i < input_size; i ++ )
    histogram[ input[i] ] ++ ; // real histogram comput.

// verify histogram
long int total_count = 0;
for (int i = 0; i < 256; i ++ )
    total_count += histogram[i];
if ( total_count != input_size )
    fprintf(stderr, "Error! ..." );
```


- Naïve "massively parallel" algorithm:
 - One thread per bin (e.g., 256)
 - Each thread scans the complete input and counts the number of occurrences of its "own" character
 - At the end, each thread stores its character count in its histogram slot

- Disadvantage: not so massively parallel ...

- New approach: one thread per input character
- The setup on the host side:

```
set up device arrays d_input, d_histogram
cudaMemset( d_histogram, 0, 256 * sizeof(int) );
int threadsPerBlock = 256;
int nBlocks = #(multiprocessors on device) * 2;
computeHistogram <<< nBlocks, threadsPerBlock >>>
( d_input, input_size, d_histogram );
```

- Notes:
 - Letting **threadsPerBlock** = 256 makes things much easier in our case here
 - Letting **nBlocks** = (number of multiprocessors [SMXs] on the device) * 2 is a good rule of thumb, YMMV
 - On current hardware (Kepler) → ~ 16384 threads

- The kernel on the device side:

```
__global__ void
computeHistogram( unsigned char * input,
                  long int input_size,
                  unsigned int histogram[256] )
{
    int i = threadIdx.x + blockIdx.x * blockDim.x;
    int stride = blockDim.x * gridDim.x;
    while ( i < input_size )
    {
        histogram[ input[i] ] += 1;
        i += stride;
    }
}
```

- Problem: **race condition!!**

Solution: Atomic Operations

- The kernel with atomic add:

```
__global__ void
computeHistogram( unsigned char * input,
                  long int input_size,
                  unsigned int histogram[256] )
{
    int i = threadIdx.x + blockIdx.x * blockDim.x;
    int stride = blockDim.x * gridDim.x;
    while ( i < input_size )
    {
        atomicAdd( & histogram[input[i]], 1 );
        i += stride;
    }
}
```

- Prototype of atomicAdd():

```
T atomicAdd( T * address, T val )
```

- where **T** can be **int**, **float** (and few others)

- Semantic: *while atomicAdd performs its operation on address, no other thread can access this memory location! (neither read, nor write)*
- Problem: this algorithm is much slower than the sequential one!
 - Lesson: always measure performance against CPU!
- Cause: **congestion**
 - *Lots of threads waiting for a few memory locations to become available*



- Remedy: partial histograms in shared memory

```
computeHistogram( unsigned char * input,
                  long int input_size,
                  unsigned int histogram[256] )
{
    __shared__ unsigned int partial_histo[256];
    partial_histo[ threadIdx.x ] = 0;
    __syncthreads();

    int i = threadIdx.x + blockIdx.x * blockDim.x;
    int stride = blockDim.x * gridDim.x;
    while ( i < input_size ) {
        atomicAdd( & partial_histo[input[i]], 1 );
        i += stride;
    }
    __syncthreads();
    atomicAdd( & histogram[threadIdx.x],
              partial_histo[input[i]], 1 );
}
```

- Note: now it's obvious why we chose 256 threads/block

- All programming languages / libraries / environments providing for some kind of parallelism/concurrency have one or more of these atomic operations:
 - `int atomicExch(int* address, int val)`:
Read old value at address, store val in address, return old value
 - `int atomicMin(int* address, int val)`:
Read old value at address, compute minimum of old and val, store result in address, return old value
 - `int atomicAnd(int* address, int val);`
 - Atomic add
 - And atomic compare-and-swap ...

- The fundamental atomic operation "**Compare And Swap**":
 - In CUDA: `int atomicCAS(int* address, int compare, int val)`
 - Performs this little algorithm atomically:

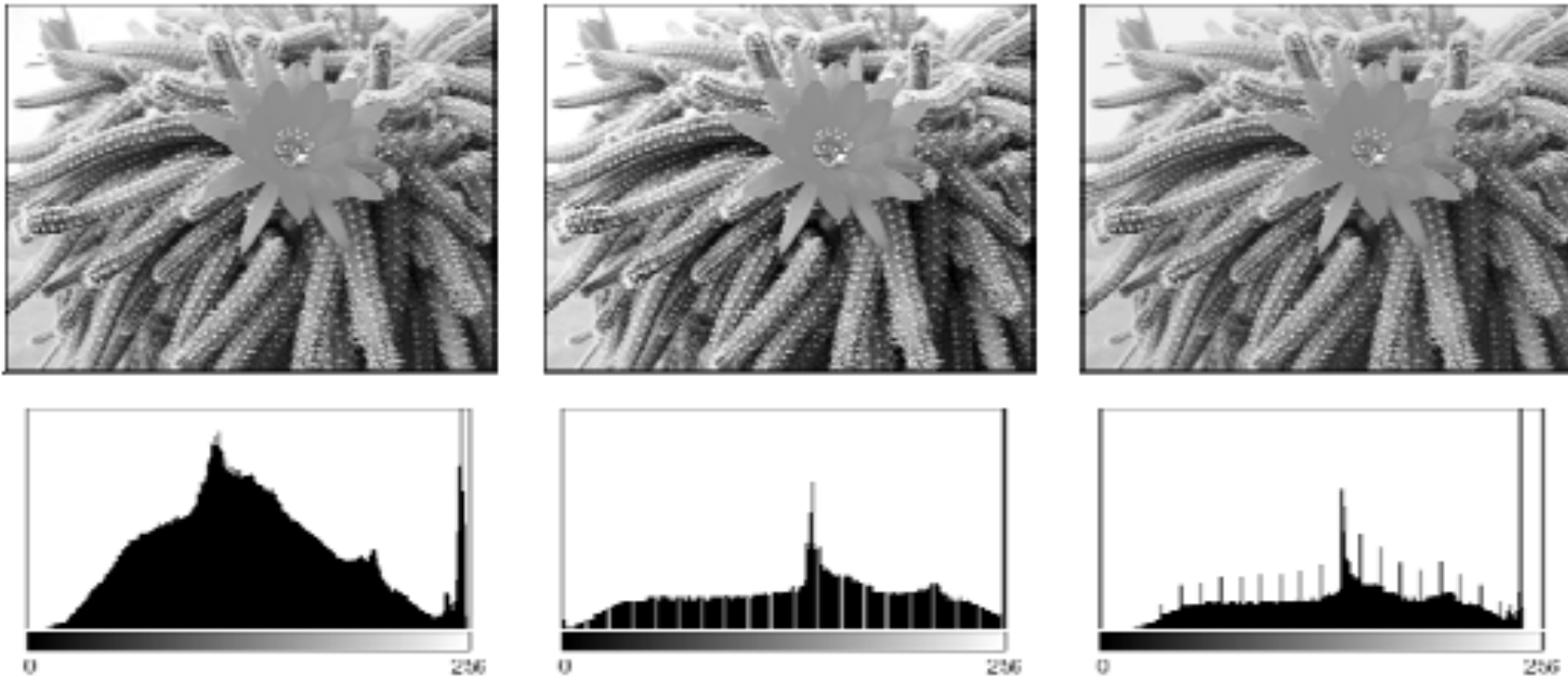
```
atomic_compare_and_swap( address, compare, new_val ) :  
    old ← value in memory location address  
    if compare == old:  
        store new_val → memory location address  
    return old
```

- Theorem (w/o proof):
All other atomic operations can be implemented using atomic compare-and-swap.

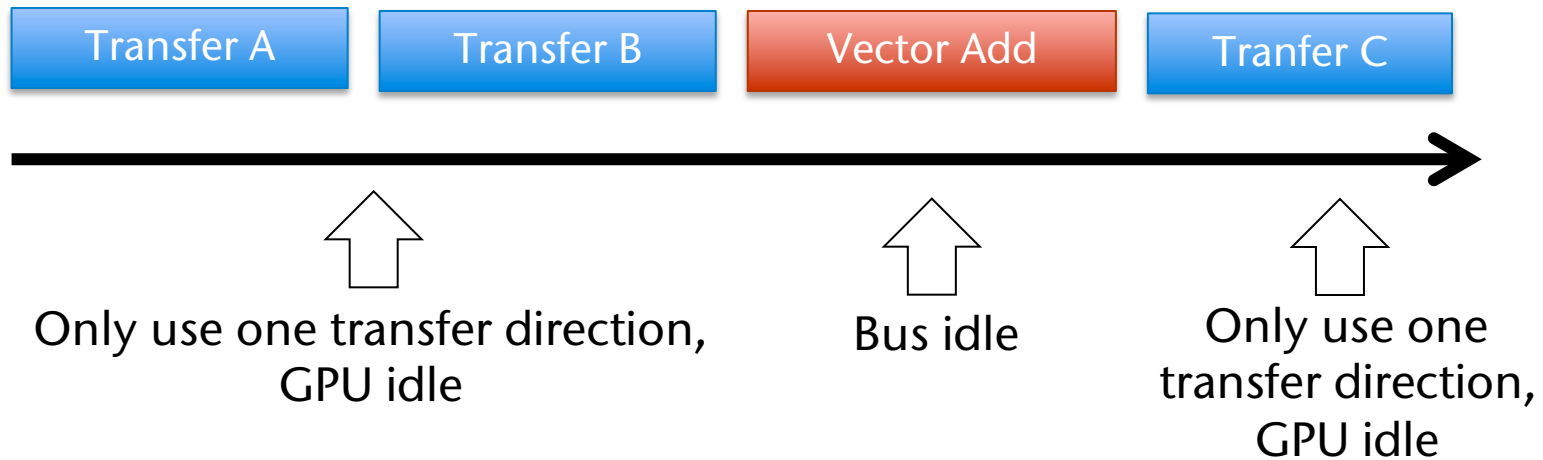
- Example:

```
atomic_add( address, incr ):  
  current_val := value in memory location address  
  repeat  
    new_val      := current_val + incr  
    assumed_val := current_val  
    current_val := compare_and_swap( address,  
                                     assumed_val,  
                                     new_val )  
  until assumed_val == current_val
```

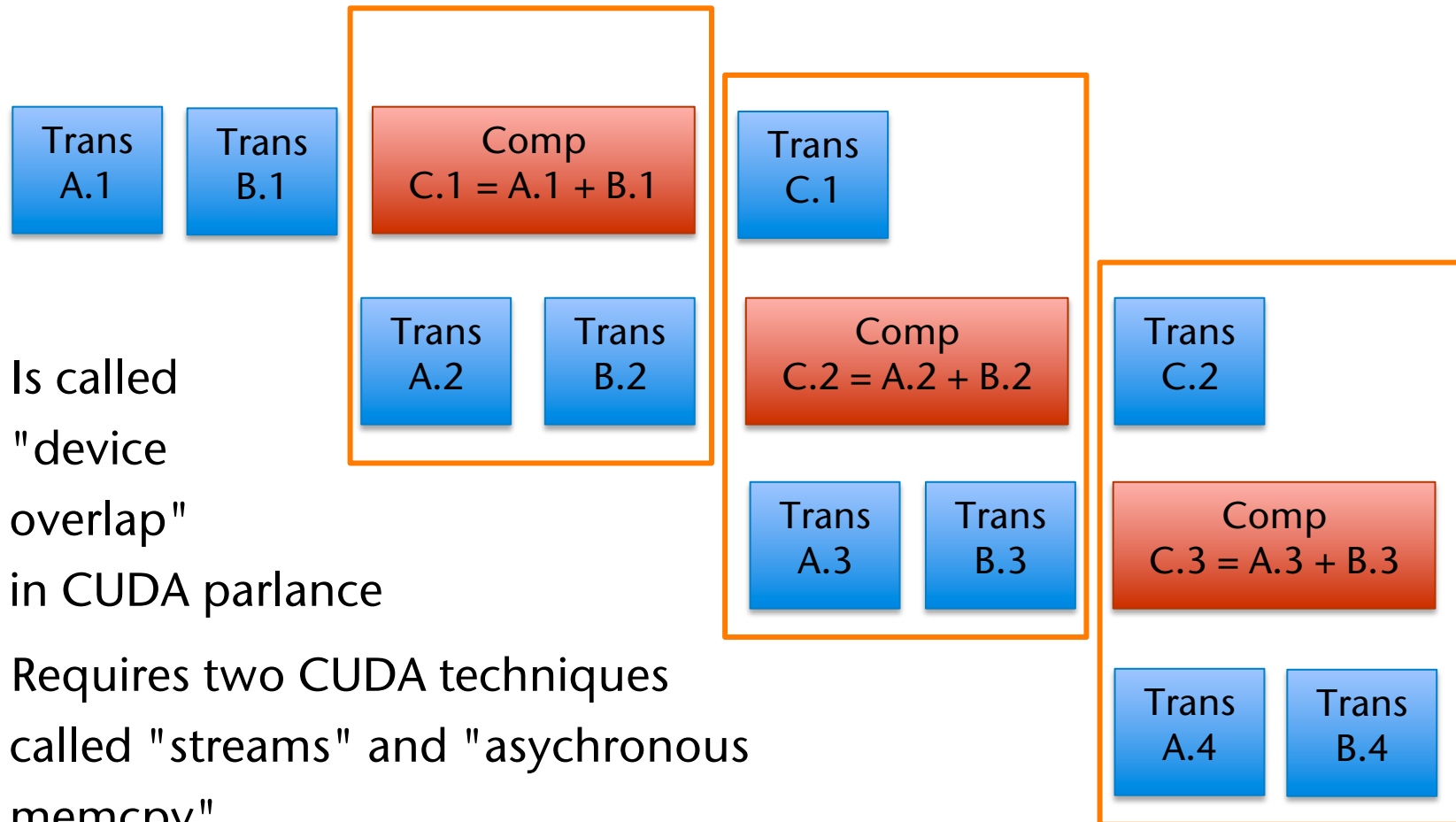
Image Restoration Using Histograms



- Problem with performance, if lots of transfer between GPU↔CPU:



- Solution: **pipelining** (the "other" *parallelism paradigm*)



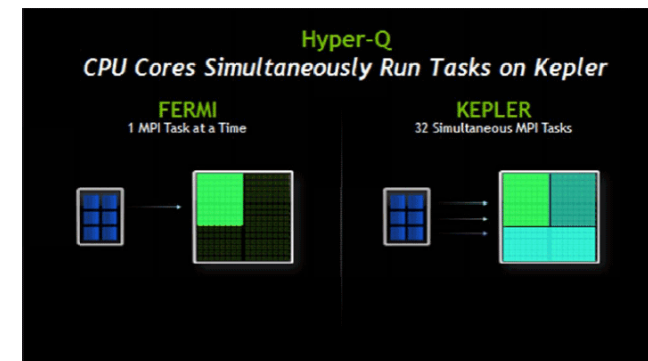
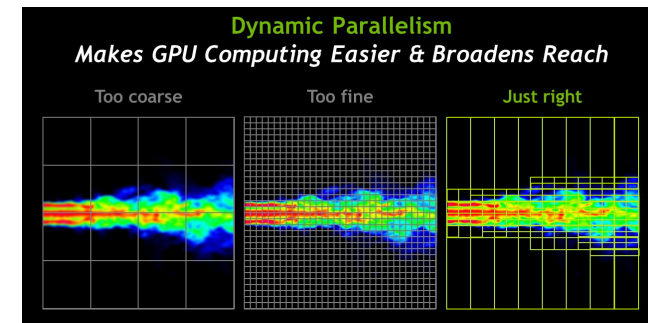
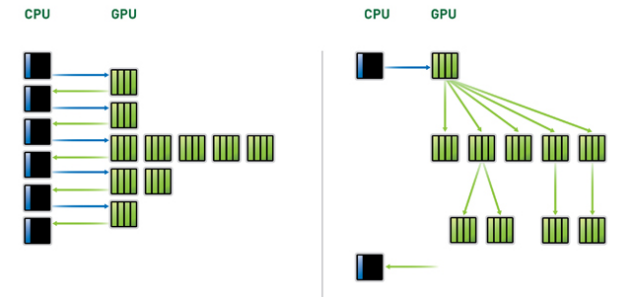
- Is called "device overlap" in CUDA parlance
- Requires two CUDA techniques called "streams" and "asynchronous memcpy"

For More Information on CUDA ...

- *CUDA C Programming Guide* (zur Programmiersprache)
- *CUDA C Best Practices Guide* (zur Performance-Steigerung)
- */Developer/NVIDIA/CUDA-5.0/doc/html/index.html*
(zum Runtime API)

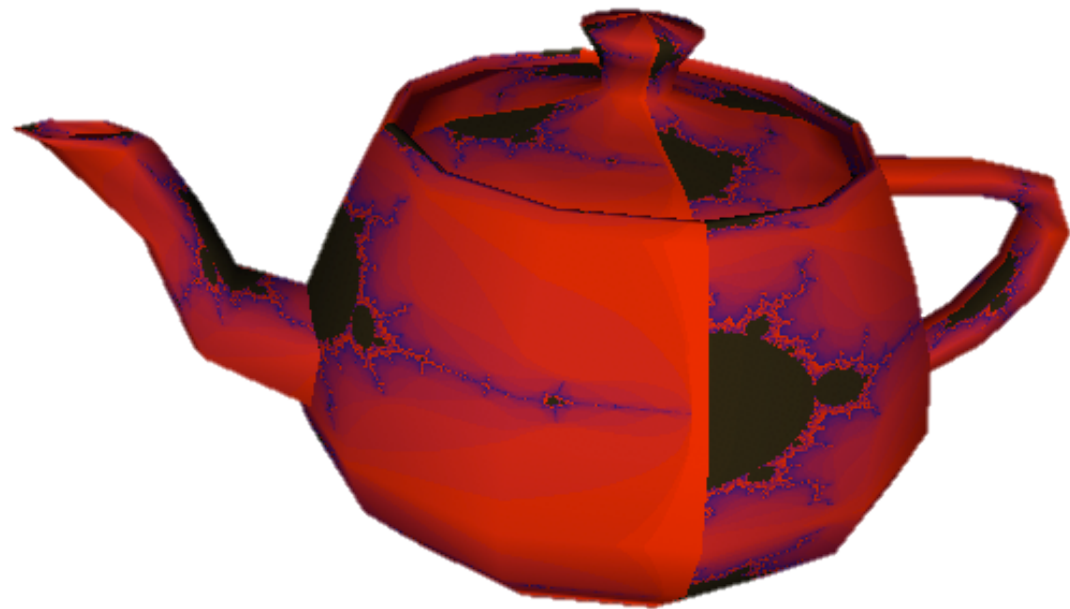
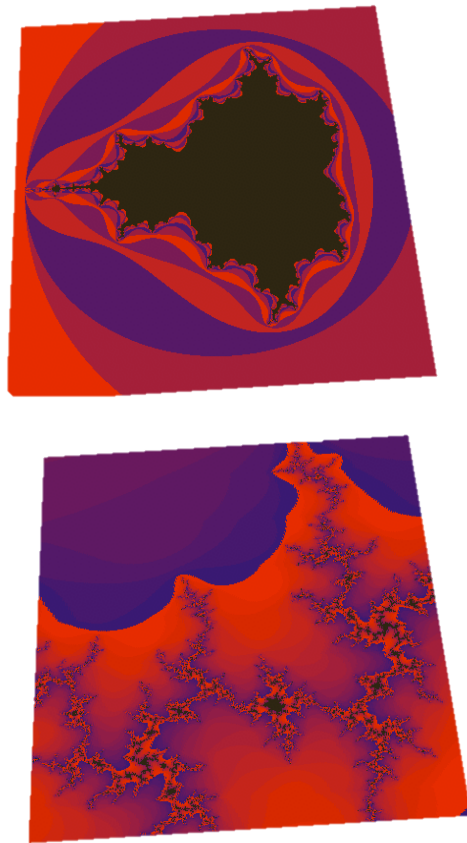
Concepts we Have Not Covered Here

- Dynamic parallelism (threads can launch new threads)
 - Good for irregular data parallelism (e.g., tree traversal, multi-grids)
- Running several tasks at the same time on a GPU (via MPI; they call it "Hyper-Q")
- See:
 - "Introduction to CUDA 5.0" on the course web page
 - "CUDA C Programming Guide" at docs.nvidia.com/cuda/index.html



- Graphics Interoperability:
 - Transfer images directly from CUDA memory to OpenGL's framebuffer
- Dynamic shared memory
- Asynchronous memory copies between host ↔ device
- Dynamic memory allocation in the kernel
 - Can have serious performance issues
- Pinned CPU memory (
- CUDA Streams
- Multi-GPU programming, GPU-to-GPU memory transfer
- Zero-copy data transfer
- Libraries: CUBLAS, Thrust, ...
- Voting functions (`__all()`, `__any()`)

- With Graphics Interoperability, you can render results from CUDA directly in a 3D scene, e.g. by using them as textures



Demo in `demos/shader/mandelbrot`

