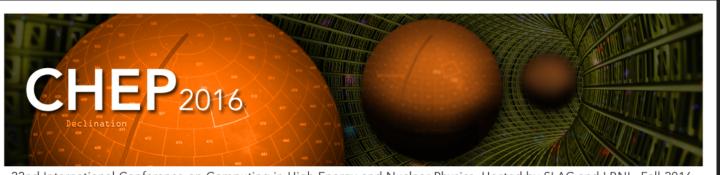


# Parallel metric trees for multi-billion bodies simulations

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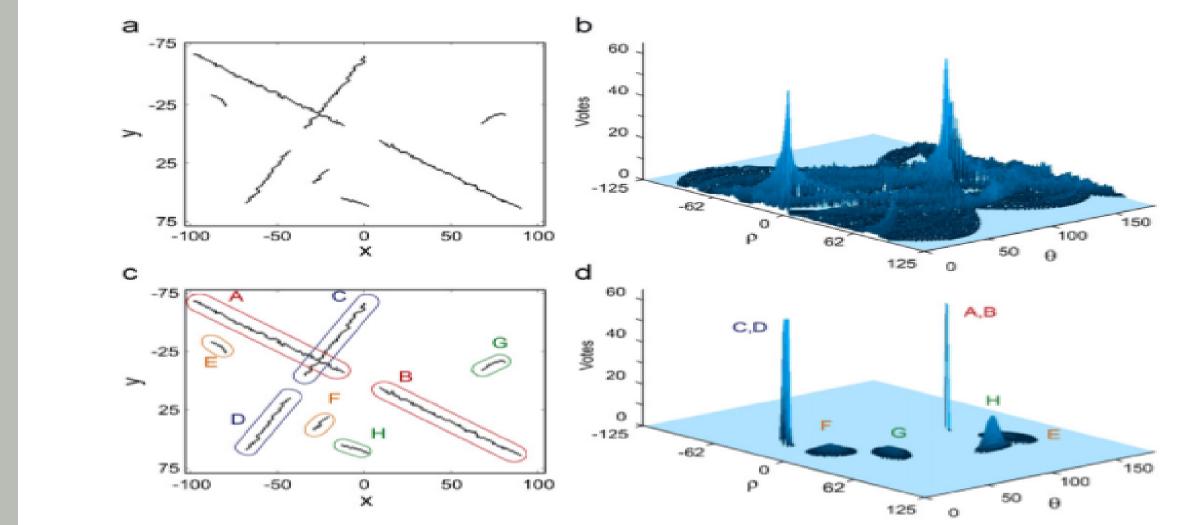
<sup>22</sup>nd International Conference on Computing in High Energy and Nuclear Physics, Hosted by SLAC and LBNL, Fall 201

#### A research question

- Theoretical foundations and tool
  - Computational Geometry
    Vantage point trees
    Metric trees
  - Parallel algorithms
    Contraction algorithms
    Accumulation trees
  - Parallel architectures
    Multi-core
    GPGPU and extended device computing
    Distributed memory archittecures
- The research question

#### Trees for high dimensional data

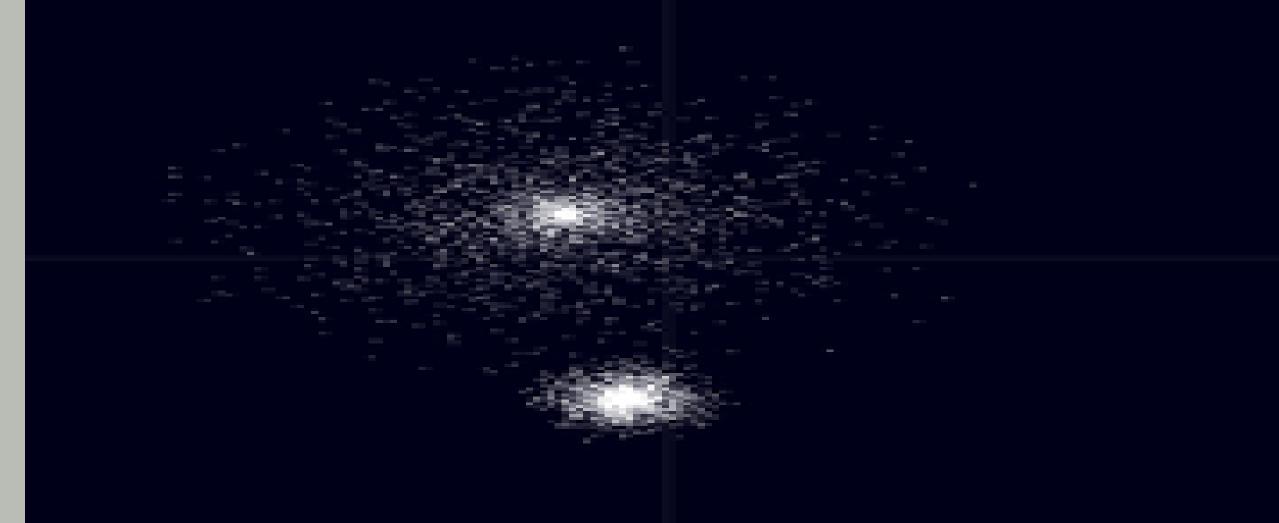
- ► Target
  - Indexing to reduce width, exponentially reduce volume.



How effective can the algorithms and data structures of theorectical parallel computing be when applied to practical problems in high dimensional domains.

# High dimensional data processing examples

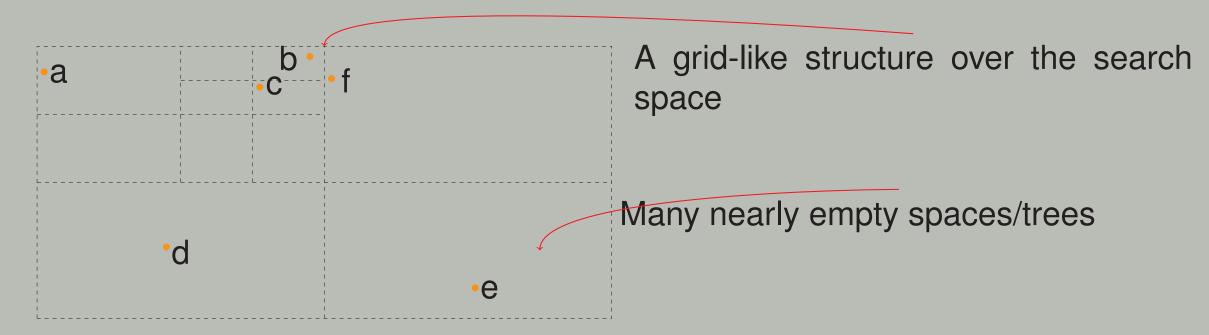
Clustering well separated regions to improve computational efficiency



- Clustering well separated blocks of points to improve line detection.
  Possible figure here showing OpenCV HT failure
- Higher order visualization and sequence matching in DNA or data mining in time series.

Sergey Brin (before Google): *A ball of radius 2 in 20 dimension Euclidean space is a million times larger in volume than a ball of radius* Metric trees can reduce partition the space.

- Balancing of construction to optimize sharing of parallel workload.
  Data structures to separate high dimensional data
- Octrees, and quadtrees tend to be inefficient in space and parallel work balancing;
- k-d-trees can become unbalanced or have irregularly split regions;



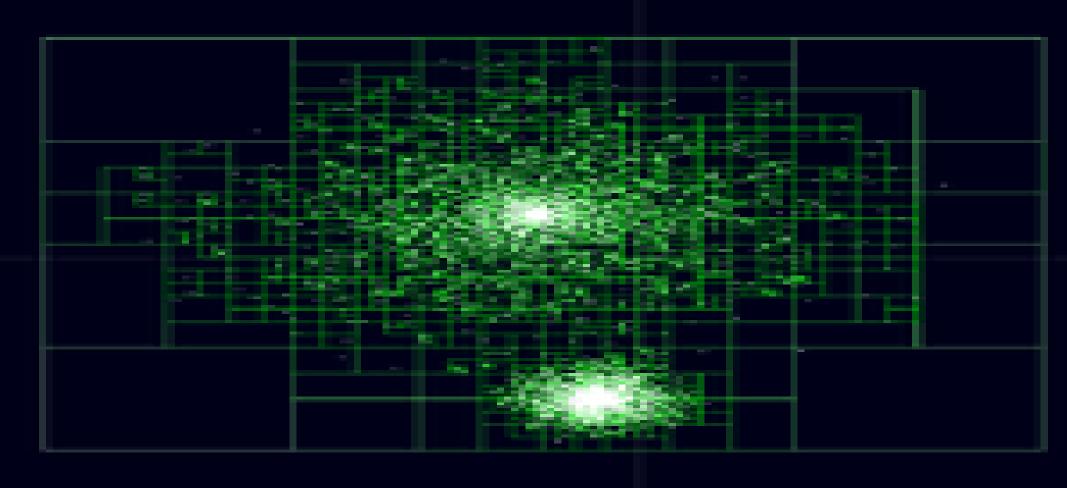
Metric and vantage-point tree might offer round regions and binary trees good for upward and downward accumulation.

### **Metric trees**

- Binary search tree.
- Indexed by distance.
- A node defines a sphere of the points living within a distance from its centre.
- Axioms that regulate the construction of a mertic tree
  - Each node is define by a central point C, a radius r, left and right subtrees.
- ► Points on the left subtree are within a distance *r* from *C*.
- Points on the right subtree are outside distance r from C.

# Computational Geometry data structures

Data separation principles



- Mathematically well founded
- Implemented using metric trees, or quad-trees.
- Example: n-body tree codes.
- A particle is well separated from a region when  $s/d < \theta$  for
- s, the width of the region
- d, distance between the paricle and the centre of mass of the region  $\theta$ , a given parameter.

- Central point is any point in the indexed set.
- Radius is the median of distances to that central point.
- Observations:
- Each node is weight balanced.
- Each node can be split with linear work, if we assume that a median can be computed with linear work.

## **Point separation**

- ► Two sets of points *P* and *Q* are separated when  $\frac{max(diam(Q), diam(R))}{d(Q, R)} < \theta$ 
  - d(Q, R)
- A separated pairs decomposition of a set P is a set of pairs  $(A_i, B_i)$  where
  - each  $A_i, B_i \subset P$
  - $A_i$  and  $B_i$  are  $\theta$ -separated.
- Parallel complexity
  - An  $O(n \lg n)$  work parallel algorithm with depth  $O(\lg n)$  can be defined based the observation that:
  - ▶ if sets P and Q are not separated then the separation set of P and Q is given by the separation sets of P<sub>0</sub> and Q and P<sub>1</sub> and Q, where P<sub>0</sub> and P<sub>1</sub> are children of Q.

# N-body tree code

Algorithm: each Physical time step simulated by a sequence of three paralle bulk steps.

#### Problems

- Parallel algorithm to extract pairs separated regions and particles.
- Parallel accumulation algorithms. Accumulation can be:

FMM Taylor expansion;

Accumulation of force and mass in Barnes-Hut;

Binning (vote counting) in algorithms for variations of Hough Transform.

- In parallel: build a metric tree.
- Complexity: Computation with depth lg n.
- In parallel: build separated sets.
- In parallel: for each pair (A<sub>i</sub>, B<sub>i</sub>), apply cell to cell computation to evalute force, centre of mass, position.

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