### The Inverted Multi-Index

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joint work with Artem Babenko Yandex



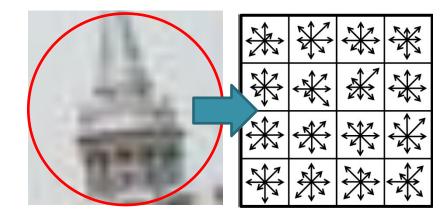
# From images to descriptors



*Interesting point detection:* 



*Interesting point description:* 



Set of 128D descriptors

# **Query process**

### *Image set:*



Dataset of visual descriptors









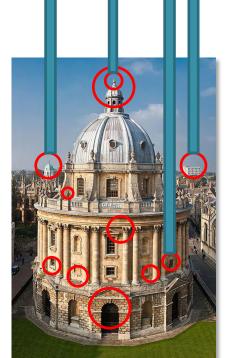
### Main operation:

Finding similar descriptors

### *Important extras:*

- + geometric verification
- + query expansion

Query:



### **Demands**

### Initial setup:

Dataset size: few million images

Typical RAM size: few dozen gigabytes

Tolerable query time: few seconds

Each image has ~1000 descriptors

### Search problem:

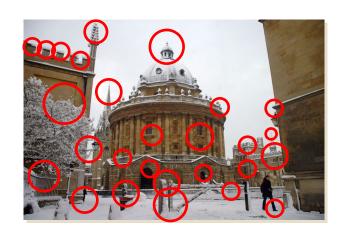
Dataset size: few billion features

Feature footprint: ~ a dozen bytes

Tolerable time: few milliseconds per feature

nearest neighbor search problem we are tackling





## Meeting the demands

**Main observation**: the vectors have a specific structure: correlated dimensions, natural image statistics, etc...

### **Technologies:**

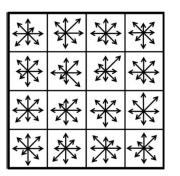
- Dimensionality reduction
- Vector quantization
- Inverted index
- Locality-sensitive hashing
- Product quantization
- Binary/Hamming encodings

### Best combinations (previous state-of-the-art):

- Inverted index + Product Quantization [Jegou et al. TPAMI 2011]
- Inverted index + Binary encoding [Jegou et al. ECCV 2008]

### New state-of-the-art for BIGANN:

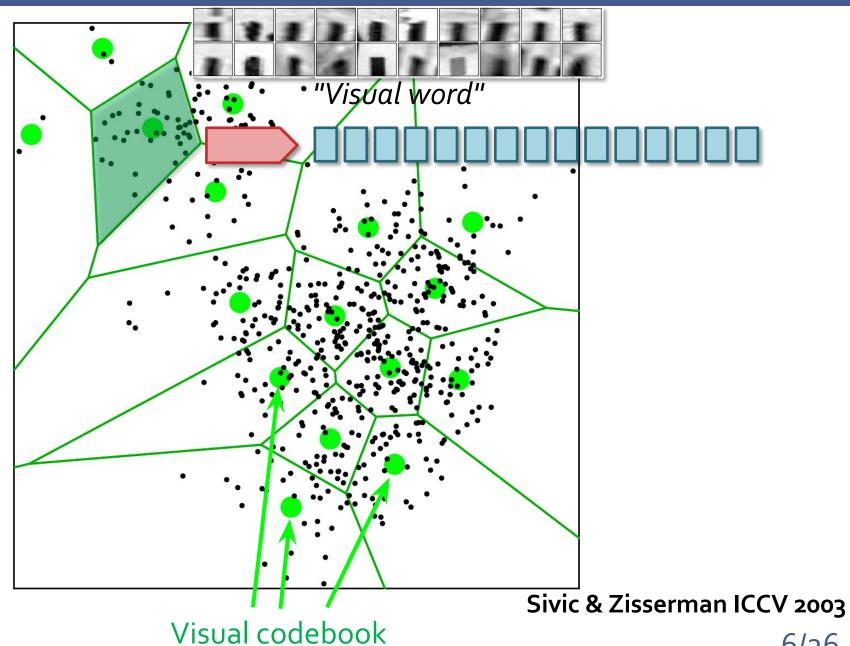
Inverted multi-index + Product Quantization [CVPR 2012]



#### Our contribution:

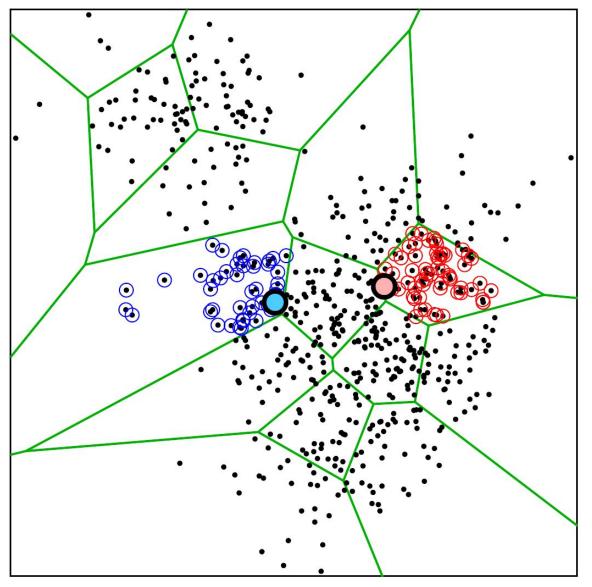
#### **Inverted Multi-Index**

# The inverted index

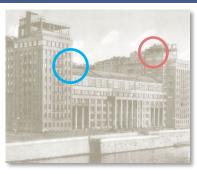


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# Querying the inverted index



Query:

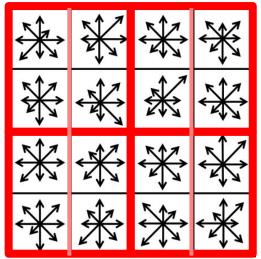


- Have to consider several words for best accuracy
- Want to use as big codebook as possible



 Want to spend as little time as possible for matching to codebooks

## Product quantization



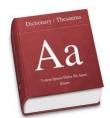
### [Jegou, Douze, Schmid // TPAMI 2011]:

- 1. Split vector into correlated subvectors
- 2. use separate small codebook for each chunk

### **Quantization vs. Product quantization:**

For a budget of 4 bytes per descriptor:





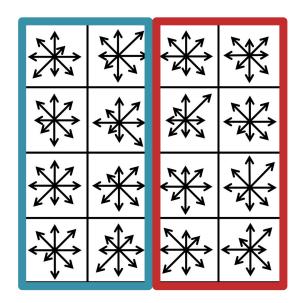
- 1. Can use a single codebook with 1 billion codewords
- 2. Can use 4 different codebooks with 256 codewords each



IVFADC+ variants (state-of-the-art for billion scale datasets) = inverted index for indexing + product quantization for reranking

## The inverted multi-index

Our idea: use product quantization for indexing

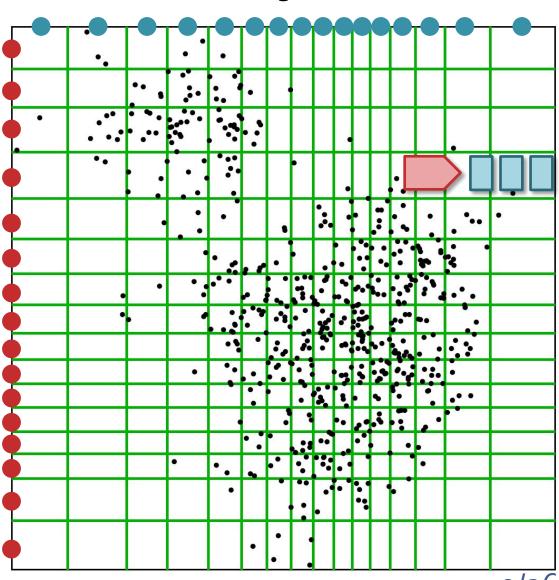


### Main advantage:

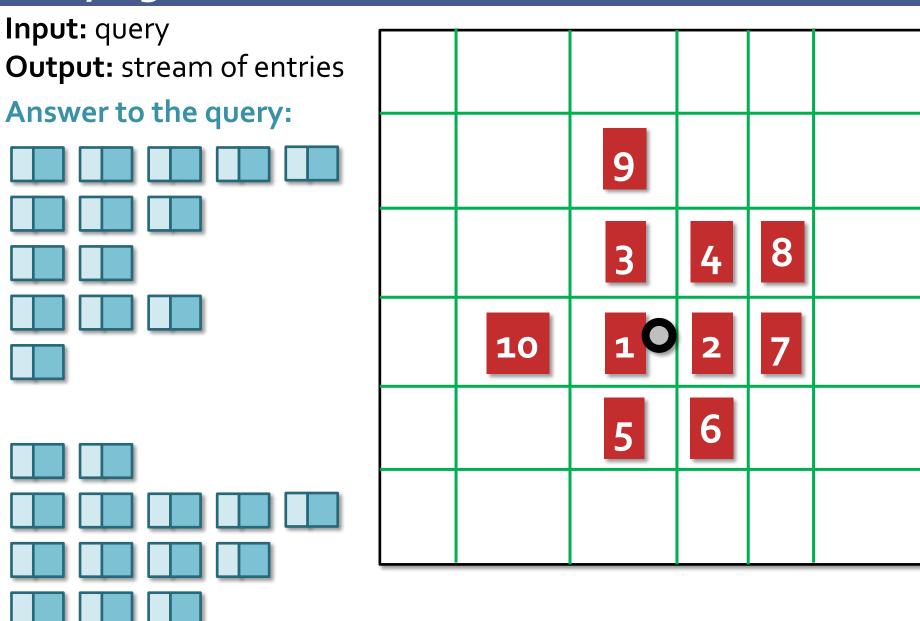
For the same K, much finer subdivision achieved

#### Main problem:

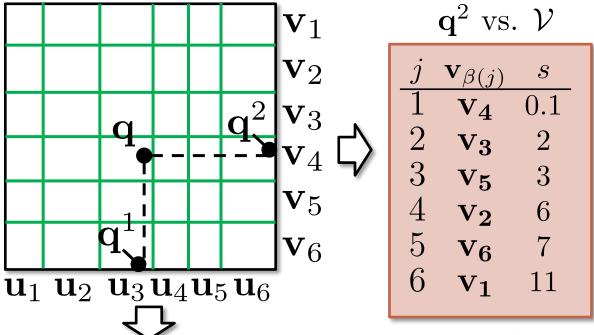
Very non-uniform entry size distribution



# Querying the inverted multi-index



## Querying the inverted multi-index – Step 1



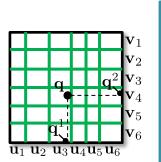
 $\mathbf{q}^1$  vs.  $\mathcal{U}$ 

i	$\mathbf{u}_{lpha(i)}$	r
$\overline{1}$	$\mathbf{u_3}$	0.5
2	$\mathbf{u_4}$	0.7
3	$\mathbf{u_5}$	4
4	$\mathbf{u_2}$	6
5	$\mathrm{u}_1$	8
6	$\mathbf{u_6}$	9

	inverted index	inverted multi-index
number of entries	K	K <sup>2</sup>
operations to match to codebooks	2K+O(1)	2K+O(1)

# Querying the inverted multi-index — Step 2

### **Step 2:** the multi-sequence algorithm



$$\mathbf{q}^1 \text{ vs. } \mathcal{U} \qquad \mathbf{q}^2 \text{ vs. } \mathcal{V}$$

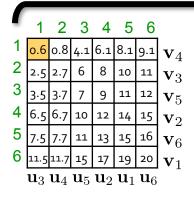
i	$\mathbf{u}_{lpha(i)}$	r
1	$u_3$	0.5
2	$\mathbf{u_4}$	0.7
3	$\mathbf{u_5}$	4
4	$\mathbf{u_2}$	6
5	$\mathrm{u}_1$	8
6	$\mathbf{u_6}$	9

$$\mathbf{q}^2$$
 vs.  $\mathcal{V}$ 

$\underline{j}$	$\mathbf{v}_{\beta(j)}$	s
1	${f v_4}$	0.1
2	${f v_3}$	2
3	${f v_5}$	3
4	$\mathbf{v_2}$	6
5	${f v_6}$	7
6	$\mathbf{v_1}$	11



$[\mathbf{u}_{lpha(i)} \; \mathbf{v}_{eta(j)}]$	(i, j)	r(i) + s(j)
$\overline{}[\mathrm{u_3}\;\mathrm{v_4}]$	(1,1)	$0.6 \ (0.5+0.1)$
$[\mathbf{u_4} \; \mathbf{v_4}]$	(2,1)	$0.8 \ (0.7+0.1)$
$[\mathbf{u_3} \ \mathbf{v_3}]$	(1,2)	2.5 (0.5+2)
$[\mathbf{u_4} \ \mathbf{v_3}]$	(2,2)	2.7 (0.7+2)
$[\mathbf{u_3} \; \mathbf{v_5}]$	(1,3)	3.5 (0.5+3)
$[\mathbf{u_4} \ \mathbf{v_5}]$	(2,3)	3.7 (0.7+3)
$[\mathbf{u_5} \; \mathbf{v_4}]$	(3,1)	4.1 (4+0.1)
$[\mathbf{u_5} \; \mathbf{v_3}]$	(3,2)	6 (4+2)
$[\mathbf{u_3} \ \mathbf{v_2}]$	(1,4)	6.5 (0.5+6)



1	2	3	4	5	6		
o.6	0.8	4.1	6.1	8.1	9.1		
2.5	2.7	6	8	10	11		
3.5	3.7	7	9	11	12		
6.5	6.7	10	12	14	15		
7.5	7.7	11	13	15	16		
11.5	11.7	15	17	19	20		
$\mathbf{u}_3  \mathbf{u}_4  \mathbf{u}_5  \mathbf{u}_2  \mathbf{u}_1  \mathbf{u}_6$							

1	2	3	4	5	6
0.6	o.8	4.1	6.1	8.1	9.1
2.5	2.7	6	8	10	11
3.5	3.7	7	9	11	12
6.5	6.7	10	12	14	15
7.5	7.7	11	13	15	16
11.5	11.7	15	17	19	20
$\mathbf{u}_3$	$\mathbf{u}_4$	$\mathbf{u}_5$	$\mathbf{u}_2$	$\mathbf{u}_1$	$\mathbf{u}_6$

_1	2	3	4	5	6
0.6	0.8	4.1	6.1	8.1	9.1
2.5	2.7	6	8	10	11
3.5	3.7	7	9	11	12
6.5	6.7	10	12	14	15
7.5	7.7	11	13	15	16
11.5	11.7	15	17	19	20
$\overline{\mathbf{u}_3}$	$\overline{\mathbf{u}_4}$	$\mathbf{u}_5$	$\mathbf{u}_2$	$\mathbf{u}_1$	$\mathbf{u}_6$

_1_	2	3	4	5	Ь
0.6	0.8	4.1	6.1	8.1	9.1
2.5	2.7	6	8	10	11
3.5	3.7	7	9	11	12
6.5	6.7	10	12	14	15
7.5	7.7	11	13	15	16
11.5	11.7	15	17	19	20
$\mathbf{u}_3$	$\mathbf{u}_4$	$\mathbf{u}_5$	$\mathbf{u}_2$	$\mathbf{u}_1$	$\mathbf{u}_6$

_1_	2	3	4	5	6
0.6	0.8	4.1	6.1	8.1	9.1
2.5	2.7	6	8	10	11
3-5	3.7	7	9	11	12
6.5	6.7	10	12	14	15
7.5	7.7	11	13	15	16
11.5	11.7	15	17	19	20
$\overline{\mathbf{u}_3}$	$\overline{\mathbf{u}_4}$	$\overline{\mathbf{u}_5}$	$\overline{\mathbf{u}}_2$	$\overline{\mathbf{u}_1}$	$\mathbf{u}_{\epsilon}$

$$(\mathbf{1},\mathbf{1}) o \mathbf{W_{3}}$$

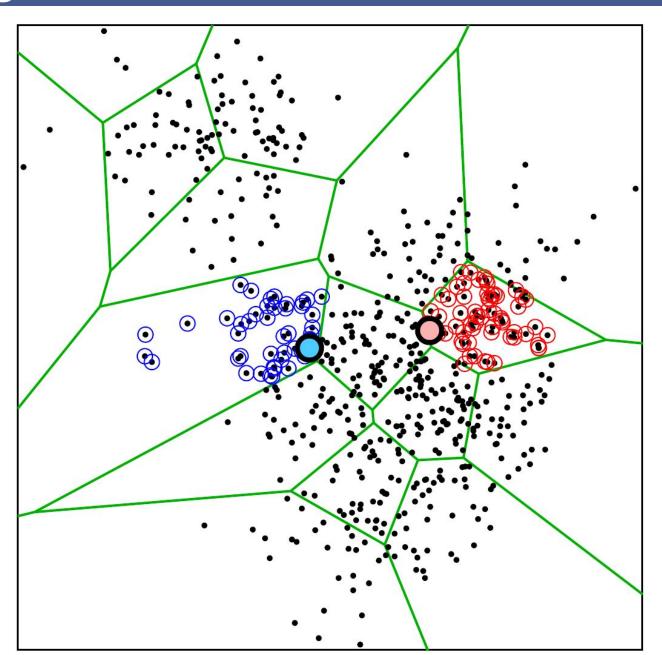
$$(\mathbf{2},\mathbf{1}) o \mathbf{W}_{\mathbf{4},\mathbf{2}}$$

$$(\mathbf{1},\mathbf{2}) o \mathbf{W_{3,3}}$$

$$(\mathbf{2},\mathbf{2}) o \mathbf{W_{4,3}}$$

$$({\bf 1},{\bf 1}) \to {\bf W_{3,4}} \quad ({\bf 2},{\bf 1}) \to {\bf W_{4,4}} \quad ({\bf 1},{\bf 2}) \to {\bf W_{3,3}} \quad ({\bf 2},{\bf 2}) \to {\bf W_{4,3}} \quad ({\bf 1},{\bf 3}) \to {\bf W_{3,5}}$$

# Querying the inverted multi-index



## **Experimental protocol**

#### **Dataset:**

- 1 billion of SIFT vectors [Jegou et al.]
- 2. Hold-out set of 10000 queries, for which Euclidean nearest neighbors are known

#### Comparing index and multi-index:

Set a candidate set length T

#### For each query:

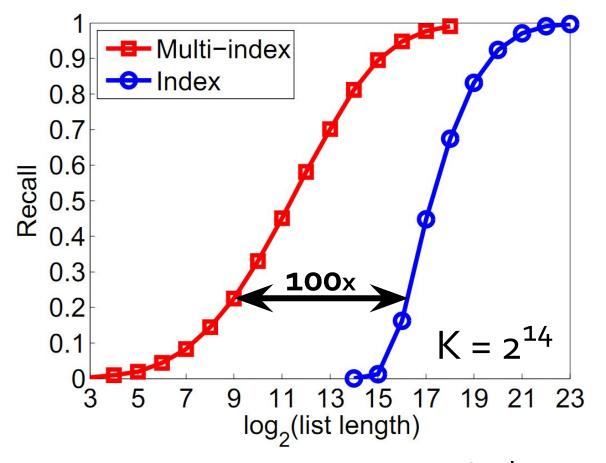
- Retrieve closest entries from index or multi-index and concatenate lists
- Stop when the next entry does not fit
  - ☐ For small T inverted index can return empty list
- Check whether the true neighbor is in the list

Report the share of queries where the neighbor was present (recall@T)

## Performance comparison

Recall on the dataset of 1 billion of visual descriptors:

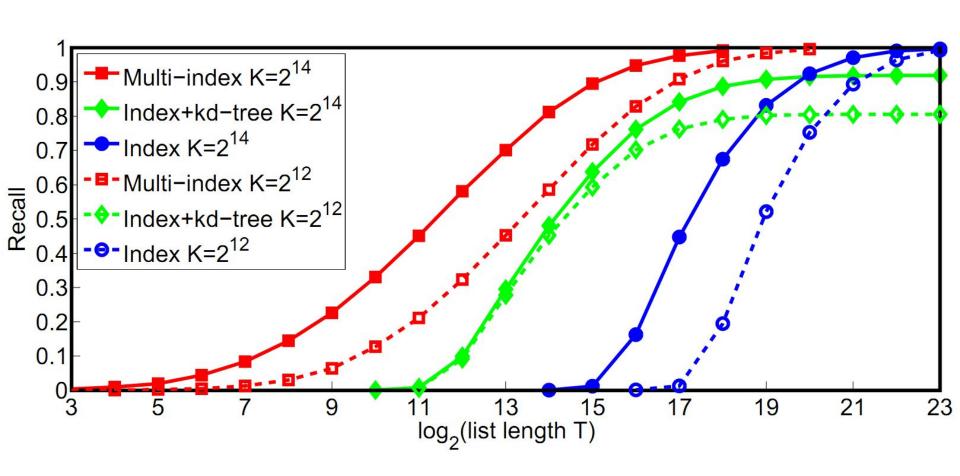
"How fast can we catch the nearest neighbor to the query?"



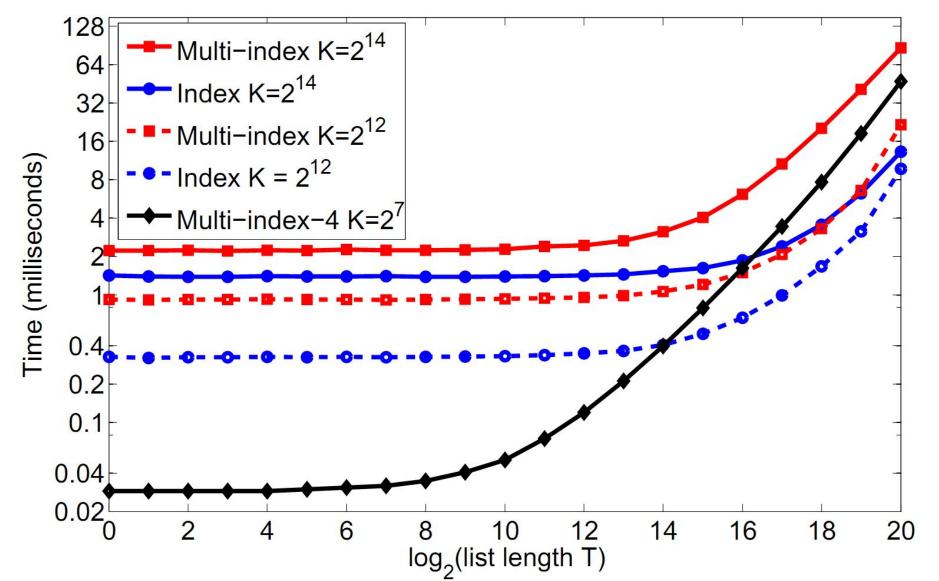
Time increase: 1.4 msec -> 2.2 msec on a single core (with BLAS instructions)

# Performance comparison

Recall on the dataset of 1 billion 128D visual descriptors:

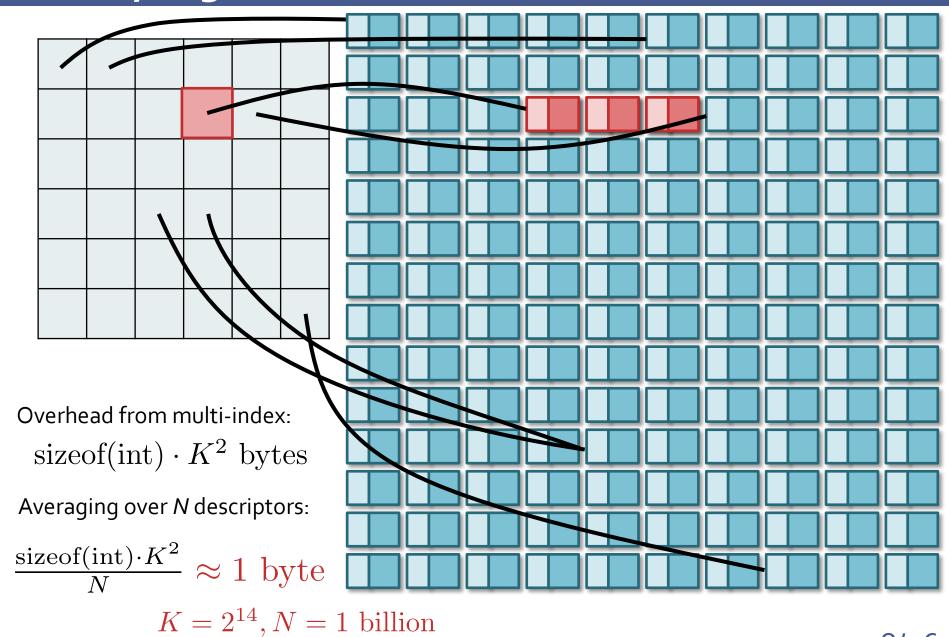


# Time complexity



For same K index gets a slight advantage because of BLAS instructions

# Memory organization



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# Why two?

For larger number of parts:

Memory overhead becomes larger

$$sizeof(int) \cdot K^2$$
 bytes sizeof(int)  $\cdot K^4$  bytes

 Population densities become even more non-uniform (multi-sequence algorithm has to work harder to accumulate the candidates)

In our experiments, 4 parts with small K=128 may be competitive for some datasets and reasonably short candidate lists (*e.g. duplicate search*). Indexing is blazingly fast in these cases!

## Multi-Index + Reranking

• "Multi-ADC": use *m* bytes to encode the original vector using product quantization

faster (efficient caching possible for distance computation)

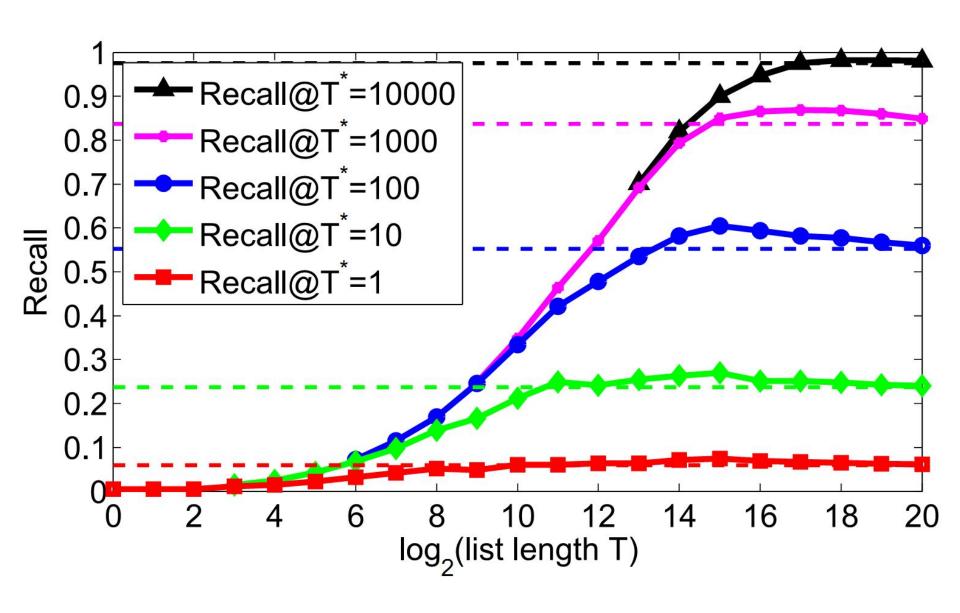
- "Multi-D-ADC": use m bytes to encode the remainder between the original vector and the centroid
  - ☐ Same architecture as IVFADC of Jegou et al., but replaces index with multi-index

more accurate

### Evaluation protocol:

- Query the inverted index for T candidates
- 2. Reconstruct the original points and rerank according to the distance to the query
- 3. Look whether the true nearest neighbor is within top  $T^*$

### Multi-ADC vs. Exhaustive search



## Multi-D-ADC vs State-of-the-art

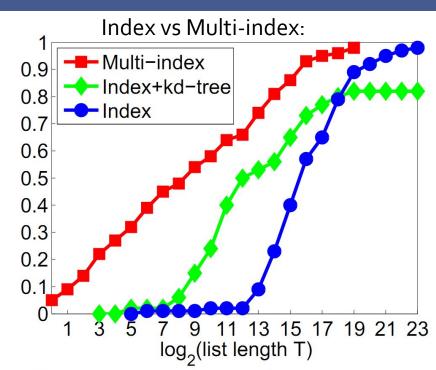
Combining multi-index + reranking:



System	List len. $T$	R@1	R@10	R@100	Time
BIGANN	N, 1 billion	SIFTs, 8	3 bytes p	per vecto	or
IVFADC	8 million	0.112	0.343	0.728	155
State-of-the-art	[Jegov et al.]	(0.088)	(0.372)	(0.733)	(74*)
Multi-D-ADC	10000	0.158	0.472	0.706	6
Multi-D-ADC	30000	0.164	0.506	0.813	13
Multi-D-ADC	100000	0.165	0.517	0.860	37

## Performance on 80 million GISTs

Same protocols as before, but on 80 million GISTs (384 dimensions) of Tiny Images [Torralba et al. PAMI'08]

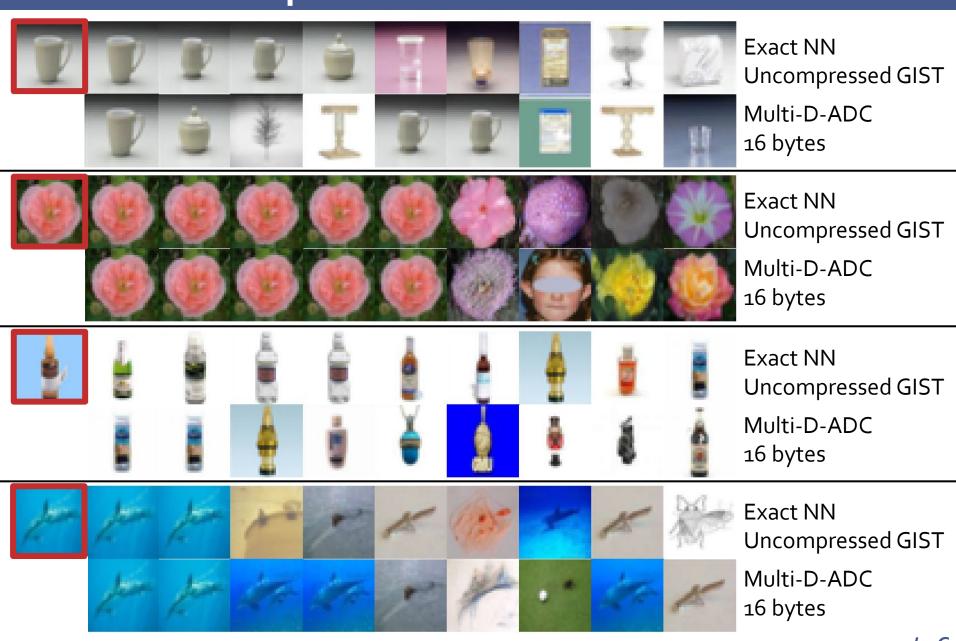


#### Multi-D-ADC performance:

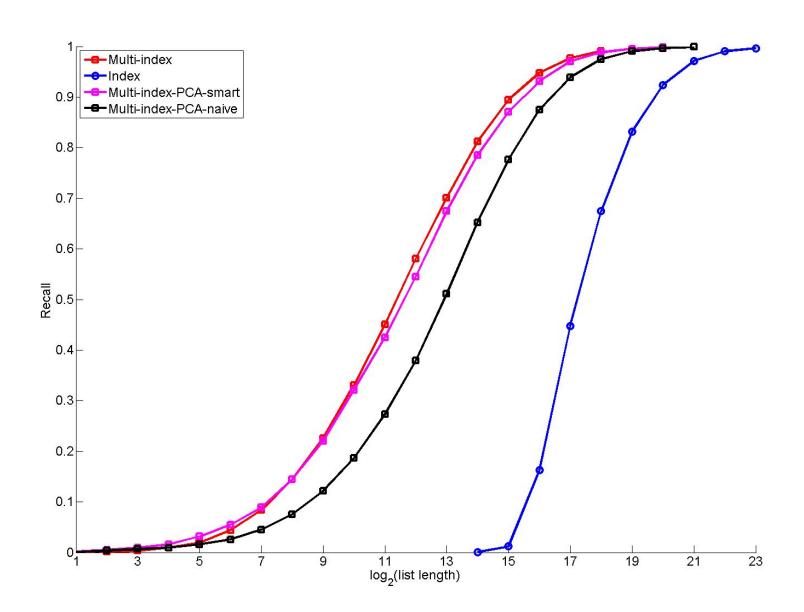
Tiny Images, 80 million GISTs, 8 bytes per vector							
Multi-D-ADC 10000 0.06 0.40 0.59 19							
Multi-D-ADC	30000	0.06	0.41	0.63	41		
Multi-D-ADC	100000	0.06	0.41	0.66	119		

Tiny Images, 80 million GISTs, 16 bytes per vector					
Multi-D-ADC	10000	0.06	0.49	0.64	19
Multi-D-ADC	30000	0.06	0.56	0.76	46
Multi-D-ADC	100000	0.06	0.56	0.85	139

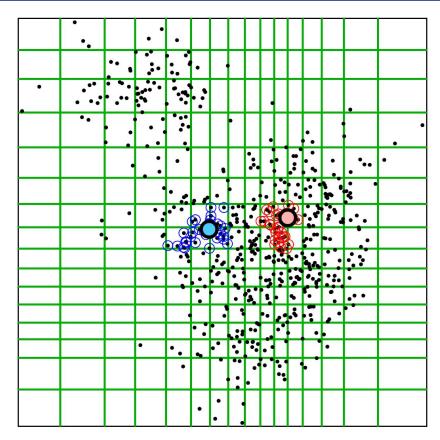
# Retrieval examples



# Multi-Index and PCA (128->32 dimensions)



## Conclusions



- A new data structure for indexing the visual descriptors
- Significant accuracy boost over the inverted index at the cost of the small memory overhead
- Code available (will soon be online)

## Other usage scenarios

#### (Mostly) straightforward extensions possible:

- Large-scale NN search' based approaches:
  - Holistic high dimensional image descriptors: GISTs, VLADs, Fisher vectors, classemes...
  - Pose descriptors
  - ☐ Other multimedia
- Additive norms and kernels: L1, Hamming, Mahalanobis, chi-square kernel, intersection kernel, etc.

## Visual search

What is this painting?





Collection of many millions of fine art images

The closest match:



Van Gogh, 1890
"Landscape with Carriage and Train in the Background"
Pushkin museum, Moscow
<u>Learn more about it</u>