

Question :-

- We have a collection of N images $[1000 \times 1000]$ pixels.
- We are given a "query" image $[1000 \times 1000]$.
- Goal: quickly find out the "closest" image to the query image.

How?

High level idea

- Transform each image into a vector in $1000^2 = 10^6$ dimensions.

$$\begin{matrix} d \\ d \end{matrix} \quad \begin{matrix} 0 & .2 & .1 \\ .15 & .7 & .8 \\ 0 & .5 & .4 \end{matrix} \quad \xrightarrow{\quad D = d^2 \quad} \quad (0, .2, .1, .15, .7, .8, 0, .5, .4)$$

- Same for query image

- Output the nearest vector to the query vector.
(say l_2 distance)

Given N vectors $v_1, v_2, \dots, v_N \in \mathbb{R}^d$

1"

and given query $q \in \mathbb{R}^D$
find $\underset{i=1}{\overset{N}{\operatorname{arg\min}}} \|v_i - q\|_2^2$

(Approx.)

NEAREST NEIGHBOR SEARCH

Gives N "base" vectors in D dimensions,
pre-process and build a data
structure.

so that when queries are presented,
we can quickly retrieve the
(approximate) closest vector to the query

Naive Solution

when query is presented, compare
distances to all base points
and output the closest.

operations per query : $3 \cdot N \cdot D + N$
 \downarrow
find min.

D subtractions,
Multiplications
Additions per point

In our example

$$D = 10^6$$

Today's lecture

generic "dimension reduction" technique
 which can bring down $10^6 \rightsquigarrow 10^3$
 with very little "error".

JOHNSON-LINDENSTRAUSS LEMMA

Given n vectors in \mathbb{R}^d , v_1, v_2, \dots, v_n
 there is a mapping $f: \mathbb{R}^d \rightarrow \mathbb{R}^k$

such that

$$\forall u_i, u_j$$

$$(1 - \varepsilon) \leq \frac{\|f(u_i) - f(u_j)\|_2}{\|u_i - u_j\|_2} \leq (1 + \varepsilon)$$

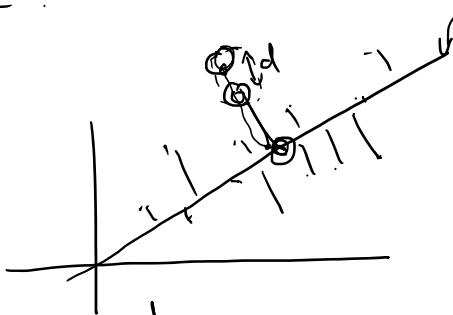
with $k = \Theta\left(\frac{\log n}{\varepsilon^2}\right)$.

Moreover we can find this efficiently
 with good probability

Goal for today

Ideas: PCA ?

- Unclear
- It's more of a
 - global error reduction



- we want pairwise preservation

Idea 2: what about just picking
k random coordinates?

$v = (x_1, x_2, \dots, x_d)$ then $f(v) = (x_{i_1}, x_{i_2}, \dots, x_{i_k})$
where i_1, i_2, \dots, i_k are
randomly chosen from $[d]$.

Same i_1, i_2, \dots, i_k are used for all vectors.

[sample with replacement, to make
analysis easy]

Good News

for any u and $v \in \mathbb{R}^d$

$$E[(\|f(u) - f(v)\|)^2] = \frac{k}{d} \|u - v\|^2$$

good, b/c then we can think of

$$\hat{f}(u) = \sqrt{\frac{d}{k}} (u_{i_1}, u_{i_2}, \dots, u_{i_k})$$

and then we'll have

$$E[(\|\hat{f}(u) - \hat{f}(v)\|)^2] = \|u - v\|^2$$

Proof of Good News

Fix u, v and let $z = u - v$

$$\|z\|^2 = \sum_{i=1}^d (u_i - v_i)^2 = \sum_{i=1}^d z_i^2$$

Now, consider i_1 and focus on

$$\begin{aligned} E_{i_1}[(u_{i_1} - v_{i_1})^2] &= \sum_{l=1}^d P_{i_1}(i_1 = l) \cdot (u_l - v_l)^2 \\ &= \frac{1}{d} \cdot \sum_{l=1}^d (u_l - v_l)^2 \\ &= \frac{1}{d} \|u - v\|_2^2 \end{aligned}$$

$$\Rightarrow E \left[\|f(u) - f(v)\|_2^2 \right] = \sum_{l=1}^k E \left[(u_{i_l} - v_{i_l})^2 \right] = \frac{k}{d} \|u - v\|_2^2$$

If we can show some "concentration inequality" that $\|f(u) - f(v)\|$ is close to its average whp, then this scheme works!

and then union bound for all $1 \leq i < j \leq n$.

Q: Is this likely to have concentration?

A: Sadly, no

Here's an example :-

$$v_1 = (1, 0, 0, \dots, 0)$$

$$v_2 = (0, 1, 0, \dots, 0)$$

$$v_n = (0, 0, \dots, 1)$$

Unless we pick one of 1 or 2
coordinate is 0,

$$\|f(v_1) - f(v_2)\| = 0$$

\Rightarrow even if $k < n-1$, we will
always make huge
error on some pair

$$\text{real distance} = \sqrt{2} \quad \}$$

$$\text{"our imagined" distance} = 0 \quad \}$$

badness because most coordinates are
same, few coords give all
the distance -

How to "spread" this over all coords?

TRICK:

- Nothing sacred about standard basis
choose a random rotation $R \in \mathbb{R}^{d \times d}$

first rotate the data

$$v_i \rightarrow R v_i \quad (\text{still in } d\text{-dim})$$

then sample k words from $R v_i$

- Intuition, can be made rigorous

- Can also be simplified greatly using Gaussians

$$f(u) = \lambda \begin{bmatrix} g_{11} & g_{12} & \dots & g_{1d} \\ \vdots & & & \\ g_{k1} & g_{k2} & \dots & g_{kd} \end{bmatrix} \begin{bmatrix} u_1 \\ u_2 \\ \vdots \\ u_d \end{bmatrix}$$

G

04/04/2021

$$f(u) = \lambda \begin{bmatrix} g_{11} & g_{12} & \dots & g_{1d} \\ g_{21} & g_{22} & \dots & g_{2d} \\ \vdots & & & \\ g_{k1} & g_{k2} & \dots & g_{kd} \end{bmatrix} \begin{bmatrix} u_1 \\ u_2 \\ \vdots \\ u_d \end{bmatrix}$$

G_r = random $k \times d$ gaussian matrix
 where each g_{ij} is an
 independent gaussian r.v
 with mean 0 & variance 1.

(e) $g_{ij} \sim N(0, 1)$.
 λ will be chosen suitably later.

THEOREM

For suitable λ and k , this mapping
 preserves all pairs of distances
 upto $(\pm \varepsilon)$ factor.

Q How do we prove this?

We'll fix a pair of vectors u_i, u_j
 and let $z = u_i - u_j$

Next we show that

$$\text{By random choice of } G_r \left[\frac{\|f(u_i) - f(u_j)\|_2^2}{\|u_i - u_j\|_2^2} \notin (1-\varepsilon, 1+\varepsilon) \right] \leq \frac{1}{n^2}$$

Then we just do union bound over all
 pairs u_i, u_j .

$$\Pr_{G_1} \left[\exists i, j \text{ which is not preserved} \xrightarrow{\text{to } (1 \pm \varepsilon)} \right] \leq \frac{\binom{n}{2}}{n^2} \leq \frac{1}{2}$$

With prob $\frac{1}{2}$, experiment was successful
(ie) all pairs preserved !

If not, just repeat 😊

Need to show that a single pair is preserved w.h.p.

Fix u & v and let $z = u - v$

$$f(u) = \lambda \cdot G_1 \cdot u$$

$$f(v) = \lambda \cdot G_1 \cdot v$$

$$f(u) - f(v) = \lambda \cdot G_1(u - v) = \lambda G_1 z = f(z)$$

Need to show that

$$1 - \varepsilon \leq \frac{\|f(z)\|_2^2}{\|z\|_2^2} \leq 1 + \varepsilon \quad \text{with prob} \geq 1 - \frac{1}{n^2}$$

let $z = (z_1, z_2, \dots, z_d)$.

and consider $\hat{z} = \frac{z}{\|z\|_2}$

$$\text{Then } z = \|z\|_2 \cdot \hat{z}$$

Moreover,

$$f(z) = \lambda \cdot G \circ z = \lambda \|z\| \cdot G \hat{z}$$

$$= \|z\| \cdot f(\hat{z})$$

w.p. $\gamma, 1 - \frac{1}{n^2}$

So, it suffices to show that

$$(1-\varepsilon) \leq \|f(\hat{z})\|_2^2 \leq (1+\varepsilon)$$

$$\Rightarrow \|z\|^2(1-\varepsilon) \leq \|f(z)\|_2^2 \leq (1+\varepsilon) \|z\|^2$$

Consider $\hat{z} = (\hat{z}_1, \hat{z}_2, \dots, \hat{z}_d)$

We know $\|\hat{z}\|_2^2 = 1 \Rightarrow \sum \hat{z}_i^2 = 1$

Let's look at $f(\hat{z}) = \lambda \cdot G \circ \hat{z}$

$$= \lambda \begin{bmatrix} \sum_{j=1}^d g_{1j} \hat{z}_j \\ \vdots \\ \sum_{j=1}^d g_{2j} \hat{z}_j \\ \vdots \\ \sum_{j=1}^d g_{kj} \hat{z}_j \end{bmatrix}$$

Each entry in $f(\hat{z})$ looks like

$$\lambda \cdot N(0, \sum_{j=1}^d \hat{z}_j^2) = \lambda N(0, 1).$$

Moreover because g_{ij} and $g_{ij'}$ are independent for all i, j, j' ,

independent for all
 i, j, i', j' ,
 these entries are themselves independent!

$$f(\hat{z}) = \lambda \begin{pmatrix} F_1 \\ F_2 \\ \vdots \\ F_k \end{pmatrix} \quad \text{where } F_i \sim N(0, 1).$$

$$\text{What is } E[\|f(\hat{z})\|^2] = \lambda \sum_{i=1}^k E[F_i]^2 = \lambda k.$$

We want $f(\hat{z})$ to be close to 1.

$$\Rightarrow \text{set } \boxed{\lambda = \frac{1}{k}}$$

$$\Rightarrow E[\underbrace{\|f(\hat{z})\|^2}_{}] = 1.$$

Let $\gamma = k \cdot \|f(\hat{z})\|^2$ be the random variable.

$$\gamma = \sum_{i=1}^k F_i^2$$

γ is the sum of many independent random variables.
 Moreover each F_i is a gaussian $N(0, 1)$

$\boxed{\chi^2 - \text{distribution}}$

'n - margin'

{ γ actually sharply concentrates around its mean,
 Try to prove this using ideas from Chernoff bounds proof

More or less,
 γ behaves like $N(\mu(\gamma), \sigma^2(\gamma))$

$$E[\gamma] = \sum E[F_i^2] = k$$

$$\text{Var}[\gamma] = \sum_{i=1}^k \text{Var}[F_i^2]$$

$$\text{Var}[F_i^2] = E[F_i^4] - E[F_i^2]^2$$

$$= \underbrace{E[F_i^4]}_{\text{constant}} - 1.$$

$$\left\{ \begin{array}{l} E[F_i^4] = O(1) \text{ [constant]} \\ \text{for gaussian distribution.} \end{array} \right.$$

$$\Rightarrow \text{Var}[\gamma] = ck \text{ for constant } c.$$

{ γ behaves like $N(k, ck)$. }
 ↓
 very big cheat, crude approx.
 Basically, we'll use Chernoff-like
 11 bounds on γ .

tail bounds on γ .

$$\Pr \left[|\gamma - E[\gamma]| \geq t \right] \leq \exp \left(-\frac{t^2}{\sigma^2} \right)$$

Can prove this type of inequality for γ .

$$E[\gamma] = k$$

$$\sigma^2(\gamma) = Ck.$$

$$\text{Set } t = \varepsilon \cdot k = \varepsilon E[\gamma]$$

$$\Pr \left[\gamma \notin (1-\varepsilon, 1+\varepsilon) E[\gamma] \right] \leq \exp \left(-\frac{\varepsilon^2 k}{Ck} \right) = \exp \left(-\frac{\varepsilon^2}{C} \right).$$

So just set $k = \frac{2C \cdot \ln n}{\varepsilon^2}$

$$\Pr (\text{bad event}) \leq \exp \left(-\frac{\varepsilon^2 \cdot 2C \ln n}{C \varepsilon^2} \right) = \frac{1}{n^2}.$$

Whenever γ lies $\in [1-\varepsilon, 1+\varepsilon] E(\gamma)$

$$\Leftrightarrow f(\hat{\gamma}) = \frac{1}{k} \gamma \text{ lies in } [1-\varepsilon, 1+\varepsilon].$$

$\Leftrightarrow f(\hat{z}) = \frac{1}{k}y$ lies in $[1-\varepsilon, 1+\varepsilon]$.

$\Rightarrow \Pr(f(\hat{z}) \notin [1-\varepsilon, 1+\varepsilon]) \leq \frac{1}{n^2}$ for
 $k = \frac{2C\ln n}{\varepsilon^2}$
and $\lambda = \frac{1}{k}$.

(c, r) - Approximate Near Neighbor Search .

Motivation

- Again vectors in high dimensional space
- look to retrieve "closest" vector to given query vector .

Concrete Example :-

N documents, N is very large
(all in english)

represented as a $\{0, 1\}^D$ vector for
 D very large . Ordered as

$D = \text{all distinct english words } \{w_1, w_2, \dots, w_D\}$

$$\theta = (v_1, v_2, \dots, v_D)$$

where $v_i = 0$ if i^{th} word is
not in document
= 1 if i^{th} word is in doc.

Goal : build a data structure / Algorithm

At when query vector q arrives, we quickly retrieve any nearby base vector.

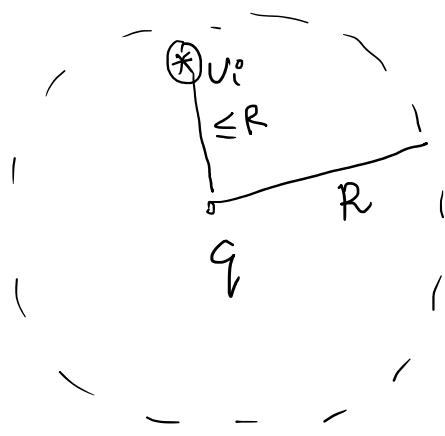
$$q = (q_1, q_2, \dots, q_D)$$

Given N base vectors v_1, v_2, \dots, v_N and $q = (q_1, q_2, \dots, q_D)$ arrives, and given parameter R , find any base vector which are at distance $\leq R$ from query q , if they exist.

Distance = Hamming distance

$$= \|u - v\|_1 = \sum_{j=1}^D |u_j - v_j|$$

= # indices where they differ



Naive approach

- Compare distances to all base vectors
Output the closest.

Query Time = $O(ND)$

Space of data structure = ND bits
(store all base vectors).

ΘN :

Can we get sub-linear dependence on N?

e.g., even if we bring query time

from ND to $\sqrt{N} \cdot D$, that is
really significant in real life

N can be $10^9, 10^{12}$ even !!

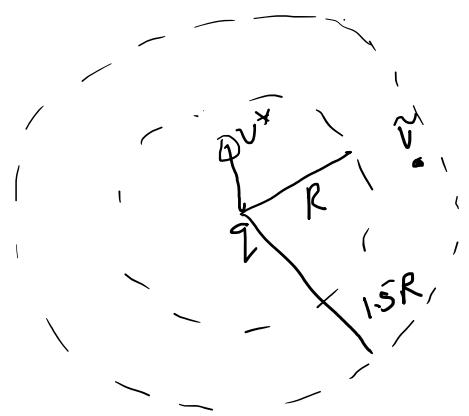
Today + Tomorrow:

Can do such a thing if we're OK
with Approximate Near Neighbor Search

Given q , target R ,

- Algo can output No if no base point is distance $\leq R$ from q .

- Output any base point at distance $\leq CR$ from q^* - for some $C > 1$.



[Algo can output \hat{v} instead of v^*]

In this case $C = 1.5$

C is fixed up-front, is like
“approximation factor of algo”.

THM

[Indyk-Motwani '98]

For above problem, can design

C -Approx NNS algorithm with

space $O(N^{1+\frac{1}{C}} + ND)$ but query time
 $= O(N^{\frac{1}{C}} D)$.

Idea: Design very generic technique called
“LOCALITY SENSITIVE HASHING”.

Imagine the following type of search Algo:-
Compute a hash function

Compute a hash function

$h: \{0,1\}^D \rightarrow [\lfloor k \rfloor]$ eg. where $k \ll N$
think of $k = \sqrt{N}$.

It maps each document vector to
one of k buckets.

Similarly it can map query to a bucket.

Alg: Only look in the query bucket
and output the closest
base vector.

Query time: $\approx \frac{N}{k} D \approx \sqrt{N} D$ if
 $k = \sqrt{N}$.

\downarrow
 h is a random hash
 \Rightarrow each bucket has
 $\approx N/k$ pts.

Issue

No reason why query's nearest neighbor
is hashed to same bucket as q .

Idea:

What if hash for ' h ' is locality sensitive?
(ie) nearby points are in
same bucket more often
than far away points!

↓

than far away points !
Then maybe such an algo would work !

LSH

A hash family $\mathcal{H} = \{h_1, \dots, h_m\}$ is
a (c, R, p_1, p_2) LSH for $p_1 > p_2$ if

for any 2 vectors x and y ,

$$\textcircled{1} \quad \Pr_{h \sim \mathcal{H}} [h(x) = h(y)] \geq p_1 \quad \text{if } d(x, y) \leq R$$

$$\textcircled{2} \quad \Pr_{h \sim \mathcal{H}} [h(x) = h(y)] \leq p_2 \quad \text{if } d(x, y) > cR$$

Property $\textcircled{2}$ ensures that buckets are small
on avg, and

$\textcircled{1}$ ensures that near neighbors of query
are in same bucket or q.

{ From (c, R, p_1, p_2) LSH to (c, R) Algo }

Idea :

$\textcircled{1}$ "boost" the gap between p_1 and p_2
by taking

① Now by taking

$$g(x) = [h_1(x), h_2(x), \dots, h_k(x)]$$

where h_1, h_2, \dots, h_k are k independent samples from \mathcal{H} .

What does g satisfy:

$$\Pr[g(x) = g(y)] \geq p_1^k \quad \text{if } d(x, y) \leq R$$

$$\Pr[g(x) = g(y)] \leq l_2^k \quad \text{if } d(x, y) > R$$

- This has driven down p_2^k
- Now, this also has driven down good case collision (p_1^k)
- Fix this by using L different hash functions

$$g_1, g_2, \dots, g_L \left\{ \begin{array}{l} \text{independently} \\ \text{for each } g_i \end{array} \right\}$$

Overall Map :-

① { Compute $g_i = (h_{i1}, h_{i2}, \dots, h_{ik})$
for $i = 1, 2, \dots, L$.

② { $x \mapsto g_i$ if $k_i \leq 1$ and

② Compute $g_i(v)$ for all $1 \leq i \leq L$ and
 all base vectors.
 When query arrives
 Compute $g_1(q), \dots, g_L(q)$.
 For $i = 1, 2, \dots, L$
 - Look at all base vectors with
 $g_i(v) = g_i(q)$.
 - If we find any at distance
 $\leq c_R$, output
 and terminate

Need to analyze time and space complexity.

Space Complexity : $N \cdot D + O(N \cdot K \cdot L)$
 hash values.

Query Time :-

$LK + L * \left(E[\# \text{ far away pts which fall into bucket of } q] \cdot D \right)$
 ↑ # hash fns for g ↑ of q ↑ dimension
 Sort of "wasteful computation"



sort of
"wasteful computation"

$$\leq L \cdot N \cdot P_2^k \cdot D$$

$$E[\text{query time}] \leq L N P_2^k \cdot D$$

Now it's all about parameter choosing

choose k st $P_2^k = \frac{1}{N}$

(ie)

$$k \log \frac{1}{P_2} = \log N$$

$$k = \frac{\log N}{\log \gamma_{P_2}}$$

{ Next choose L so that q 's near-neighbor
is in the bucket with good
probability

Let v^* be q 's neighbor within
distance $\leq r$ (if any).

What is the probability that
it falls into one of the
 L buckets of q ?

$\Pr[v^* \text{ falls in } q^s \text{ bucket in one of the } L \text{ hash fns}]$

$$\begin{aligned} & 1 - \prod_{i=1}^L (1 - p_i^k) \\ & = 1 - (1 - p_i^k)^L \quad (1 - \varepsilon)^L \\ & \approx L \cdot p_i^k \quad \approx 1 - \varepsilon L \\ & \text{Need to normalize} \rightarrow \sim \quad \text{when } \varepsilon \text{ is small} \end{aligned}$$

Set L st

$$L \cdot p_i^k = 1$$

to ensure good success probability

$$\begin{aligned} L &= \frac{1}{p_i^k} \\ &= \left(\frac{1}{p_i}\right) \frac{\log N}{\log(\gamma_p)} \end{aligned}$$

$$L = N \frac{\log(\frac{1}{p})}{\log(\frac{1}{p_2})} \quad p$$

Parameter $p = \underline{\log(\gamma_p)} < 1$

$\log(\gamma_{PL})$
 The important parameter of
 LSH.

Summary

$$k = \frac{\log N}{\log \gamma_{PL}}; L = N^p$$

Expected Query Time

$$= KL + O(N \cdot L \cdot p_L^k \cdot D)$$

$$= \frac{\log N}{\log \gamma_{PL}} \cdot N^p + O(N^p \cdot D)$$

{great if $p < 1$ } ☺

Success Prob

If \exists a near vbr, we
 find some approx NN
 with prob

$$\geq \left(1 - \left(1 - p_1^k \right)^L \right) - p_1^k \cdot L$$

$$(e^{+}, 1+x) \xrightarrow{?} 1 - e^{-1-x} .$$

$$(e^+ > 1+\epsilon) \geq (1 - \gamma_e) \cdot \boxed{B}$$

THM

(C, R, P_1, P_2) LSH with $\ell = \frac{\log \gamma_R}{\log \gamma_{P_2}}$

implies a randomized

(C, R) -ANN5 algorithm

with success prob $\geq (1 - \gamma_e)$ of
finding a near nbr @
distance $\leq CR$ (if there exists
some nbr at dist $\leq R$)

and expected query processing time

$$= O(N^\ell \left(\frac{\log N}{\log \gamma_{P_2}} + D \right))$$

\leq constant, can ignore

\Rightarrow LSH type hash fns are good as
long as ℓ is small !!

QN

for vectors in $\{0, 1\}^D$ with $d(x, y)$

$$= \sum_{i=1}^D |x_i - y_i|_1 \text{ as distance}$$

What is a good LSH?

- Want
- If x, y are close ($\leq R$)
 $\Pr[h(x) = h(y)]$ is large
 - If x, y are far apart ($\geq CR$)
 $\Pr[h(x) = h(y)]$ is small.

Dirt Simple Idea

Pick a random coordinate
 d from $\{1, 2, \dots, D\}$.

$$\boxed{h_d(x) = x_d}$$

$$H = \{h_1, h_2, \dots, h_D\}$$

P_1, P_2 ANALYSIS

If $d(x, y) \leq R$, (ie) they differ
 in at most
 R places,

$$\Pr_d [h_d(x) = h_d(y)] \geq \underbrace{\left(1 - \frac{R}{D}\right)}_{P_1}.$$

$$\Pr_d [h_d(x) = h_d(y)] \leq \underbrace{\left(1 - \frac{Ct}{D}\right)}_{P_2}.$$

$\leftarrow d \leftarrow$ $\curvearrowright P_2$
 If $d(x, y) \geq cR$

What is the f of this hash fn.

$$f = \frac{\log(1/\epsilon R)}{\log(1/\epsilon R_2)}$$

$$= \frac{\log\left(\frac{1}{1 - \frac{R}{D}}\right)}{\log\left(\frac{1}{1 - \frac{R_2}{D}}\right)}$$

Need to be formalized
 Assuming $R \ll D$,
 we'll make
 some simplifications

$$\begin{aligned}
 a) \frac{1}{1-\epsilon} &\approx 1+\epsilon \\
 b) \log(1+\epsilon) &\approx \epsilon
 \end{aligned}
 \quad \left. \begin{array}{l} \\ \end{array} \right\} \text{if } \epsilon \ll 1.$$

Hence $f \approx \frac{R^D}{D \cdot cR} = \frac{1}{c}$.

Using this hash family gives us
the (C, R) - Near Neighbor search
problem with
Space $\mathcal{O}(N^{1+1/C} + ND)$ and
Query time $\mathcal{O}(\underbrace{N^{1/C}}_{\sim} \cdot (D + \log N))$.

↓

represents a significant milestone
for "nearest vector search" and
is used in variety of applications
today !!