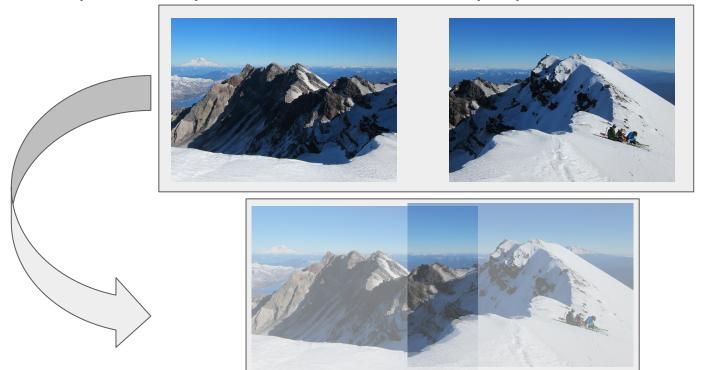
MLRF Lecture 03

J. Chazalon, LRE/EPITA, 2025

Descriptors matching and indexing

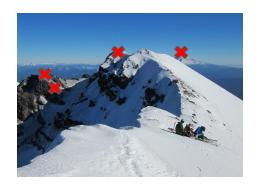
Lecture 03 part 02

How are panorama pictures created from multiple pictures?



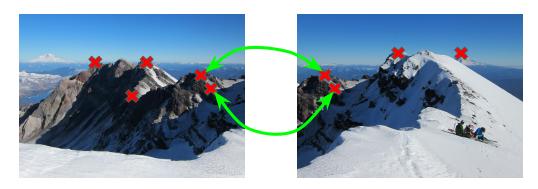
How are panorama pictures created from multiple pictures?





1. Detect small parts invariant under viewpoint change: **keypoints**

How are panorama pictures created from multiple pictures?



- 1. Detect small parts invariant under viewpoint change: **keypoints**
- 2. Find pairs of matching keypoints using a **description** of their neighborhood

How are panorama pictures created from multiple pictures?

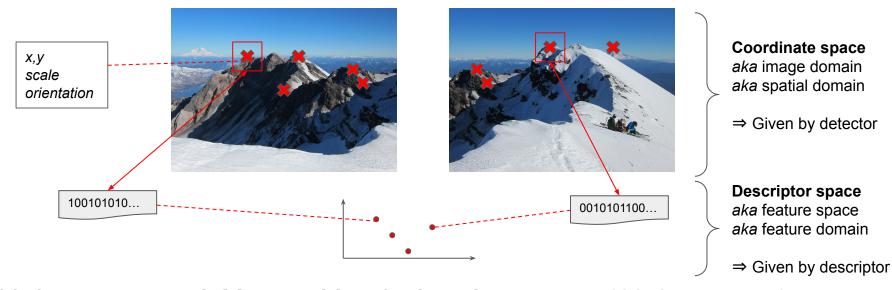






- Detect small parts invariant under viewpoint change: <u>keypoints</u>
- 2. Find pairs of matching keypoints using a **description** of their neighborhood
- 3. Compute the **most likely transformation** to blend images together

Given some keypoints in image 1, what are the more similar ones in image 2?



This is a **nearest neighbor problem** in **descriptor space** (this lecture part). This is also a **geometrical problem** in **coordinate space** (next lecture parts).

Matching

Matching problem

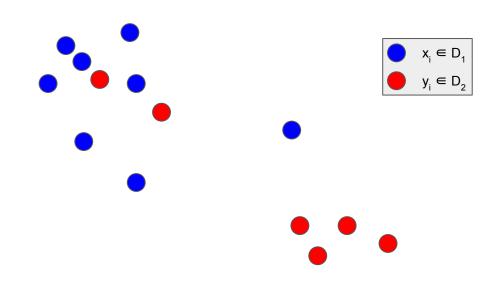
Goal: given two sets of descriptors, find the best **matching pairs**.

Need a **distance/norm**: depends on the descriptor

- Distribution (histogram)? Statistics?
- Data type?
 - Float, integers: Euclidean, cosine...
 - Binary: Hamming...

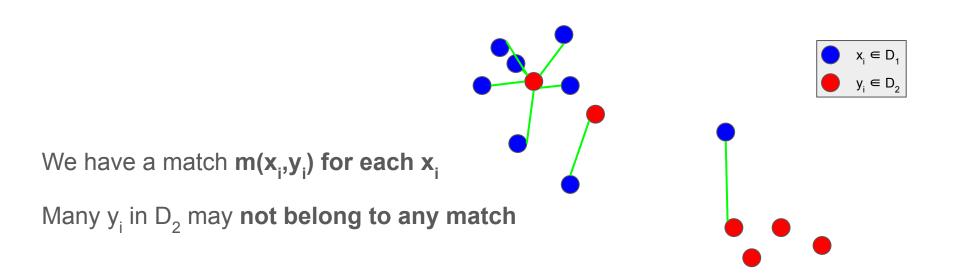
1-way matching

For each x_i in the set of descriptors D_1 , find the closest element y_i in D_2 .



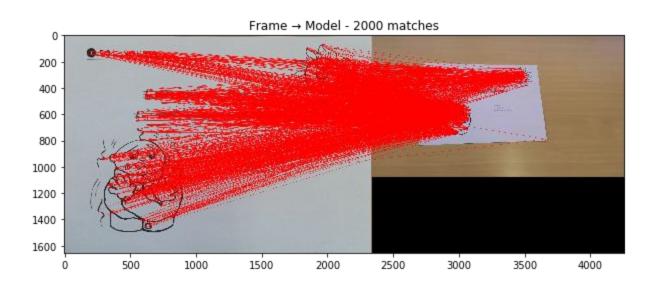
1-way matching

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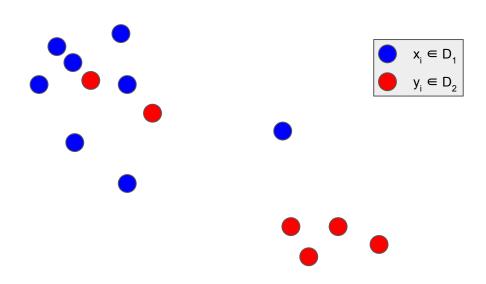
1-way matching

Example from next practice session



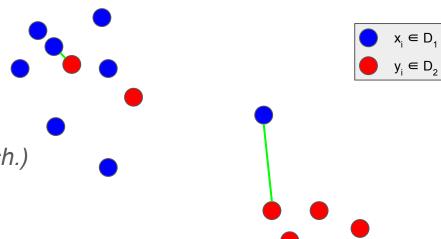
Symmetry test aka cross check aka 2-way matching

For each x_i in the set of descriptors D_1 , find the closest element y_i in D_2 such as x_i is **also the closest** element to y_i .



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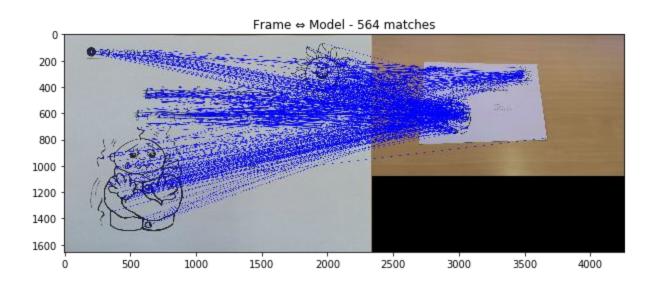
Filters a lot of noise.

(Less matches, not every x_i gets a match.)

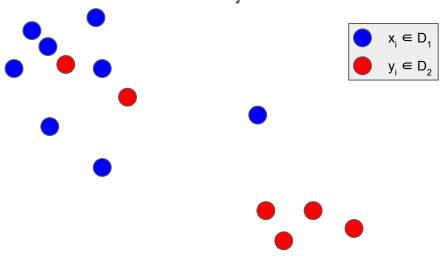
Costly to compute.

Symmetry test aka cross check aka 2-way matching

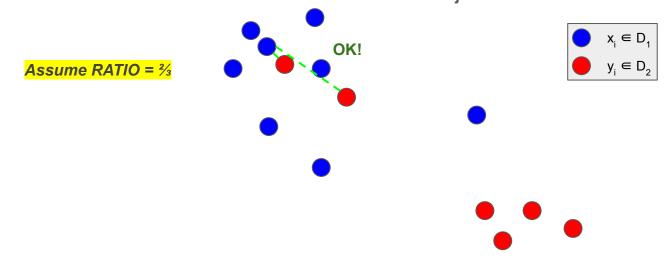
Example from next practice session



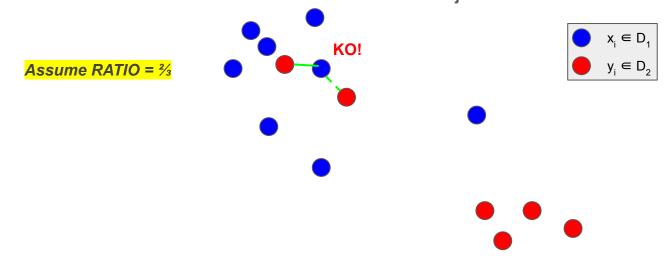
For each x_i in the set of descriptors D_1 , find the 2 closest elements y_i and y_i in D_2 .



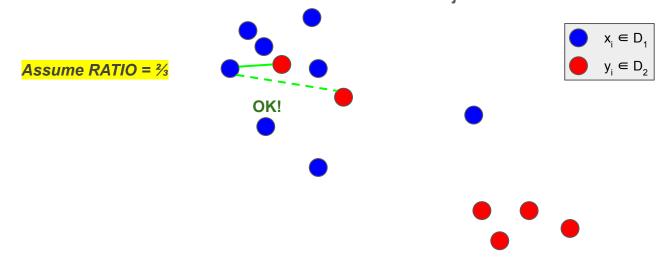
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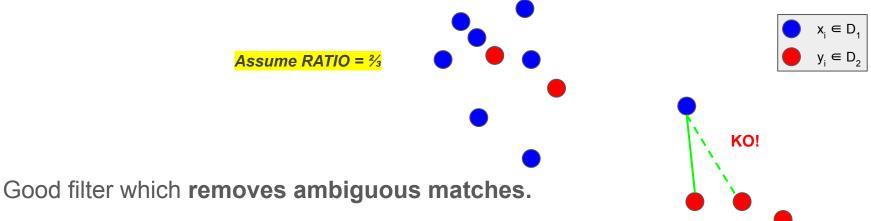


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For each x_i in the set of descriptors D_1 , find the 2 closest elements y_i and y_i in D_2 .

Keep the match $m(x_i, y_i)$ only if $dist(x_i, y_i) < RATIO * dist(x_i, y_i)$



Like 1-way matching, **potential y**, **duplicates** in matches.

Can be made symmetrical.

Ratio test: calibrate the ratio

Adjust it on a training set!

For each correct/incorrect match in your annotated database, plot the *next to next closest distance* PDF.

What is a good ratio in D. Lowe's experiment? →

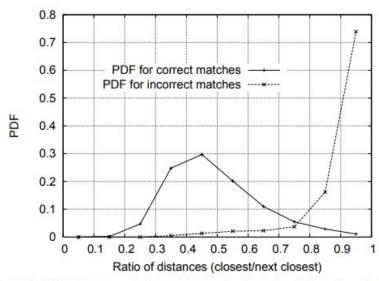
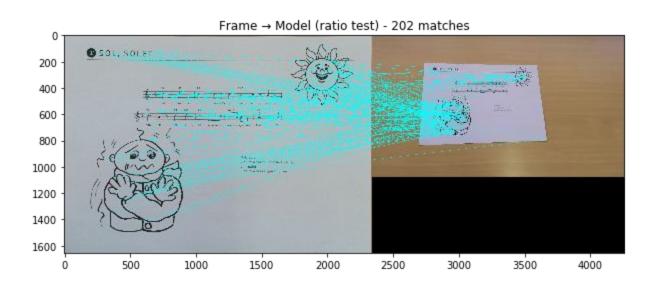
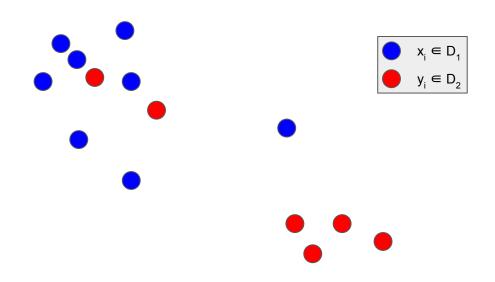


Figure 11: The probability that a match is correct can be determined by taking the ratio of distance from the closest neighbor to the distance of the second closest. Using a database of 40,000 keypoints, the solid line shows the PDF of this ratio for correct matches, while the dotted line is for matches that were incorrect.

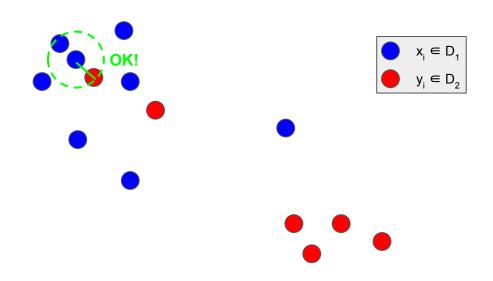
Example from next practice session



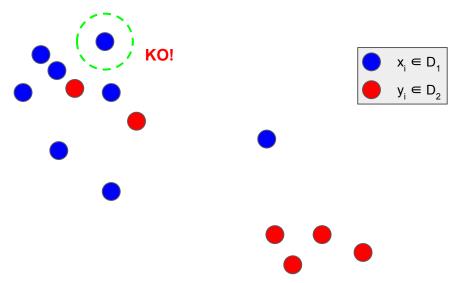
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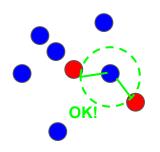


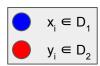
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For each x_i in the set of descriptors D_1 , find the closest element y_i in D_2 and make sure **dist(x_i, y_i) < RADIUS**

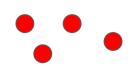
May allow multiple good matches for some x_i





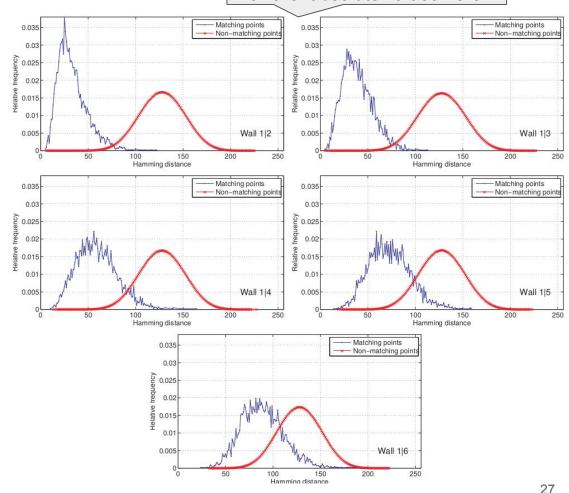
Harder to calibrate

- 1 absolute value for all descriptor space!
- Usually req. a "background model"
 - = a set D₃ with **only incorrect matches**



From BRIEF paper.

If we have a background model which give us the red curve for each case (not knowing the blue one), can we choose a good radius?

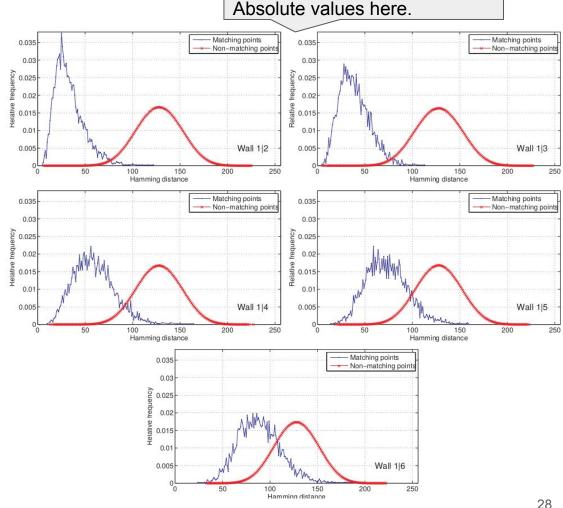


Beware: absolute values here!

From BRIEF paper.

If we have a background model which give us the red curve for each case (not knowing the blue one), can we choose a good radius?

75-125 seems good here!



Example from next practice session

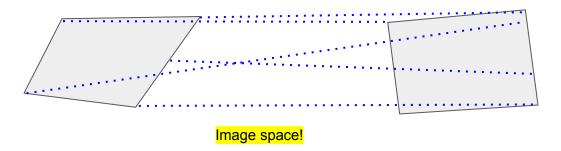
[Missing because not useful for this session]

[and tricky to handle multiple good matches]

What about the coordinates of the keypoints?

Once we have a list of matches $m(x_i, y_i)$,

we can check whether the coordinates of the keypoints of the matched descriptors describe a consistent mapping from one position to another.



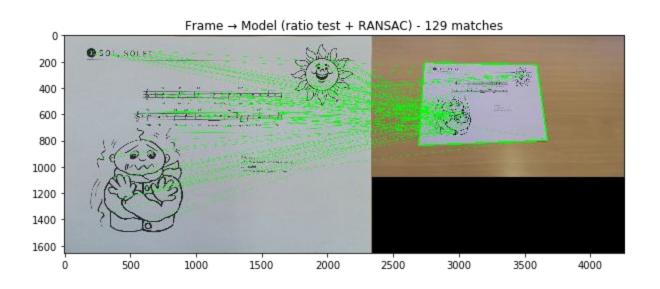
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Example from next practice session



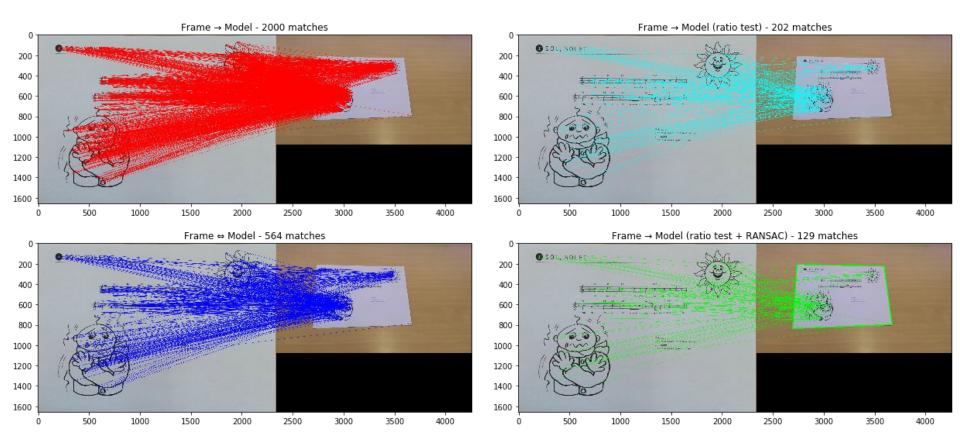
How to check the consistency of the transformation?

Difference classes for transformations.

Different methods to estimate them and check which matches agree and disagree.

⇒ Next lecture parts.

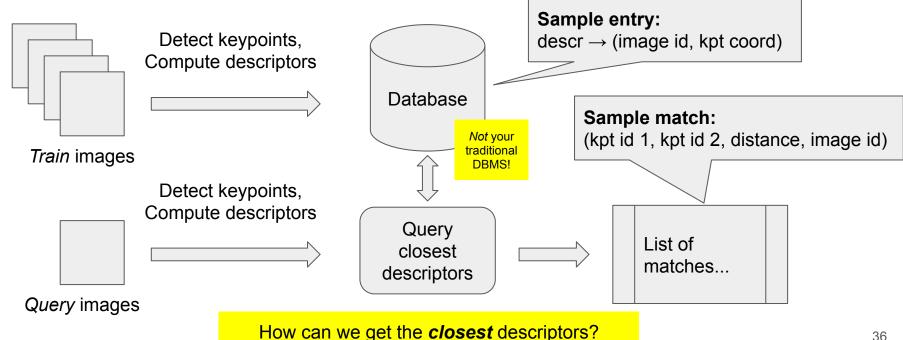
Summary of matching techniques (for practice session)



Indexing

Indexing pipeline

Use case: We have a database of images and we want to find an object from it.



Bruteforce matching aka linear matching

Simply scan all data and keep the closest elements.

Does not scale to large databases, but can be faster on small ones! Especially with fast distance measures, like Hamming.

Exact matching. Global optimum guarantee.

Supports cross check (double scan).

Indexing

Build **one or more indexes of descriptors** → descriptor data

Indexing is often approximate (especially if asking for more than 1 neighbor), because :

- 1. Databases can grow very large
- 2. Descriptor spaces have many dimensions

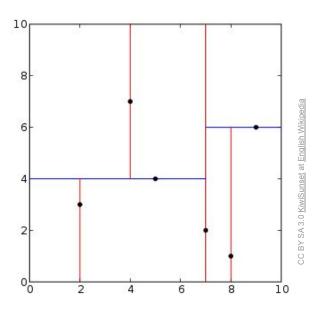
Exact matching and global optimum are **not always guaranteed**.

Also, cross check usually does not make sense and is therefore not implemented.

Usually, we start by **reducing the dimension** / **encoding our features** (next lecture)

kD-Trees

The k-d tree is a binary tree in which every leaf node is a k-dimensional point.



kD-Trees

The k-d tree is a binary tree in which every leaf node is a k-dimensional point.

Construction: for each dimension, recursively split the space to maximize data separation until a maximum size is reached

Retrieval: compute the leaf node of each query, then explore points in the leaf and in siblings / parents if criterion not satisfied (boundaries not within radius of query's acceptance hyperball)

Complexity: asymptotic O(log N) when N>>2k

In practice, kD-trees do not work for searching in <u>high</u> dimension spaces.

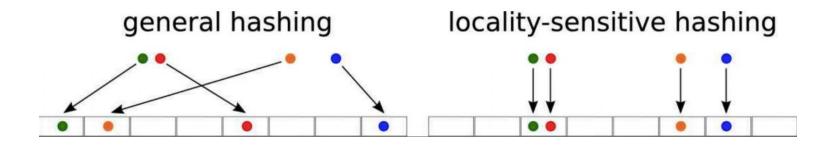
FLANN – Efficient indexing

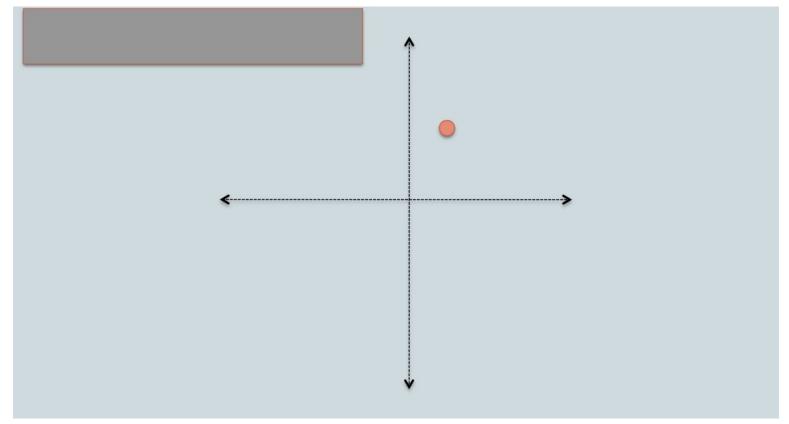
Original version: hierarchical k-Means.

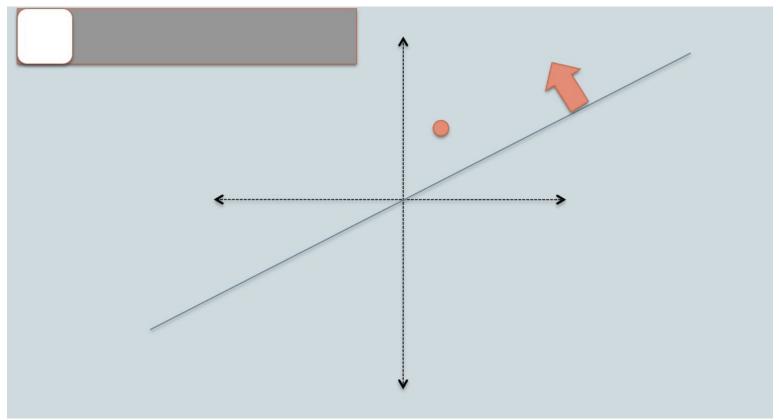
Construction: repetitive k-Means on data (then inside clusters) until minimum cluster size is reached.

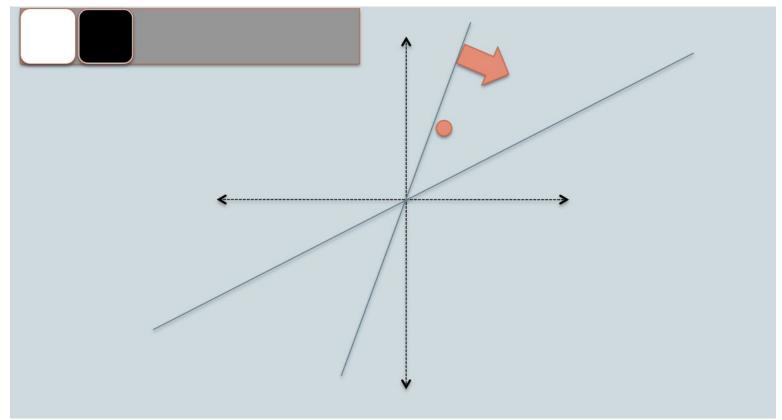
Retrieval: traverse the tree in a best-bin-first manner with backtrack queue, backtrack until enough points are returned

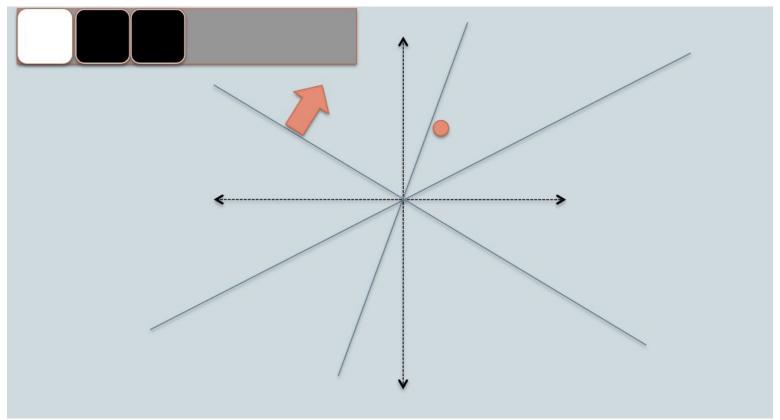
Hash items using a family of hash function which project similar items in the same bucket with high probability. **NOT cryptographic hashing!**

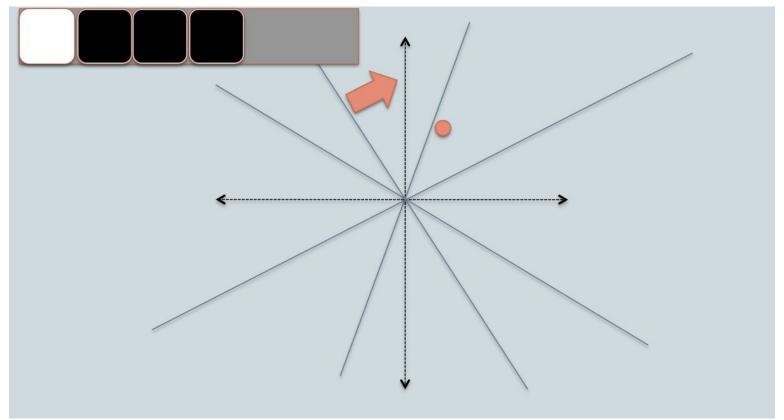


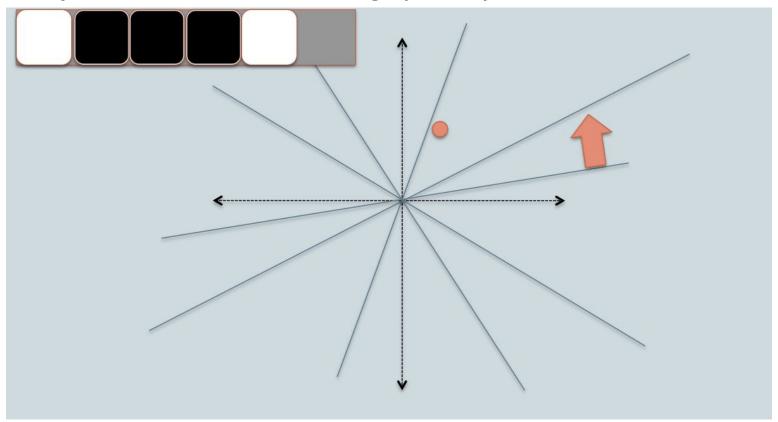


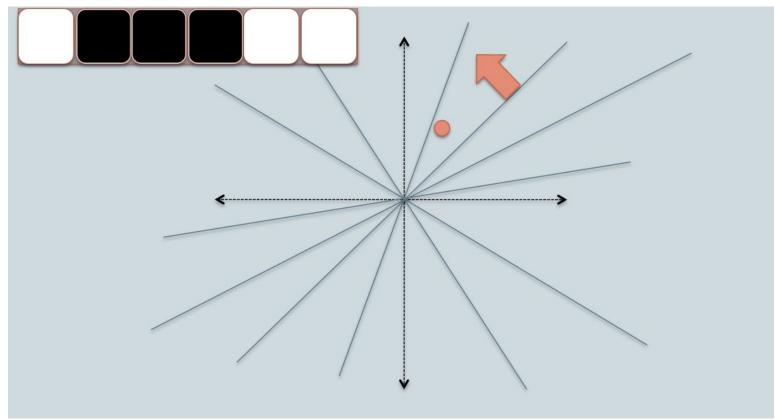


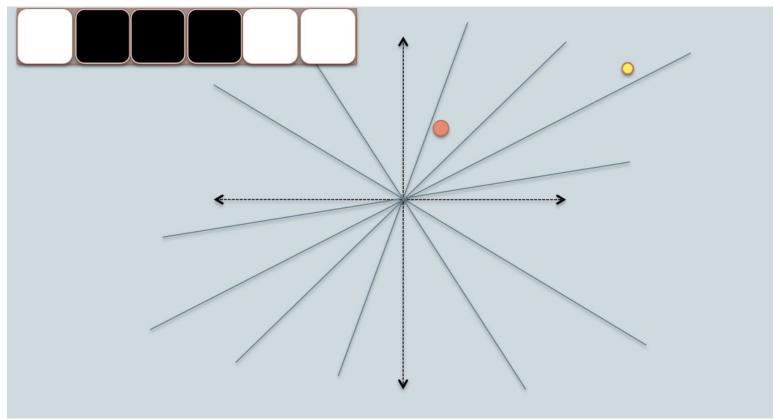


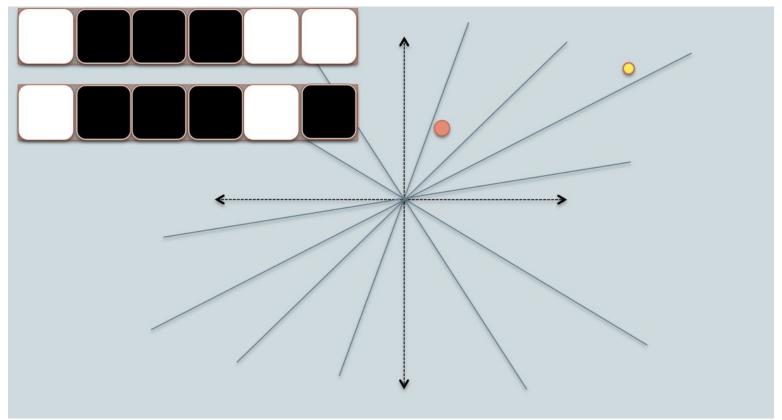


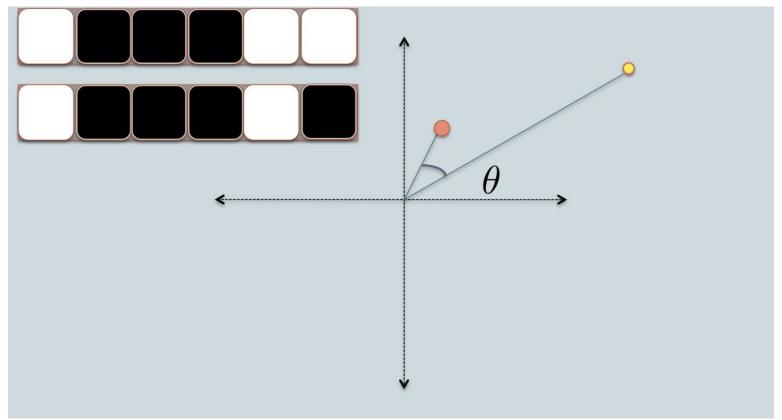


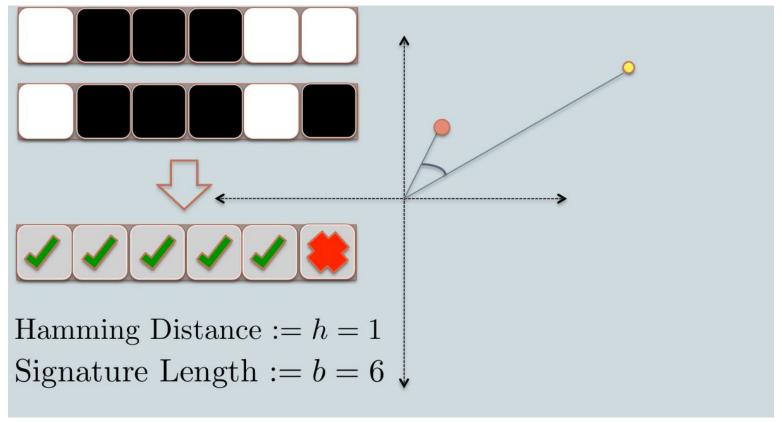


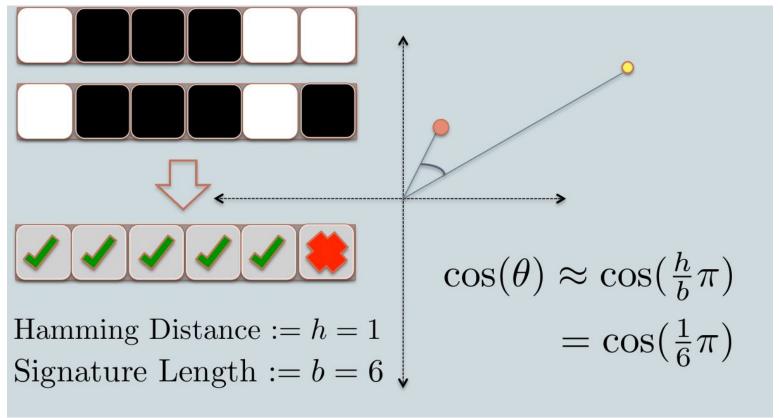


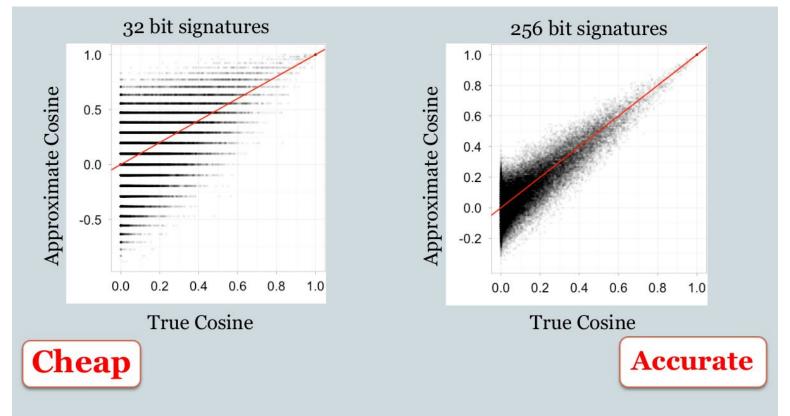












Fast and efficient with large spaces, lot of data.

Return a "good match", maybe not the best one.

kNN can be costly (scan other bins).

Indexes binary descriptors very straightforwardly using bit sampling (sample bits from the coordinates).

Random projections for other cases, or other techniques...

Which indexing?

Experiment.

Advices for practice session:

- Use bruteforce/linear for first experiments: best (but slow) results
- Use LSH for binary descriptors
 like ORB
- Use randomized kD-Trees with SIFT (integer descr. with similar dimension) for moderate dataset size,

k-Mean tree otherwise

