

Gabriele Martino

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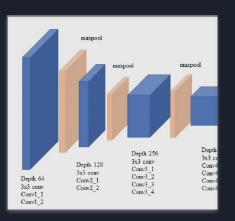
CNN

Index

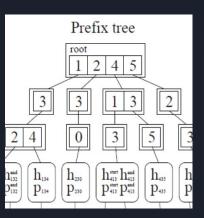
**Dataset** 

**Distractor Dataset** 

**VGG16, VGG19** 



**PPIndex** 



**Sketches** 



MrFlickr25k



### Pipeline

Features Extraction from pretrained CNN

Performance Analysis

Features
Extraction from
Fine Tuned CNN

Performance Analysis

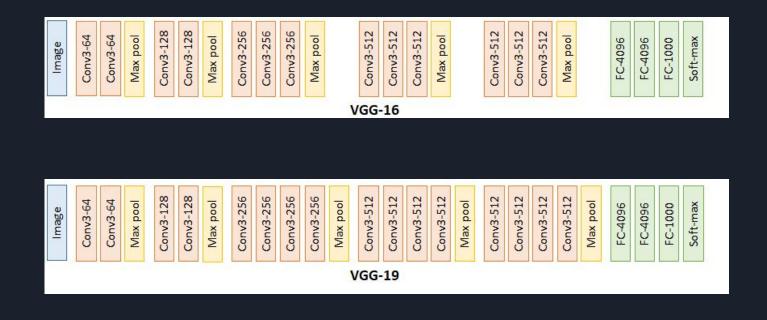
### Pipeline

Features Extraction from pretrained CNN

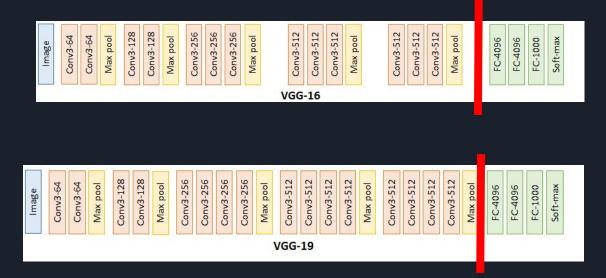
Performance Analysis Features
Extraction from
Fine Tuned CNN

Performance Analysis

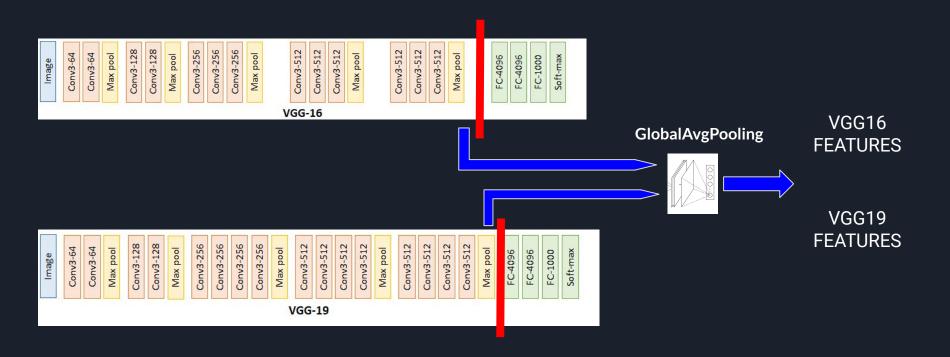
# Features Extraction from pretrained CNN (i)



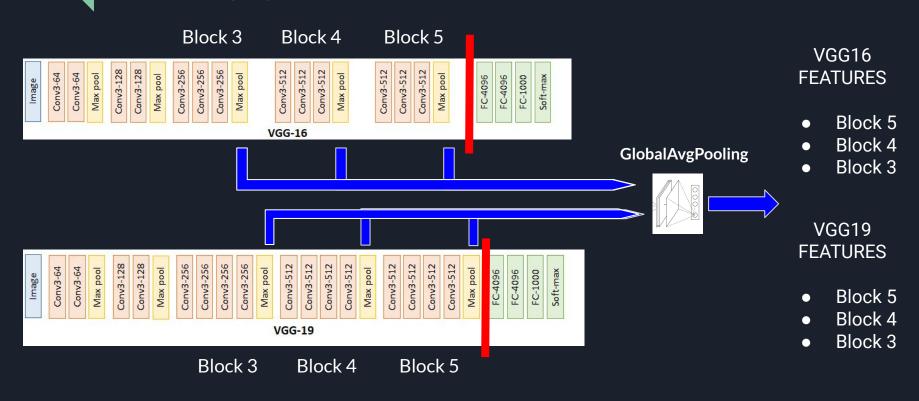
# Features Extraction from pretrained CNN (ii)



# Features Extraction from pretrained CNN (iii)



# Features Extraction from pretrained CNN (iv)



### Pipeline

Features Extraction from pretrained CNN

Performance Analysis Features Extraction from Fine Tuned CNN

Performance Analisis

### Pipeline

Features Extraction from pretrained CNN

Performance Analysis Features Extraction from Fine Tuned CNN

Performance Analysis

Exact Search on different features

Analysis on different number of Pivots

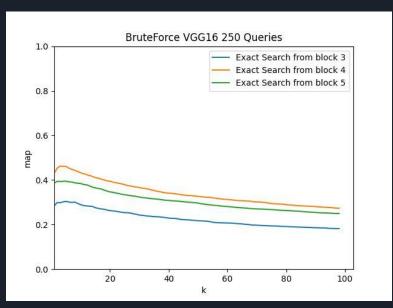
Analysis on Prefix Length

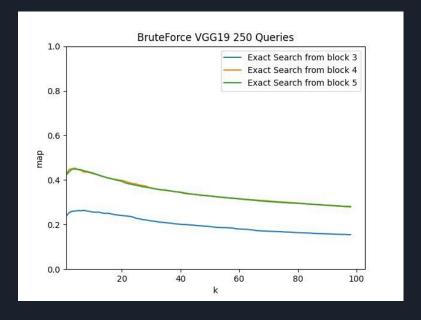
Analysis on Z

Analysis on pivot selection method

### Performance Analysis - Exact Search on different features

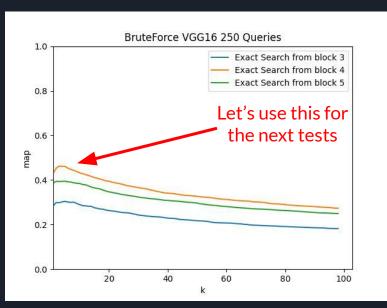
KNN search on query extracted from the dataset with a bucket of dataset + distractor. Queries are the same for all the analysis; 250, one for each class.

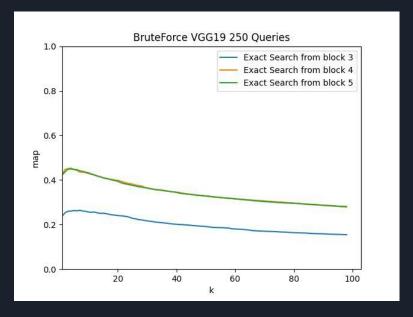




### Performance Analysis - Exact Search on different features

KNN search on query extracted from the dataset with a bucket of dataset + distractor. Queries are the same for all the analysis; 250, one for each class.





#### Performance Analysis with PPindex (ii)

#### PPIndex parameters:

- Number of pivots
- Pivots selection method
- **L**, prefix length
- **Z**, number of objects retrieved before k cutting

#### Performance Analysis with PPindex (iii)

#### Experiments:

- Number of pivots: Np∈{10, 50, 100, 200}
- Pivot selection method: random, 3Np method, Kmedoids
- [
- $Z = K + b, b \in \{20, 50, 100, 200\}$
- Same 250 queries, one for each class of the dataset, to make a real comparison
- K ∈[1,100]

#### Analysis of number of pivots (i)

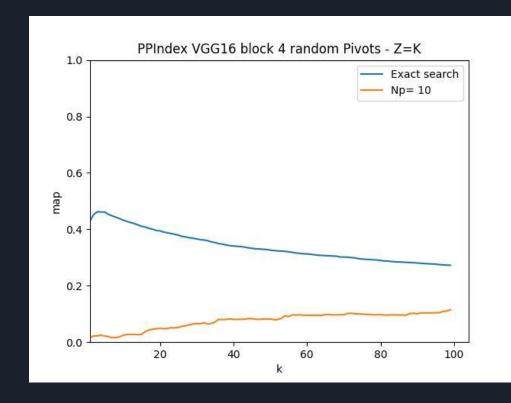
#### Analysis of number of pivots

Number of pivots: Np∈{10, 50, 100, 200}

Fixed Parameter:
Pivot Selection: Random

L = 10

Z = K



#### Analysis of number of pivots (ii)

#### Analysis of number of pivots

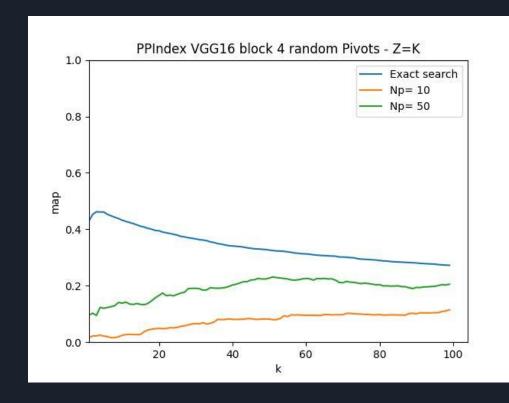
Number of pivots: Np∈{10, 50, 100, 200}

Fixed Parameter:

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L = 10

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#### Analysis of number of pivots (iii)

#### Analysis of number of pivots

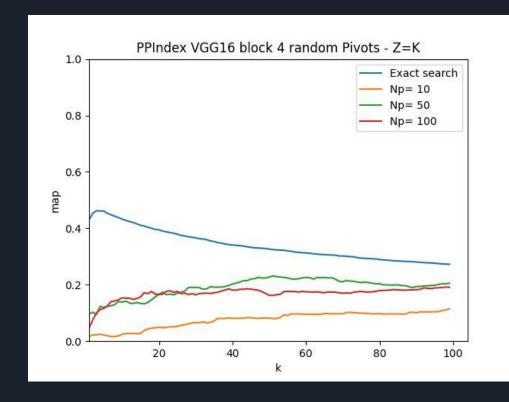
Number of pivots: Np∈{10, 50, 100, 200}

Fixed Parameter:

Pivot Selection: Random

L = 10

Z = k



### Analysis of number of pivots (iv)

Analysis of number of pivots

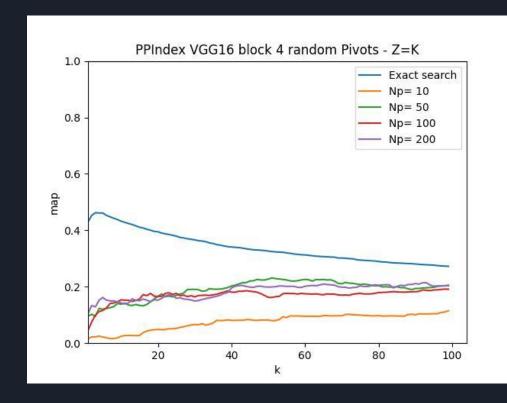
Number of pivots: Np∈{10, 50, 100, 200}

Fixed Parameter:

Pivot Selection: Random

L = 10

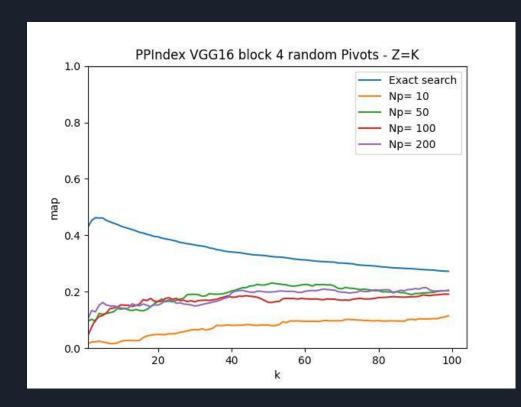
Z = K



#### Analysis of Z (i)

Let's vary Z properly using

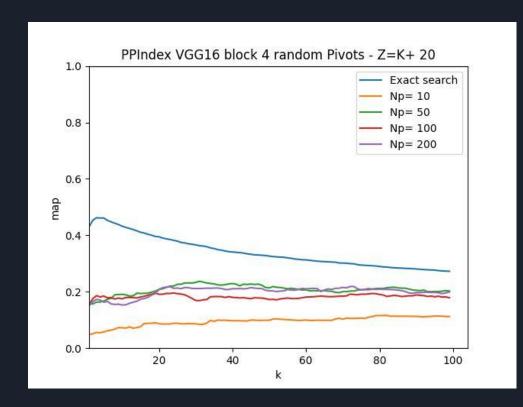
$$Z = k + b$$



#### Analysis of Z(ii)

Let's vary Z properly using

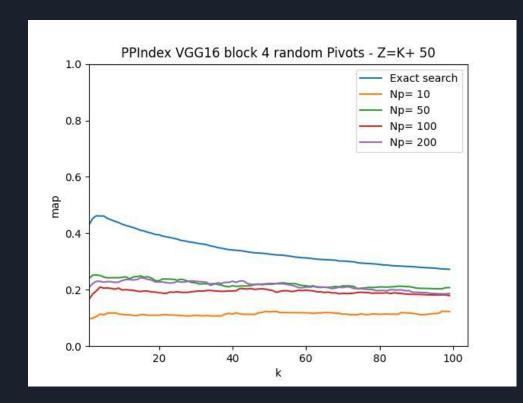
$$Z = k + b$$



#### Analysis of Z(iii)

Let's vary Z properly using

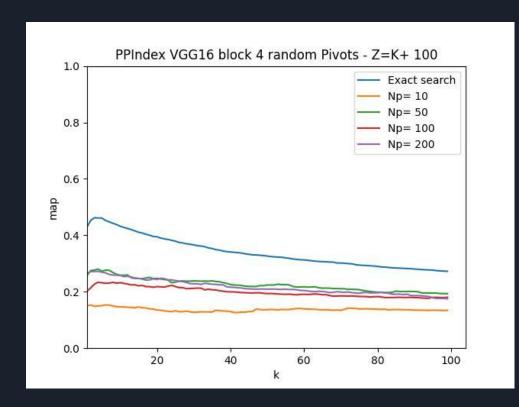
$$Z = k + b$$



#### Analysis of Z(iv)

Let's vary Z properly using

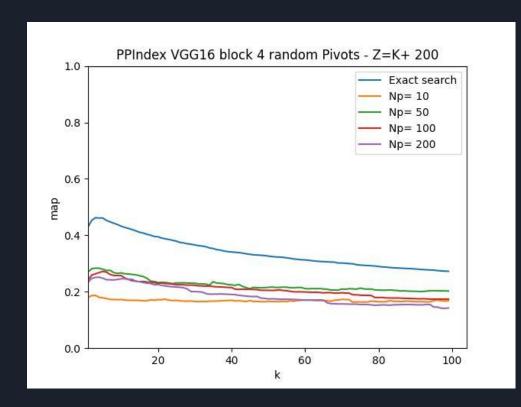
$$Z = k + b$$



#### Analysis of Z(v)

Let's vary Z properly using

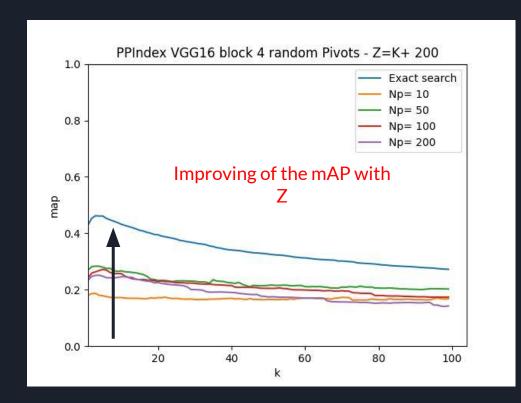
$$Z = k + b$$



### Analysis of Z(vi)

Let's vary Z properly using

$$Z = k + b$$



#### Analysis of Z - Note

Considering a dataset size of 45'000 (quite small) a low number of pivots is enough. For a small dataset the number of pivots affect a lot the results also because of the kind of search algorithm.

Given a k-NN query for an object  $q \in O$ , the basic search function of the PP-Index consists of computing the permutation prefix  $\Pi$  p and searching for the longest prefix match in the prefix tree whose subtree points to at least z candidate objects .[1]

This means that for a large number of pivots the objects are already spread a lot at the first level of the tree of the index. This could also lead to an impossibility to retrieve a certain number of objects.

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Example:

N = 45'000, Np = 250

N/Np = 180

(assuming a uniform distribution)

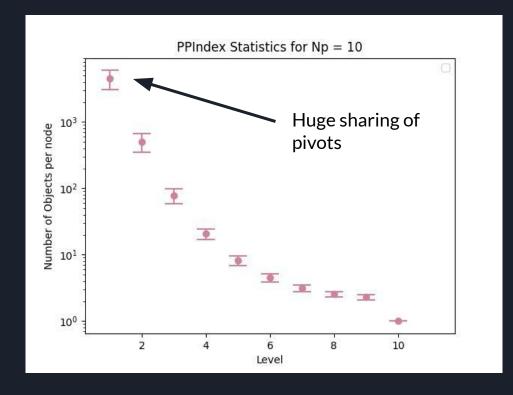
This means that at first level of the tree, each subtree contains around 180 object. Hence impossible to achieve Z = K + 200 for any K, leading to incorrect results. More on this later.

## Analysis of Prefix length and objects distribution (i)

xAxis = Tree level yAxis = objects per node of that level

Random pivots

The first value is exactly N/Np



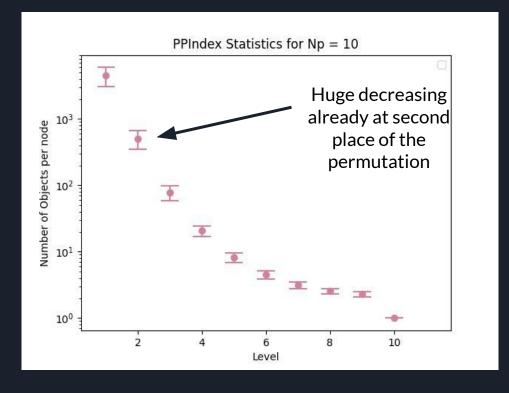
Confidence Interval at 95%

### Analysis of Prefix length and objects distribution (ii)

xAxis = Tree level yAxis = objects per node of that level

Random pivots

The first value is exactly N/Np

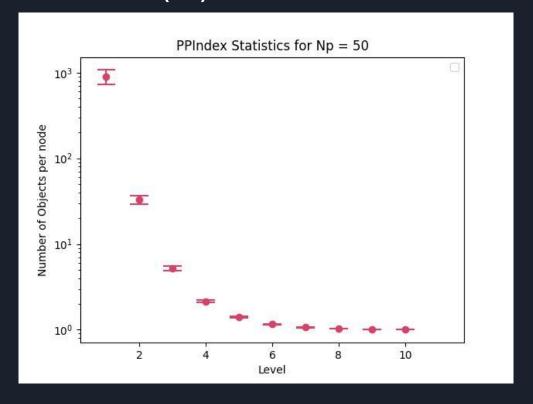


Confidence Interval at 95%

# Analysis of Prefix lenght and objects distribution (iii)

xAxis = Tree level yAxis = objects per node of that level

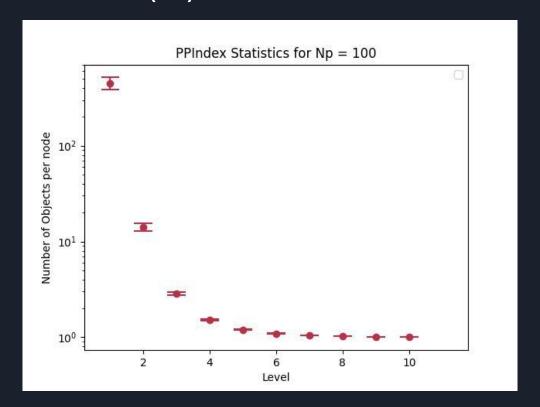
Random pivots



# Analysis of Prefix lenght and objects distribution(iv)

xAxis = Tree level yAxis = objects per node of that level

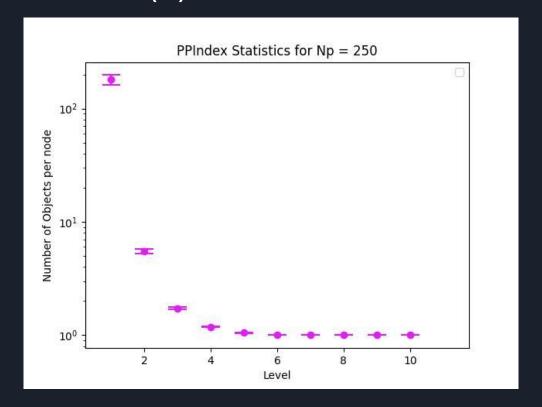
Random pivots



# Analysis of Prefix length and objects distribution(v)

xAxis = Tree level yAxis = objects per node of that level

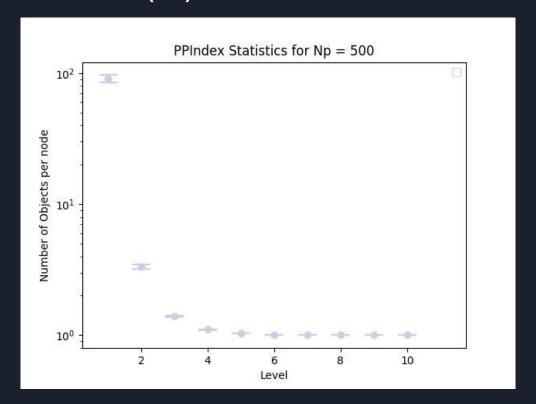
Random pivots



# Analysis of Prefix length and objects distribution(vi)

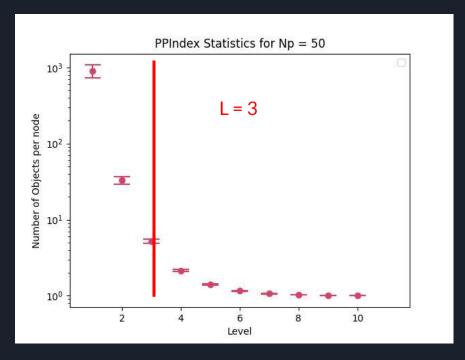
xAxis = Tree level yAxis = objects per node of that level

Random pivots

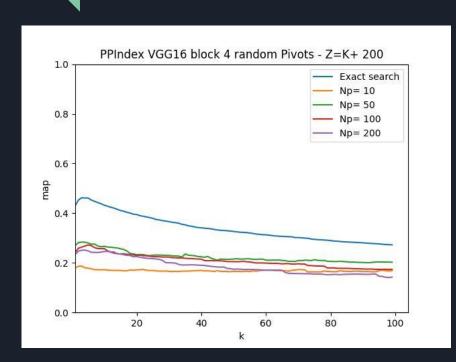


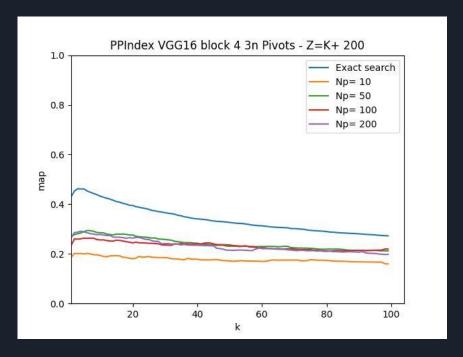
# Analysis of Prefix length and objects distribution(vi)

It's possible to extract some euristics on where to cut the prefix and so the depth of the tree, considering the exponential decreasing of the distribution of the object for each level of the tree.



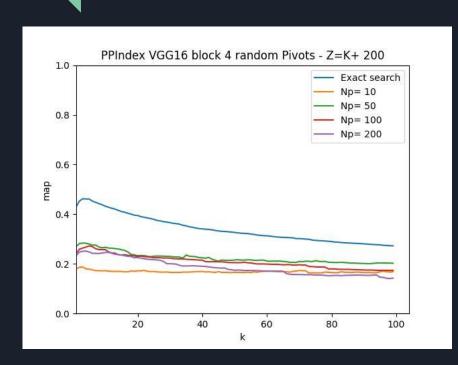
#### Pivot Selection Method Analysis (i)

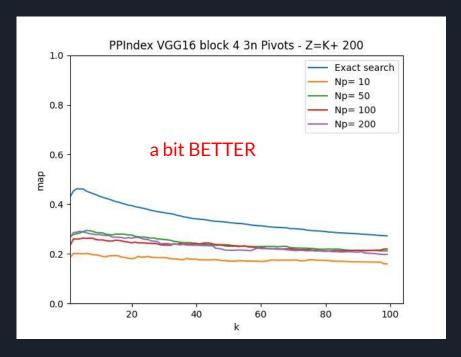




Random

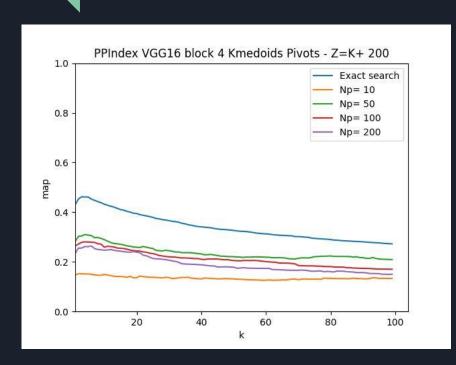
#### Pivot Selection Method Analysis (ii)

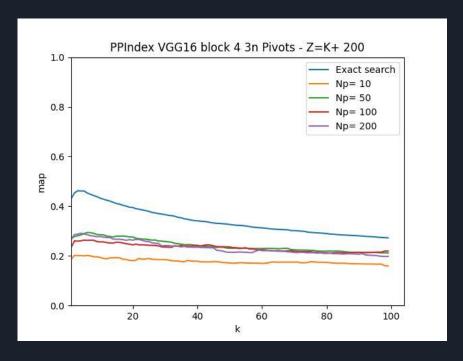




Random

#### Pivot Selection Method Analysis (iii)

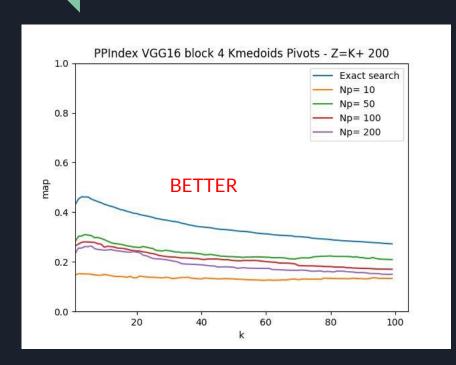


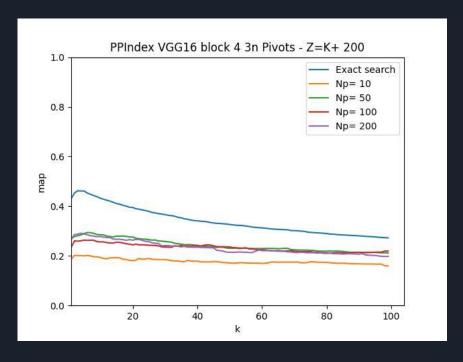


Kmedoids

3n

### Pivot Selection Method Analysis (iv)





**Kmedoids** 

Given a k-NN query for an object  $q \in O$ , the basic search function of the PP-Index consists of computing the permutation prefix  $\Pi$  p and searching for the longest prefix match in the prefix tree whose subtree points to at least z candidate objects .[1]

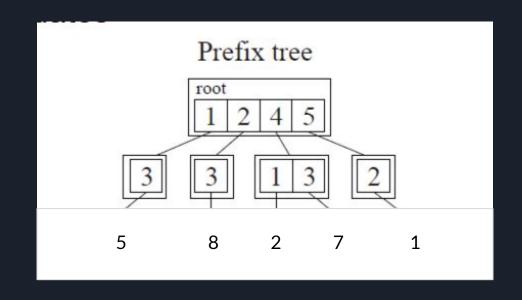
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Multiple query: at search time, p additional permutation prefixes from the query permutation prefix  $\Pi$ I q are generated, by swapping the position of some of its elements. The geometric rationale is that a permutation prefix  $\Pi$ I x differing from another permutation prefix  $\Pi$ I y for the swap of two adjacent/near elements identifies an area  $V\Pi$ I x of the similarity space adjacent/near to  $V\Pi$ I y. Performing a search with additional "swapped" permutation prefixes extends the search process to areas of the search space that are likely to contain relevant objects.

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Example:



q = [4, 1, 3, 5]

Z = 10

Object captured = 0

Example:

q = [4, 1, 3, 5]

Z = 10

Object captured = 0

Example:

Prefix tree root

q = [4, 1, 3, 5]

Z = 10

Object captured = 0

Example:

Take object inside

q = [4, 1, 3, 5]

Z = 10

Object captured = 2

Example:

q = [4, 1, 3, 5]

Z = 10

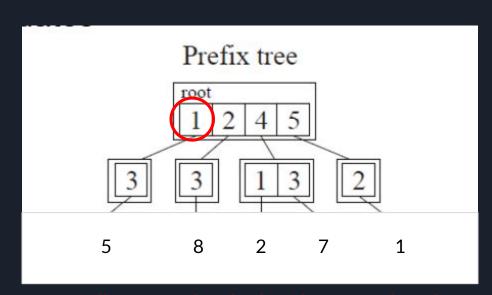
Object captured = 2 + 7

Number of objects under the node

Go to the sibling and take all the objects

Example:

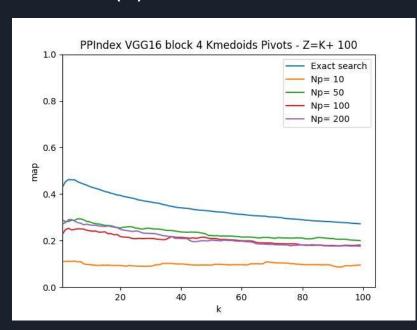
Number of objects under the node

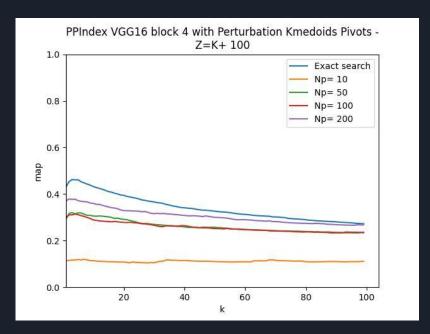


Still not enough go back to the root node and take element from the second best subtree and repeat the search

#### Small modifications to Algorithm:

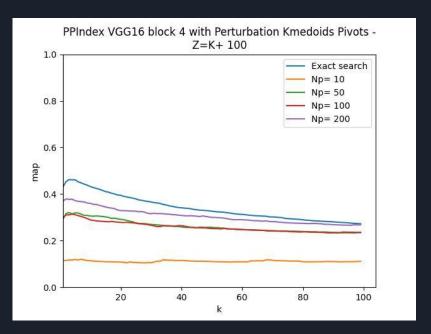
- Search the deeper matching subtree that contains at least Z candidates
- If the deeper matching subtree contains less than Z candidates, include also the candidate of the siblings of the last matching node (maybe better the siblings in matching order respect to the query)
- If the first matching node of the Prefix tree (root) does not contains enough objects ( < Z) search following the second element of the permutation of the query
- Don't cut the permutation of the query to L for the search, to allow to explore all the subtree in order of distance if necessary





This approach also overcome the problem of the increasing granularity for large number of pivots for a small dataset.

Here a greater number of pivots improve the performance



With Perturbation

It is possible to estimate the query time considering the amount of computation the algorithms does.

$$M*distanceComp + M*log\left(M\right) + log\left(L\right) + Retrieval + Z*distanceComp + Z*log\left(Z\right) \blacktriangleleft$$

Distances to pivots

M = Number of pivots

L = Prefix Lenght

It is possible to estimate the query time considering the amount of computation the algorithms does.

$$M*distanceComp + M*log\left(M\right) + log\left(L\right) + Retrieval + Z*distanceComp + Z*log\left(Z\right)$$

Ordering of Pivots for Permutations

M = Number of pivots

L = Prefix Lenght

It is possible to estimate the query time considering the amount of computation the algorithms does.

$$M*distanceComp + M*log\left(M\right) + log\left(L\right) + Retrieval + Z*distanceComp + Z*log\left(Z\right)$$

Searching in the Tree (Negligeble)

M = Number of pivots

L = Prefix Lenght

It is possible to estimate the query time considering the amount of computation the algorithms does.

$$M*distanceComp + M*log\left(M\right) + log\left(L\right) + Retrieval + Z*distanceComp + Z*log\left(Z\right)$$

Retrieval from disk

M = Number of pivots

L = Prefix Lenght

It is possible to estimate the query time considering the amount of computation the algorithms does.

$$M*distanceComp + M*log\left(M\right) + log\left(L\right) + Retrieval + Z*distanceComp + Z*log\left(Z\right)$$

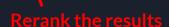
Real distance to 2 candidates

M = Number of pivots

L = Prefix Lenght

It is possible to estimate the query time considering the amount of computation the algorithms does.

$$M*distanceComp+M*log\left(M\right)+log\left(L\right)+Retrieval+Z*distanceComp+Z*log\left(Z\right)$$



M = Number of pivots

L = Prefix Lenght

To avoid using seconds as time evaluation because strictly depends on the machine, we used the adimensional time ratio, evaluating the Query Time using the PPindex and the Query Time using the NaiveSearch (Total Scan of the dataset).

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We used the Fieller's theorem, that finds the CI of a ratio of two random variables

Fieller showed that if  ${\it a}$  and  ${\it b}$  are (possibly correlated) means of two samples with expectations  $\mu_a$  and  $\mu_b$ , and variances  $\nu_{11}\sigma^2$  and  $\nu_{22}\sigma^2$  and covariance  $\nu_{12}\sigma^2$ , and if  $\nu_{11},\nu_{12},\nu_{22}$  are all known, then a (1 –  ${\it a}$ ) confidence interval ( $m_{\rm L},m_{\rm U}$ ) for  $\mu_a/\mu_b$  is given by

$$(m_L,m_U) = rac{1}{(1-g)} \left[ rac{a}{b} - rac{g
u_{12}}{
u_{22}} \mp rac{t_{r,lpha}s}{b} \sqrt{
u_{11} - 2rac{a}{b}
u_{12} + rac{a^2}{b^2}
u_{22} - g\left(
u_{11} - rac{
u_{12}^2}{
u_{22}}
ight)} 
ight]$$

where

$$g = \frac{t_{r,\alpha}^2 s^2 \nu_{22}}{b^2}$$

Here  $s^2$  is an unbiased estimator of  $\sigma^2$  based on r degrees of freedom, and  $t_{r,\alpha}$  is the  $\alpha$ -level deviate from the Student's t-distribution based on r degrees of freedom.

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 $IE = \frac{Cost(Q)}{Cost^{A}(Q)}$ 

then

where

$$g=rac{t_{r,lpha}^2s^2
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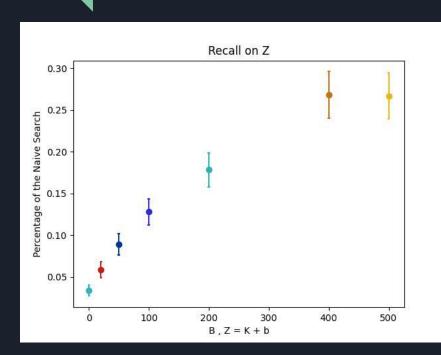
In our case the two random variable are the made with a query test of the same 250 Queries, from which we can extract the mean and the variance.

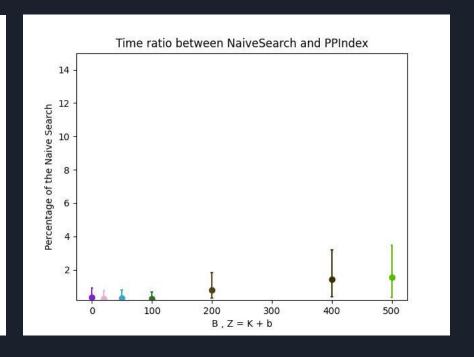
#### **Experiments Parameters:**

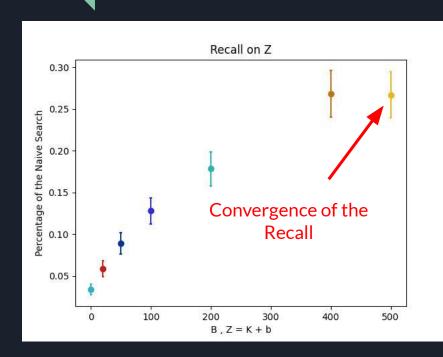
- Pivots Selection method: Kmedoids
- Number of Pivots: 50
- K = 20
- I=3
- All the dataset is in memory ( since the machine has and SSD the linear scanning of the reordered blocks could not be a lot visible)

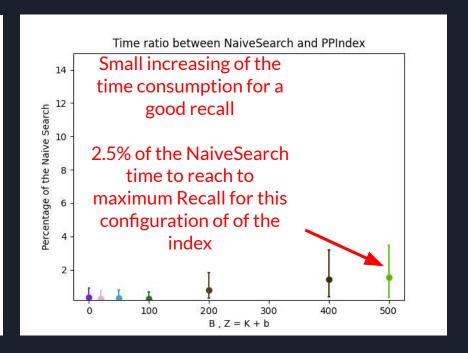
#### **Evaluation metrics:**

- Ratio of the Query Time means
- Recall between the retrieved set (regardless the ordering)

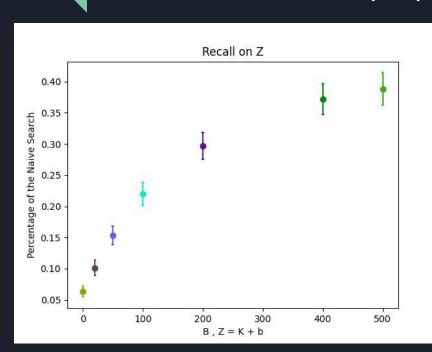


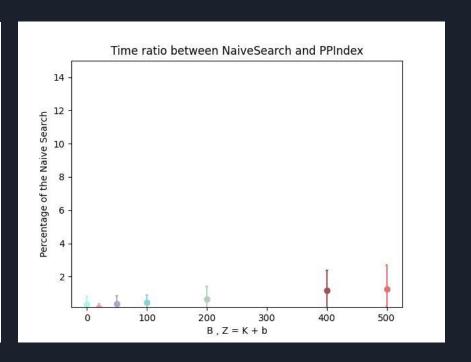






### Query Time Analysis - With Perturbation (vii)





With perturbation search with almost no increasing of the time

#### Conclusions

- Best pivots selection method: **Kmedoids**
- Number of Pivots: **around 0.5% of the size of the dataset**
- Use perturbation in the search to better explore the space
- Cut L to small value considering an exponential decreasing of the object for each level

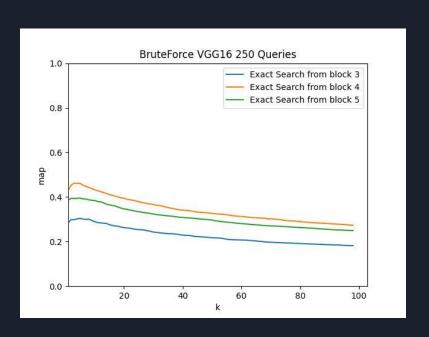
### Pipeline

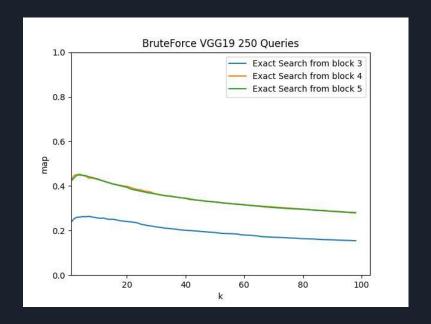
Features Extraction from pretrained CNN

Performance Analysis Features Extraction from Fine Tuned CNN

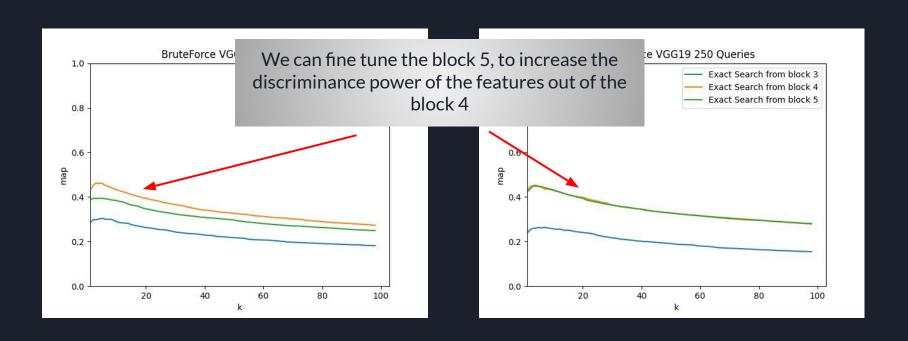
Performance Analysis

### Features Extraction from Fine Tuned CNN (i)

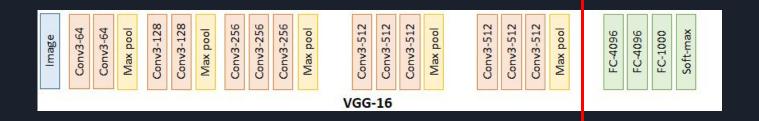




### Features Extraction from Fine Tuned CNN (i)



1) Cut the FC layers and put custom ones



1) Cut the FC layers and put custom ones

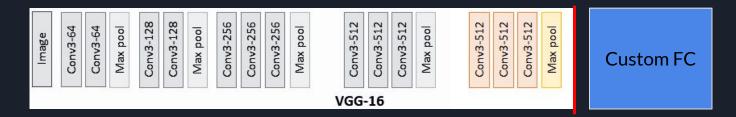


**Custom FC** 

- GlobalAveragePooling out of the CNN (512 values)
- BatchNormalization layer
- 1 Hidden layer with 1024 neurons
- Dropout
- Data augmentation for the training set (training = 0.8, validation = 0.2)
- 250 classes

Dataset of 20'000 samples

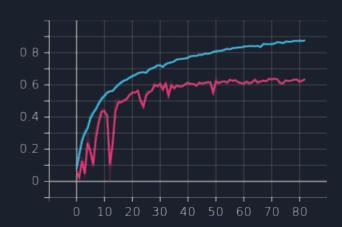
2) Freeze all the CNN and Train only the Custom FC and the final block



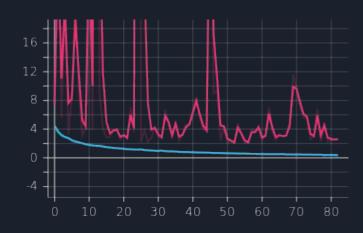
Freezed To train

Actually we should first train only the Custom FC layers, then unfreezing the last block of the VGG and using a smaller learning rate to not the destroy totally the pretrained weights, but considering that has been shown that the last layer is not so useful for the features, that our machine is powerful enough for the training from sckretch of that amount of parameters, we trained totally the last part of the network.

2) Freeze all the CNN and Train only the Custom FC and the final block. Results

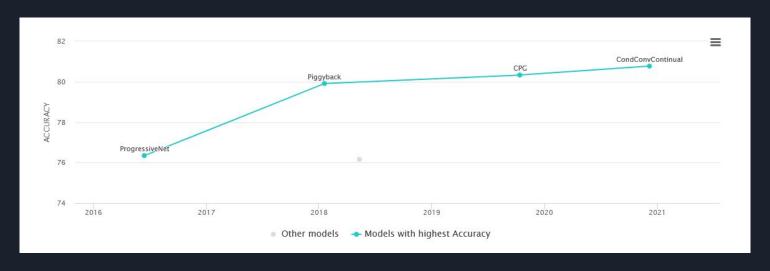


Epoch Accuracy Test accuracy: 0.62



Epoch Loss Test loss: 2.04

Not to bad considering the state of the art of the classification on this dataset



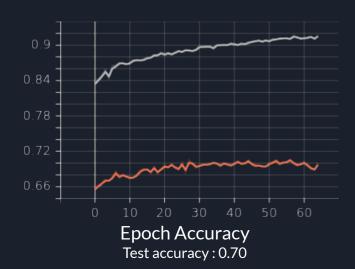
We can do BETTER!

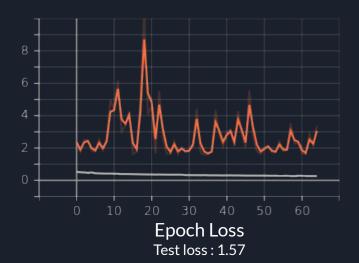
3) Unfreeze part of the CNN that we want to fine tune and retrain with a smaller learning rate



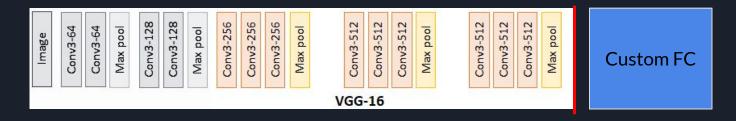
Freezed To Fine Tune

3) Unfreeze part of the CNN that we want to fine tune and retrain with a smaller learning rate. Results:



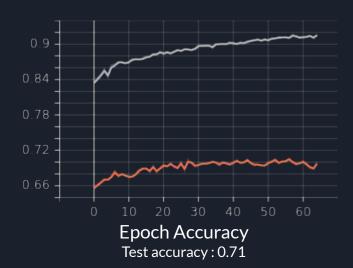


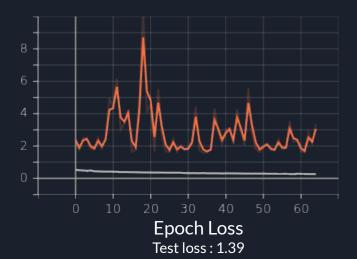
3) Unfreeze part of the CNN that we want to fine tune and retrain with a smaller learning rate



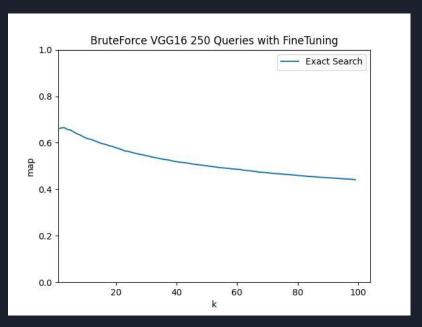
Freezed To Fine Tune

2) Freeze all the CNN and Train only the Custom FC





3) Extract Features from the CNN and evaluate mAP



### Pipeline

Features Extraction from pretrained CNN

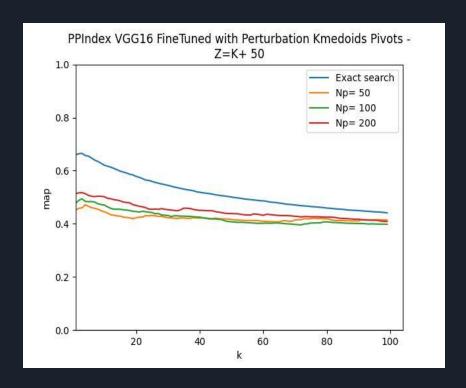
Performance Analysis Features Extraction from Fine Tuned CNN

Performance Analysis

#### Performance Analysis Fine Tuning

#### HyperParameter:

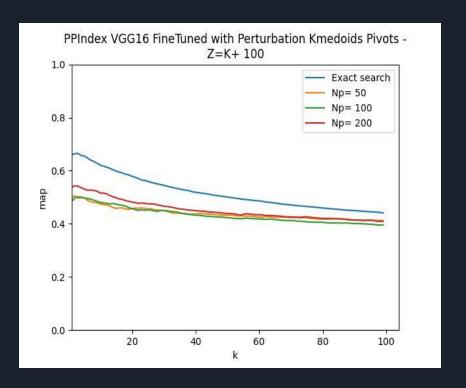
- Pivot Selection Method:
   Kmedoids
- Number of Pivots: {50, 100, 200}
- $Z = K + \{50, 100, 200\}$



#### Performance Analysis Fine Tuning

#### HyperParameter:

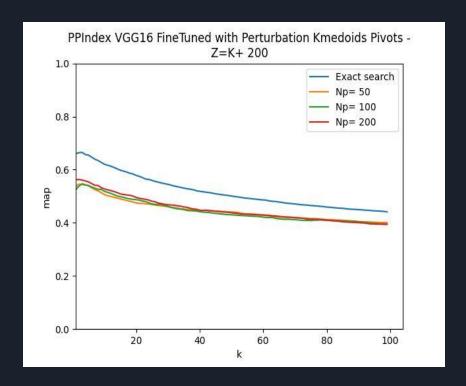
- Pivot Selection Method: Kmedoids
- Number of Pivots: {50, 100, 200}
- $Z = K + \{50, 100, 200\}$



#### Performance Analysis Fine Tuning

#### HyperParameter:

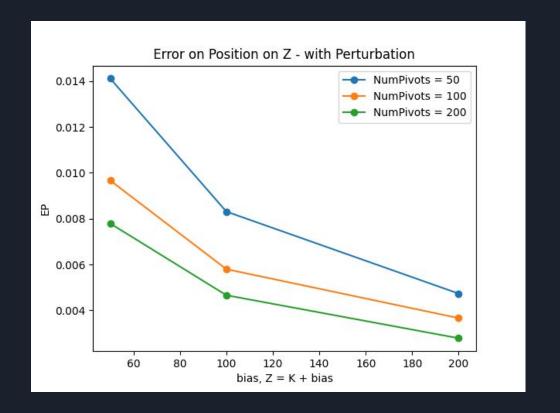
- Pivot Selection Method:
   Kmedoids
- Number of Pivots: {50, 100, 200}
- Z = K + {50, 100, 200}



#### Performance Analysis Fine Tuning - EP

#### Hyperparameters:

- Selection Method: Kmedoids
- L = 3
- Fine Tuned Features



The End

#### References

- PP-Index: Using Permutation Prefixes for Efficient and Scalable Similarity Search Andrea Esuli Istituto di Scienza e Tecnologie dell'Informazione CNR
- <u>https://keras.io/api/applications/</u>
- Fieller EC: The biological standardization of Insulin. Suppl to J R Statist Soc 1940, 7:1-64.
- <u>https://blog.keras.io/building-powerful-image-classification-models-using-very-little-data.html</u>
- https://paperswithcode.com/dataset/sketch
- Very Deep Convolutional Networks for Large-Scale Image Recognition <u>Karen Simonyan</u>, <u>Andrew Zisserman</u>
- The many Facets of Approximate Similarity Search Marco Patella Paolo Ciaccia