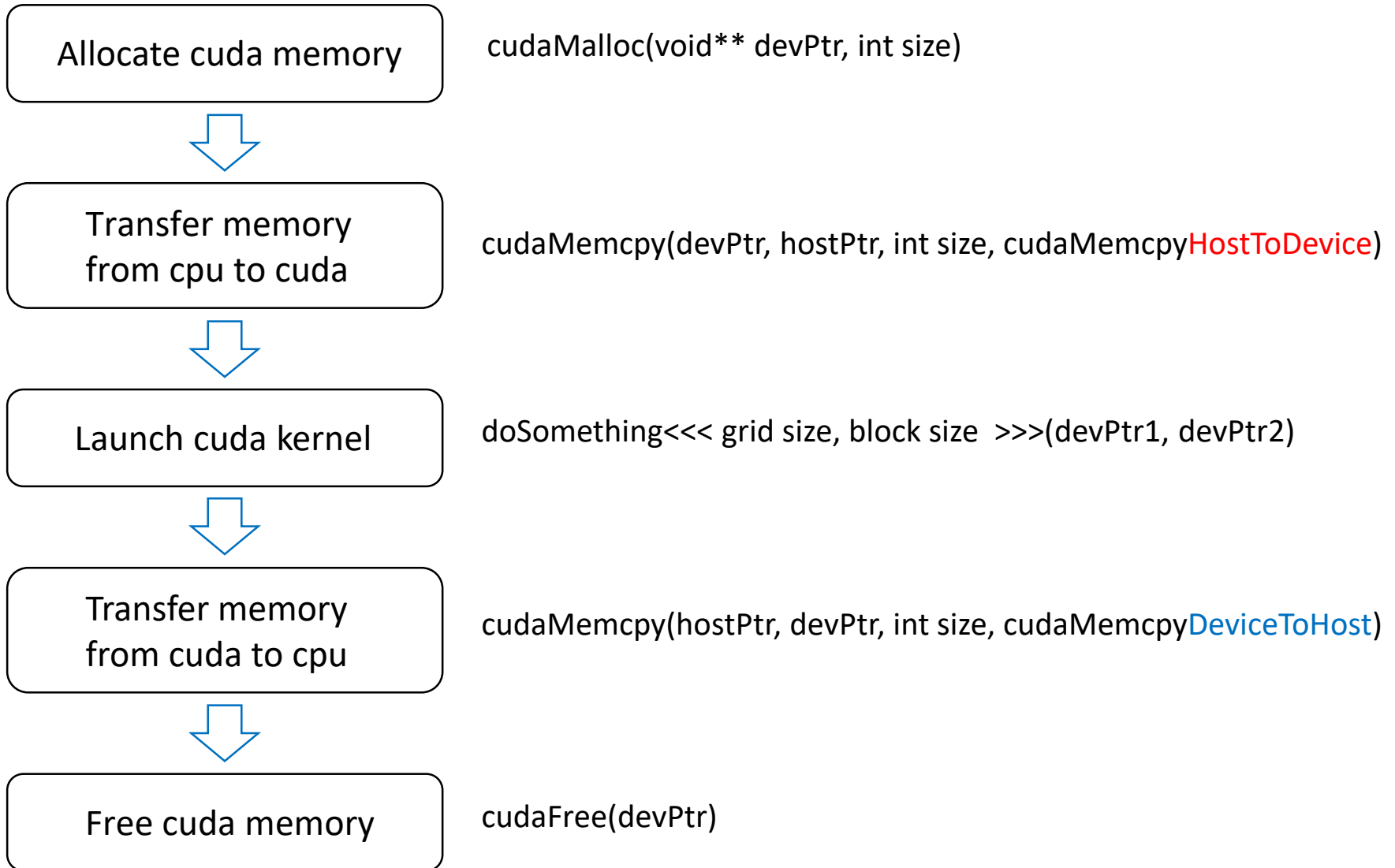


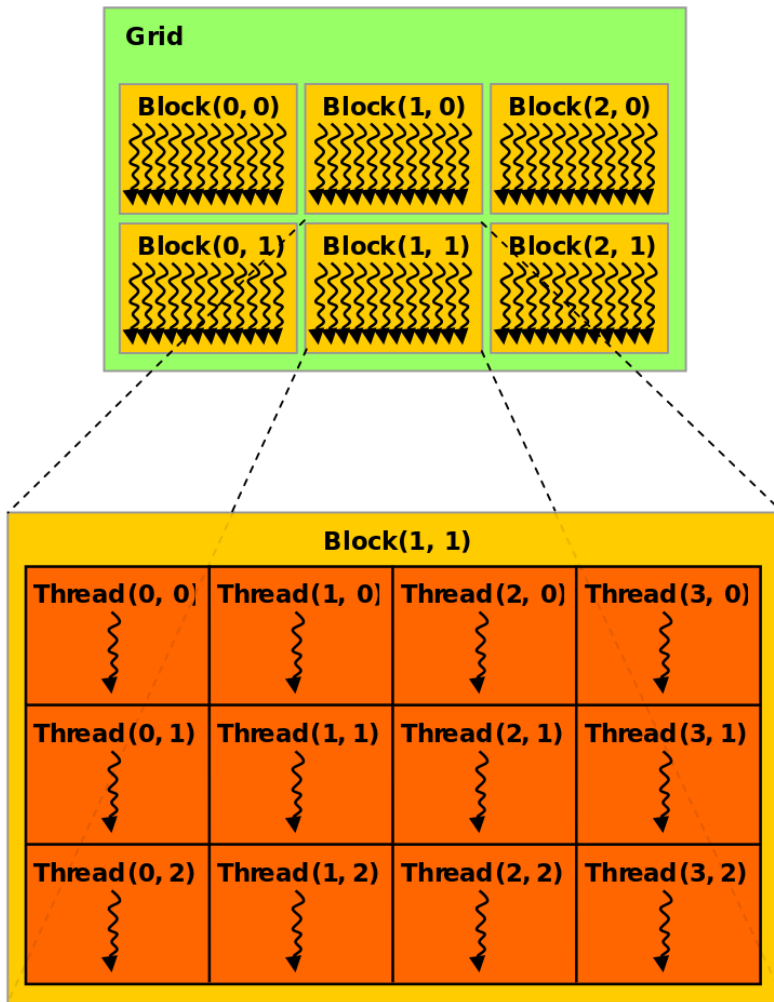
CUDA acceleration for ACTS seed finding

Beomki yeo (KAIST)

General flow of CUDA computing



CUDA kernel



- A kernel is launched with given size of grid and block

`doSomething<<< grid size, block size >>>(devPtr1, devPtr2)`

- threads in different blocks can not communicate each other, in principle.
→ Independent tasks are split into each block
- User can refer the thread ID and block ID inside kernel

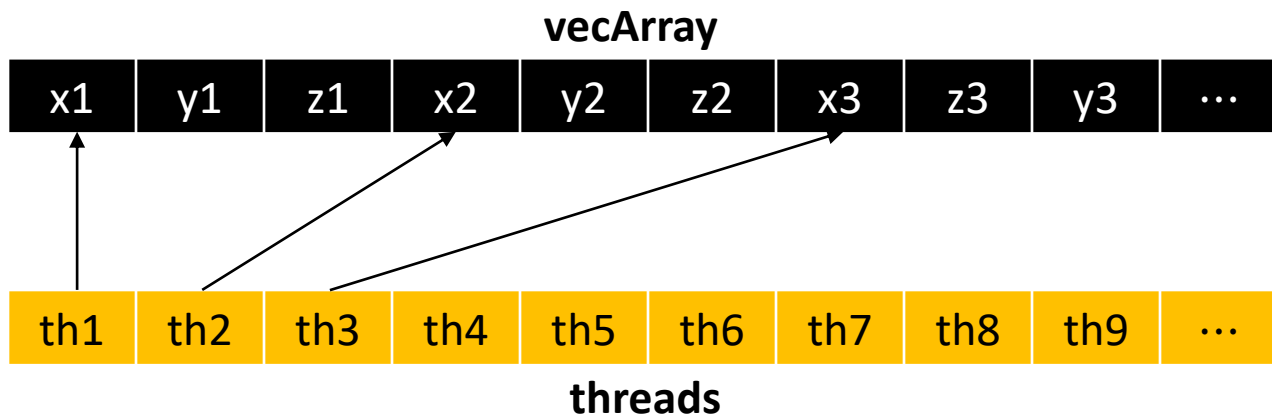
Data structure for fast memory access

- When accessed data is aligned about threads, the number of data transaction required is reduced (it gets faster)
- Each thread of CUDA can access only 1, 2, 4, 8, 16 Bytes at once

Ex1) 3D vector (x,y,z) handling (misaligned memory access)

```
3dVectorHandle(Vector3D* vecArray){  
    Vector3D v = vecArray[threadIdx.x];  
    // this is similar to...  
    // float x = vecArray[threadIdx.x][0];  
    // float y = vecArray[threadIdx.x][1];  
    // float z = vecArray[threadIdx.x][2];  
}
```

Memory is NOT aligned against thread access



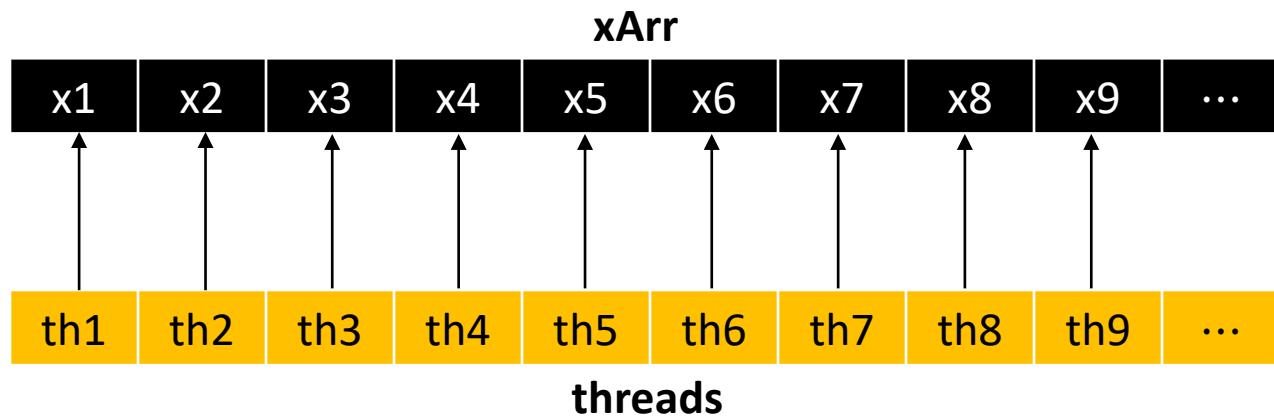
Data structure for fast memory access (cont.)

- When accessed data is aligned about threads, number of data transaction is reduced (it gets faster)
- Each thread of CUDA can access only 1, 2, 4, 8, 16 Bytes at once

Ex2) Aligned memory access

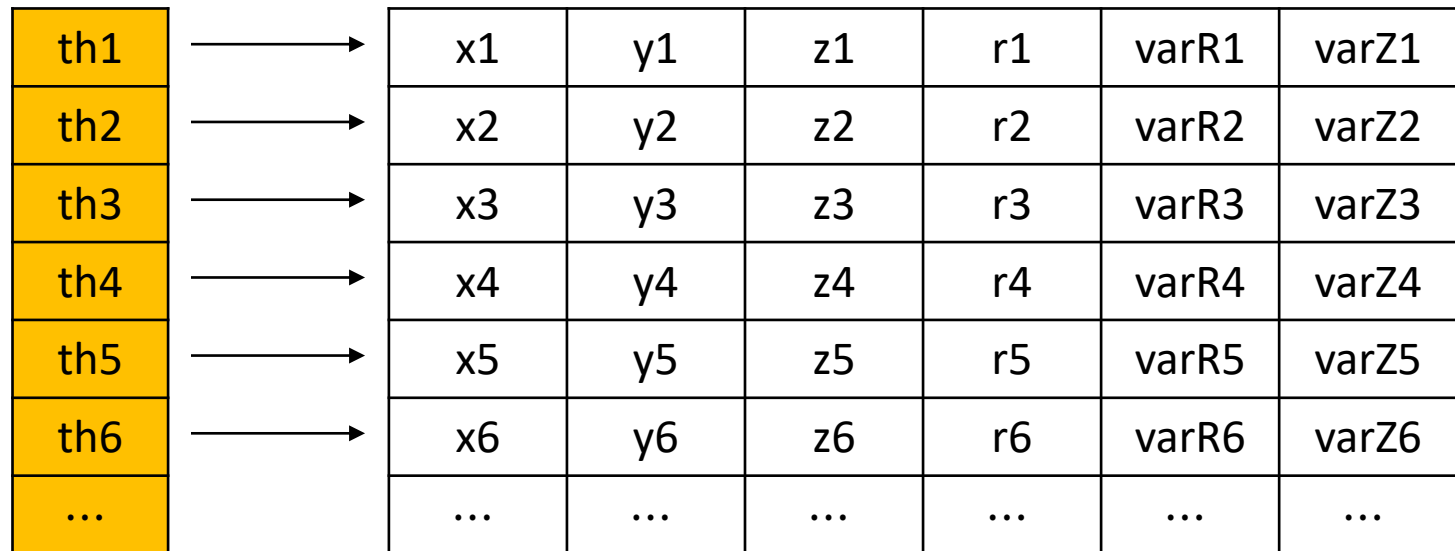
```
3dVectorHandle(float* xArr, float* yArr, float* zArr){  
    float x = xArr[threadIdx.x];  
    float y = yArr[threadIdx.x];  
    float z = zArr[threadIdx.x];  
}
```

Memory is Aligned against thread access

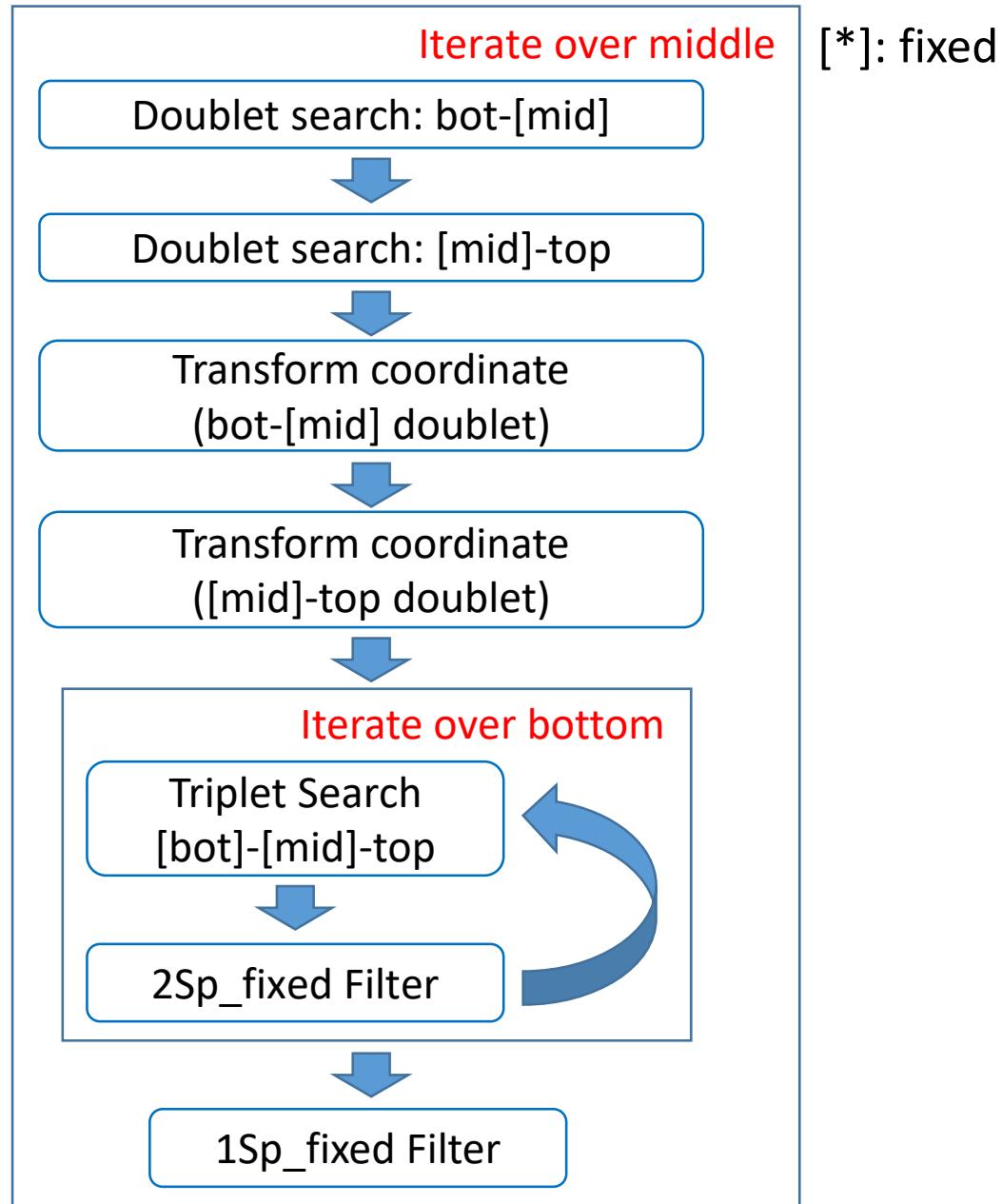


Data structure for fast memory access (cont.)

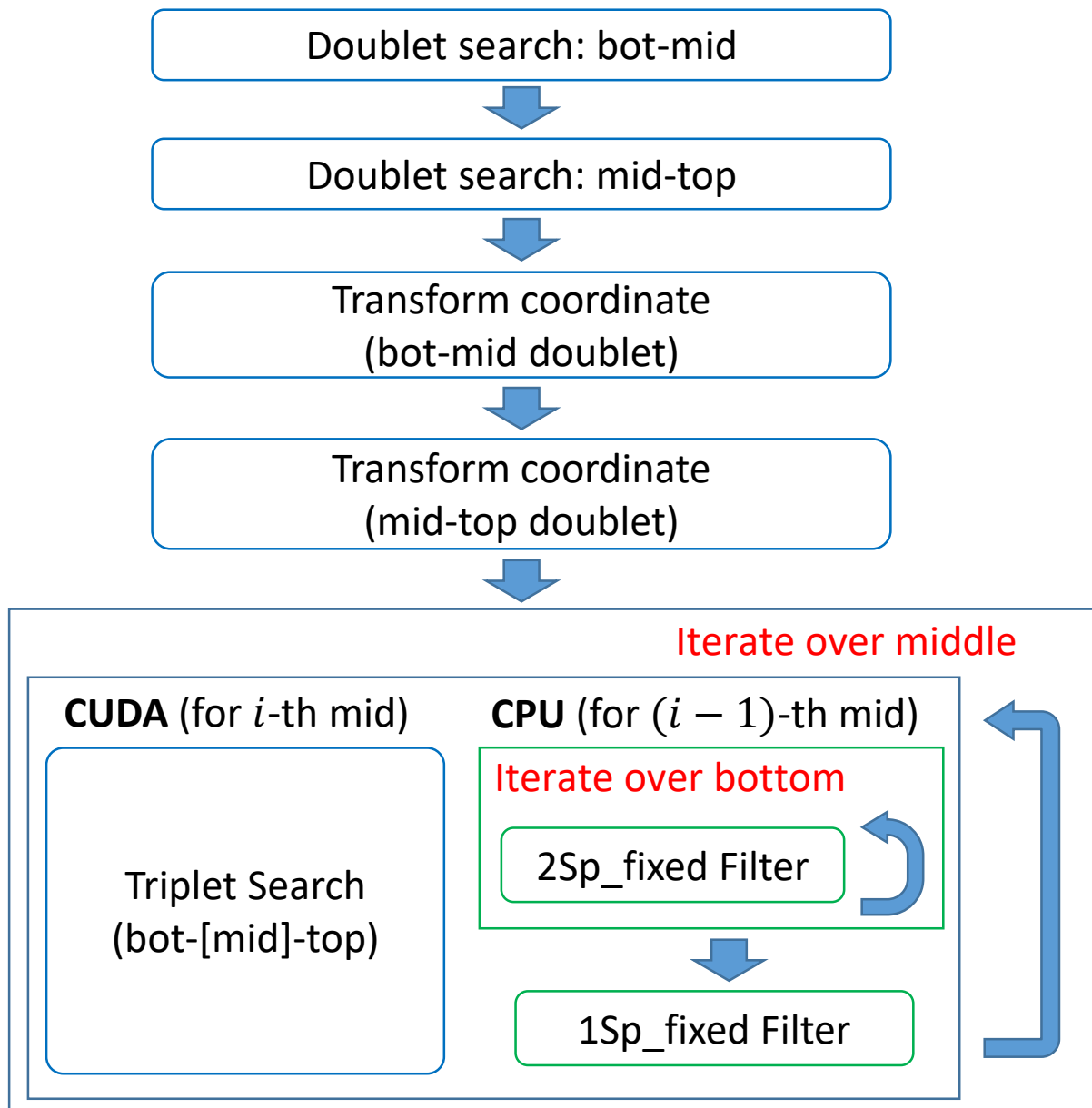
- All spacepoints data were flattened into the matrix (**column-major**) which respects the aligned data structure



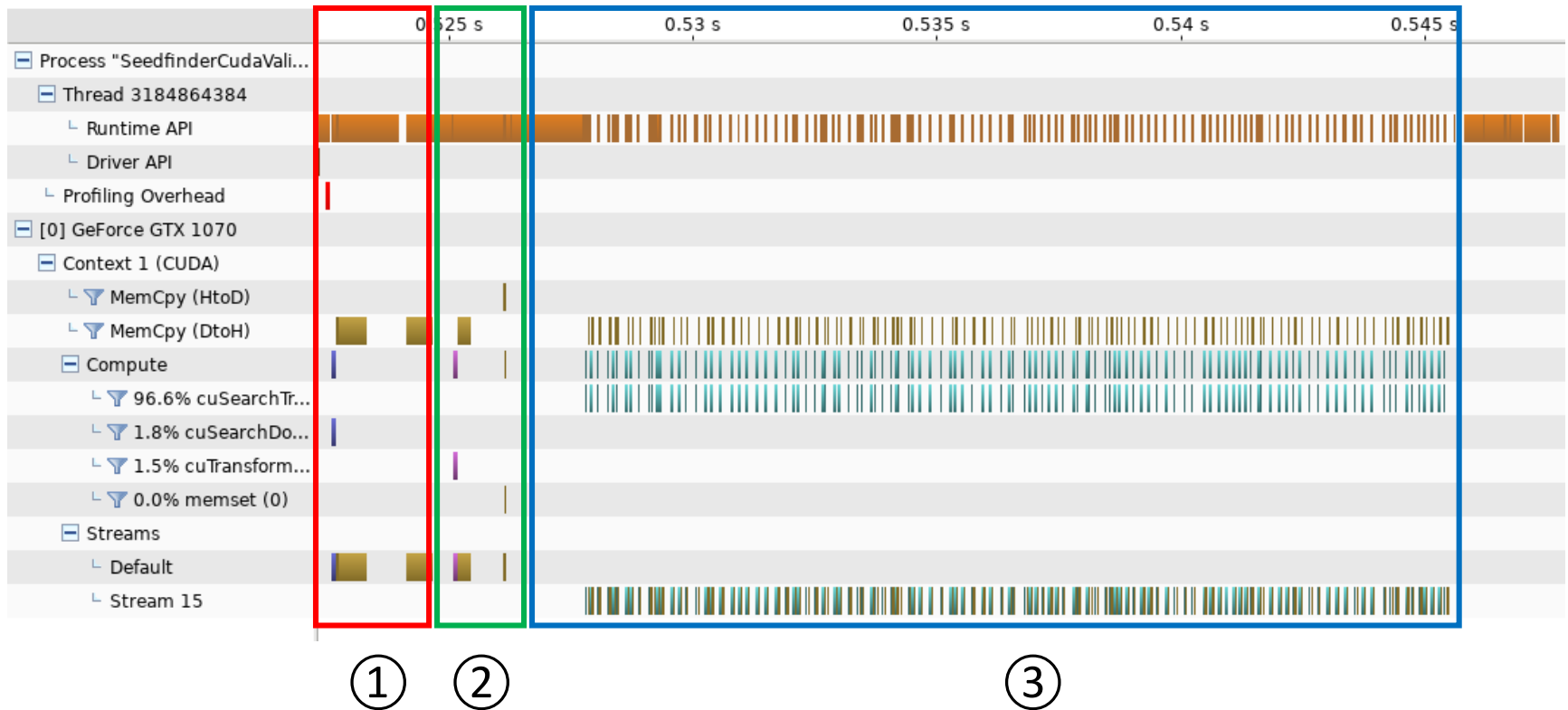
CPU algorithms for a group of space points



CUDA algorithms for a group of space points



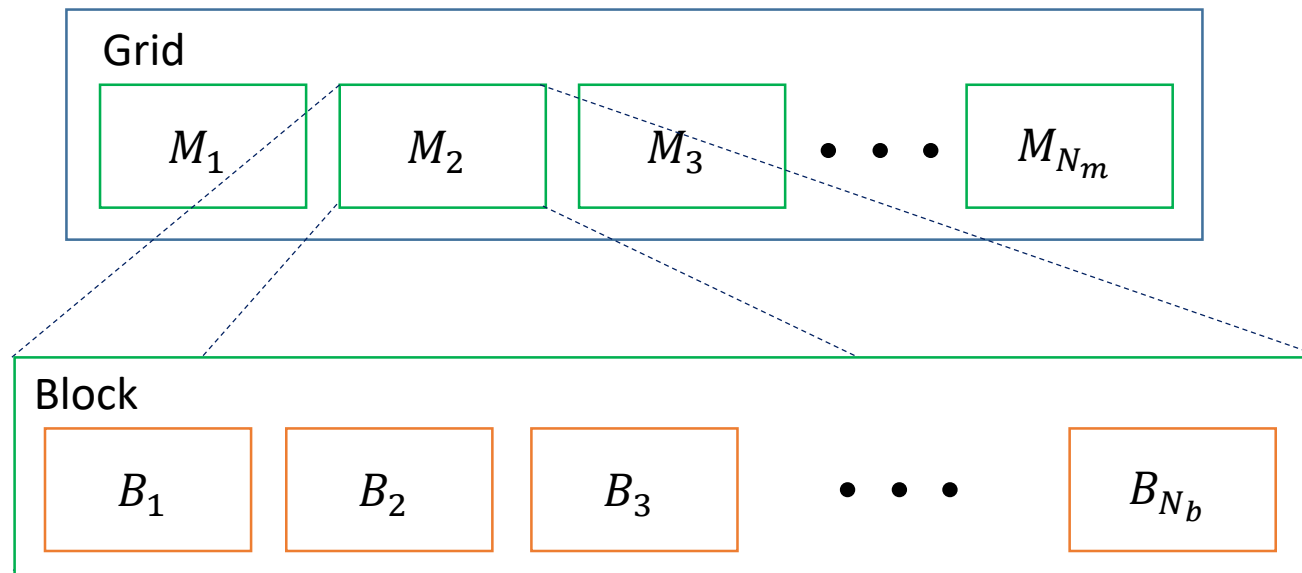
Timeline for CUDA



- ① Doublet Search
- ② Transform coordinate
- ③ Triplet Search + SeedFilter (CPU)

Doublet Search (for middle-bottom)

- **Input** : spacepoint data of middle and bottom hits
- **Output**: index of compatible hits that form doublets



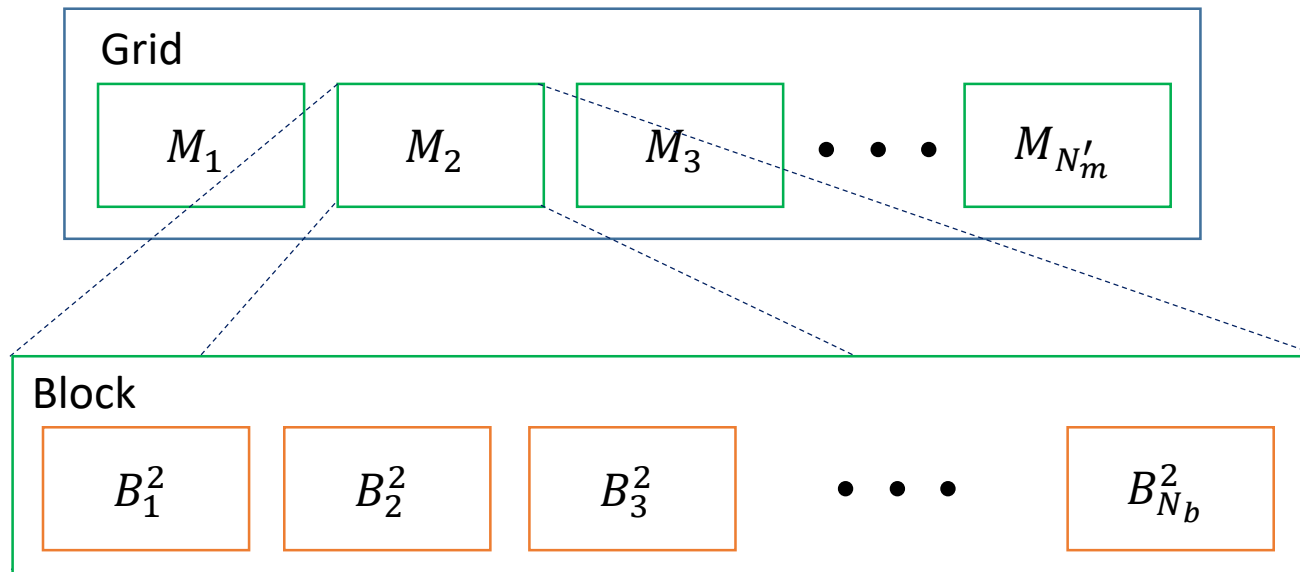
- Since the maximum number of threads per block is 1024, the same kernel is iterated over different sets of bottom hits if $N_b > 1024$

Transform coordinate (for middle-bottom)

Input : spacepoint data middle and bottom hits
index of compatible hits,

Output: (Reduced) spacepoint data transformed into circle

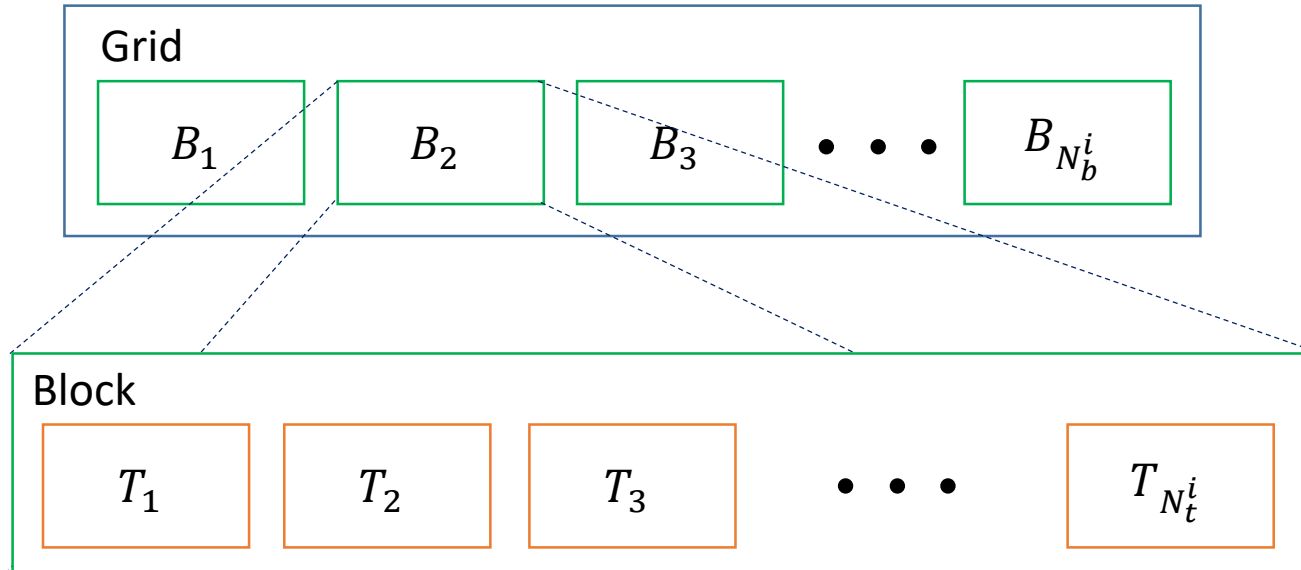
- The matrix size of the output is reduced from the original data to the number of doublets
- The structure is similar to the doublet search



Triplet search (for a middle hit)

Input for the triplet search of i -th middle hit:

(N_b^i) hit & circle matrix and (N_t^i) hit & circle matrix



output for the triplet search of i -th middle hit:

- 1) **Number of top hits which form triplets** for every bottom hit
- 2) curvature and impact parameters of triplets

Timeline of triplet search



- Triplet search for i -th mid and seed filtering for $(i - 1)$ -th mid are done asynchronously
- Seed filtering done by Cpu is the biggest bottleneck

results comparison (CUDA vs. CPU)

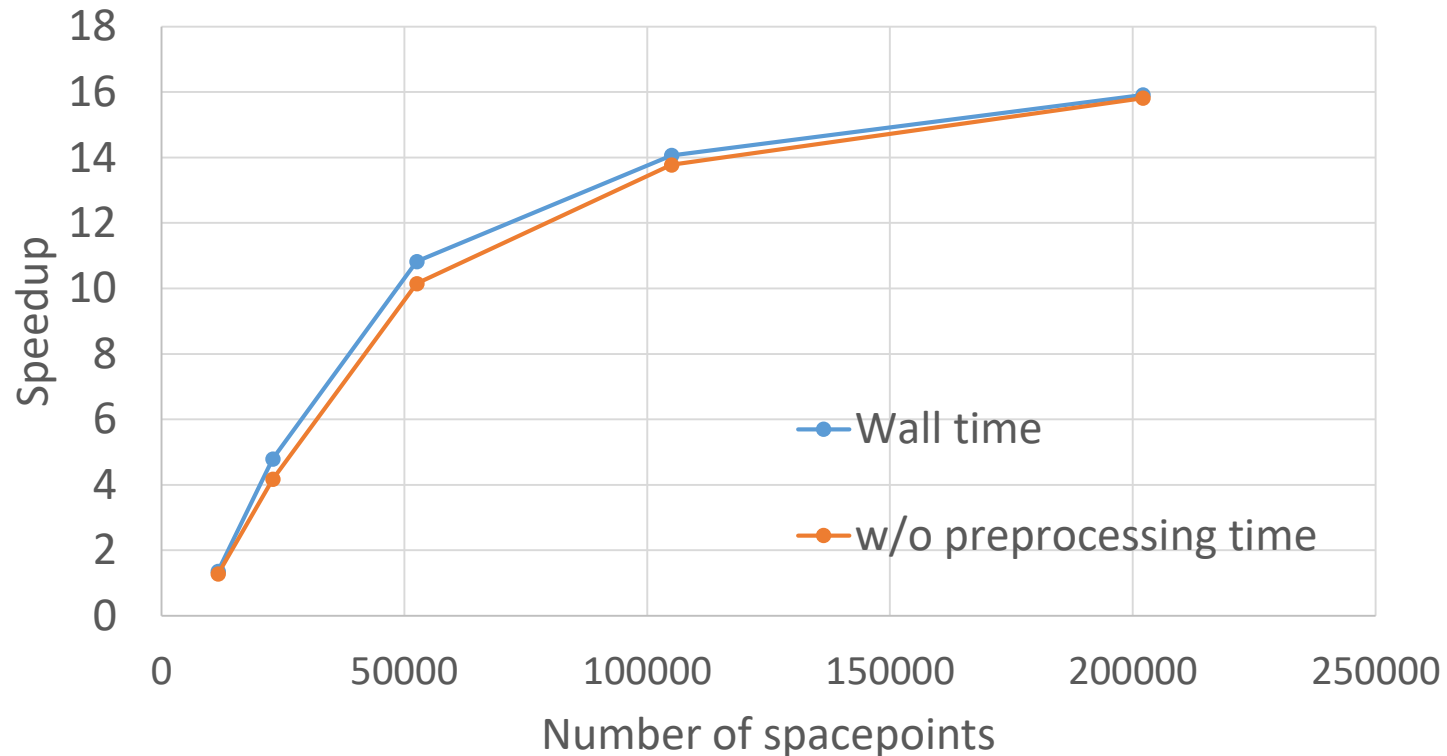
- CPU model: i7-5820K CPU @ 3.3 GHz
- GPU model: GTx 1070

# space points	# of seeds (CUDA)	# of seeds (CPU)	Seed matching ratio
10k	2690	2690	100%
20k	5512	5512	100%
50k	12805	12805	100%
100k	25572	25572	99.99%
200k	49302	49302	99.97%

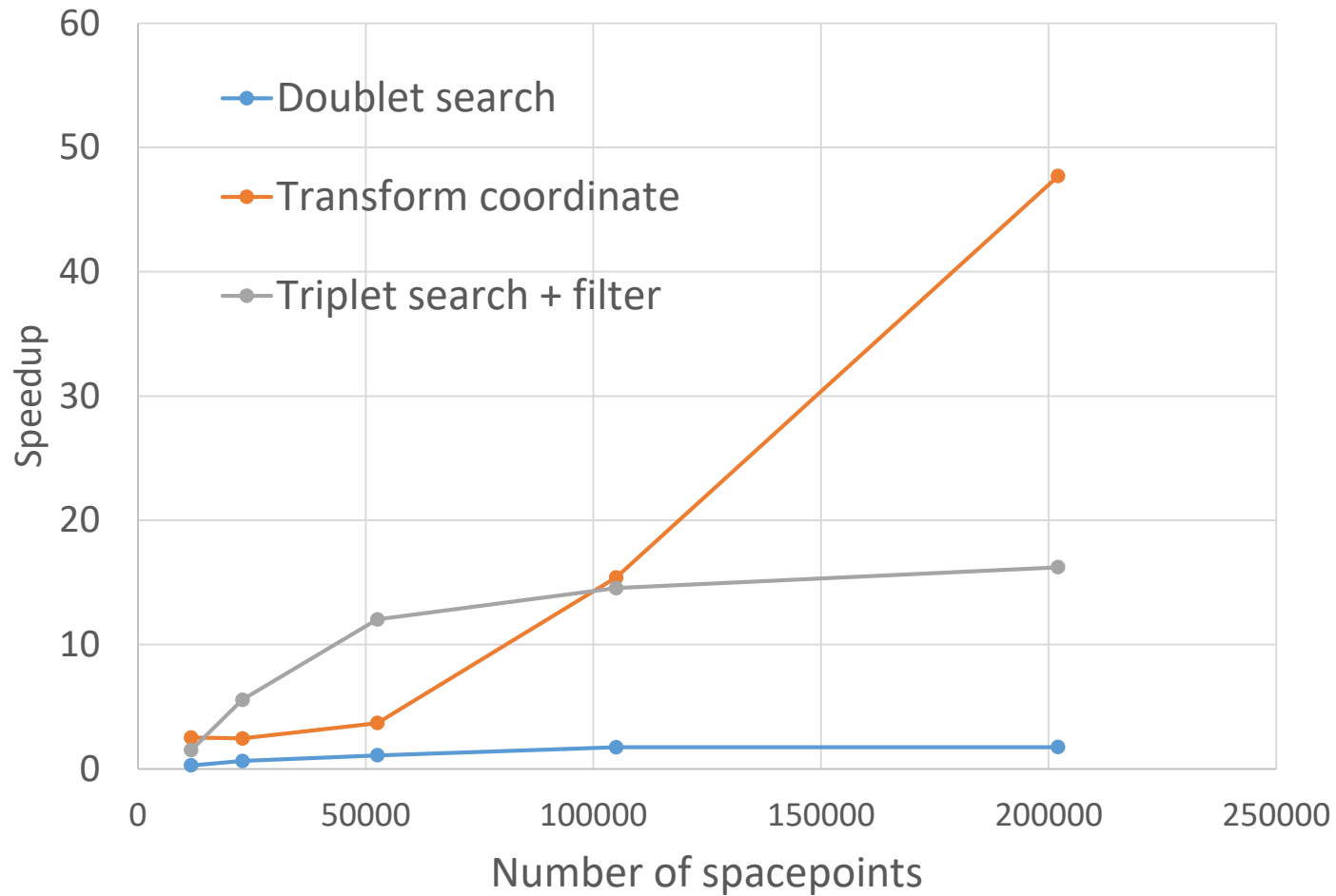
- Seed matching ratio is almost 100%
- the mismatches happen due to the different rounding policy between CPU and GPU

Wall-time speedup (Release mode (-O2))

- Speedup was measured by comparing the wall time:
(preprocessing + CPU seed finding) / (preprocessing + CUDA seed finding)
- Preprocessing includes data reading and grouping



Speedup for each algorithm (Release mode (-O2))



Summary

- Parallelization on the seed finding was done successfully
- Validation tests showed that the seed matching ratio is $\sim 100\%$
- Achieved one order of magnitude improvement in speedups for $>50k$ space points