



Challenging Differential Privacy The Case of Non-interactive Mechanisms

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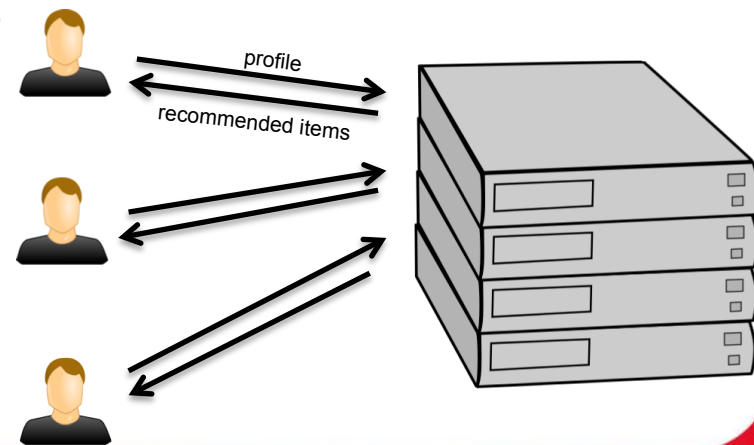
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Outline

1. Personalization and privacy
2. Theoretical analysis
3. Practical decoders
4. Experiments

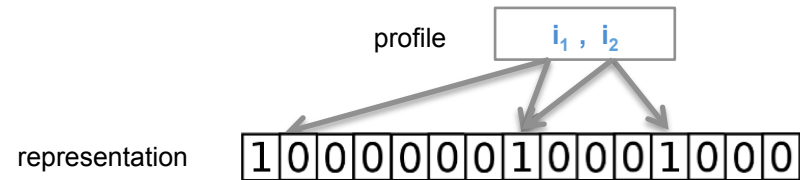
Personalized recommendation system

- Use user-item similarity for prediction and ranking
- Maintain a user profile
 - Composed of past items and preferences
- Aggregate user profiles for similarity computation
 - Needs profile information exchange

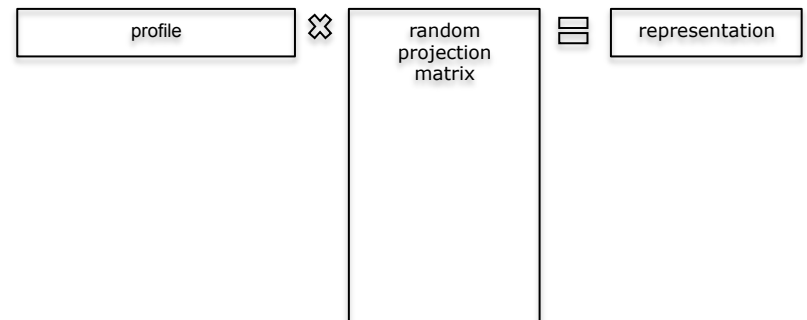


Profile representation

- Compact representation of user profile
- Examples
 - Bloom filter hash-based probabilistic data structure
 - Items stored as bits addressed by k-hash functions



- Random projection in low dimensional space (Johnson–Lindenstrauss transformation)
 - Items are vector points



Privacy

- User profile is personal data
- Sanitization mechanism
 - Modify the representation before its disclosure
- Measures of privacy
 - k -anonymity, l -diversity, t -closeness, ...
 - We choose differential privacy

Differential privacy [1]

- $\mathcal{F} : \mathcal{D}^n \rightarrow \mathcal{D}^n$: ϵ -differentially private randomized function if for all neighboring profiles \mathbf{x}' of \mathbf{x} , where $\mathbf{x}', \mathbf{x}, \mathbf{t} \in \mathcal{D}^n$

$$\mathbb{P}[\mathcal{F}(\mathbf{x}) = \mathbf{t}] \leq e^\epsilon \cdot \mathbb{P}[\mathcal{F}(\mathbf{x}') = \mathbf{t}]$$

- Achieved by randomized perturbation of data
 - Interactive: perturbed for each query
 - Non-interactive: perturbed and published
- In personalization:
 - Altering one item will not change the probability of a user profile representation \mathbf{t}

[1] Dwork, C.: Differential privacy, ICALP 2006

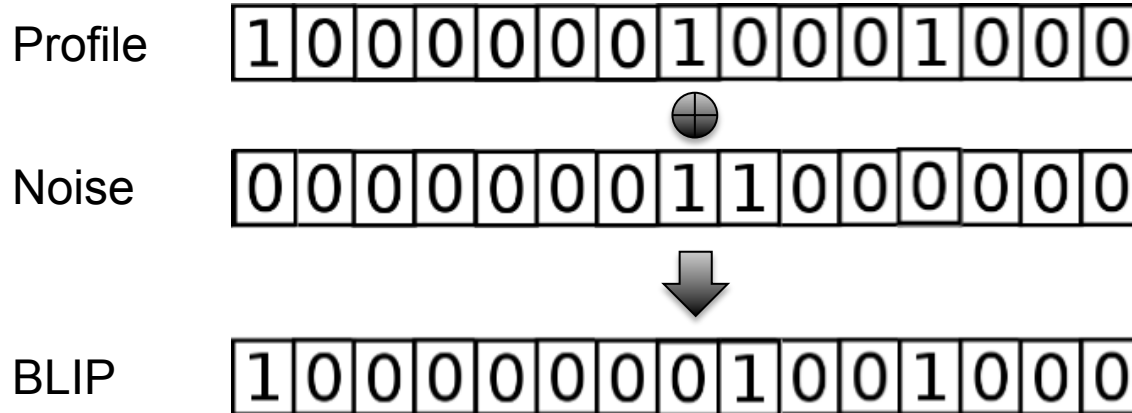
BLIP [2]

1. Profile representation: Bloom filter B_P

2. ϵ -differentially private representation: BLIP: $\tilde{\mathbf{B}}_P = \mathbf{B}_P \oplus \mathbf{Noise}$

– Randomization by binary noise, i.i.d. Bernoulli distribution

– Random flipping with probability $p_\epsilon = 1/(1 + e^{\epsilon/K})$.



[2] Alaggar, M., Gambs, S., Kermarrec, A.-M.: BLIP: Non-interactive Differentially-Private Similarity Computation on Bloom Filters, SSS 2012

JLT [3]

1. Profile representation: Johnson-Lindenstrauss Transform

$$\mathbf{Y}_P = \sum_{j \in P} \mathbf{X}_j$$

- Codeword \mathbf{X}_j is real vector of length L : $X_j(i) \stackrel{i.i.d.}{\sim} \mathcal{N}(0, 1/L)$

2. (ϵ, δ) -differentially private representation: $\tilde{\mathbf{Y}}_P = \mathbf{Y}_P + \mathbf{Noise}$

- Randomization by noise: $Noise(i) \stackrel{i.i.d.}{\sim} \mathcal{N}(0, \sigma^2)$

with $L \geq 2(\log(N) + \log(2/\delta))$, $\sigma \geq \frac{4}{\epsilon} \sqrt{\log(1/\delta)}$ and $\epsilon < \log(1/\delta)$

[3] Kenthapadi, K., Korolova, A., Mironov, I., Mishra, N.: Privacy via the Johnson Lindenstrauss transform.

Journal of Privacy and Confidentiality, 5(1), 39-71.

Our Contribution

THEORETICAL ANALYSIS

Single decoder for BLIP

- The adversary infers the presence of a single item
 - Hypothesis test:
 - Item j is not in the profile $\mathbb{P}[\tilde{\mathbf{B}}_P(\ell), X_j(\ell)] = \mathbb{P}[\tilde{\mathbf{B}}_P(\ell)]\mathbb{P}[X_j(\ell)]$
 - Item j is in the profile $\mathbb{P}[\tilde{\mathbf{B}}_P(\ell), X_j(\ell)] = \mathbb{P}[\tilde{\mathbf{B}}_P(\ell)|X_i(\ell)]\mathbb{P}[X_i(\ell)]$
 - Mutual information $I(\tilde{\mathbf{B}}_P; \mathbf{X}_j)$ measures the amount of information the BLIP is disclosing about the presence of item j .
- By testing sequentially all N items, the adversary reconstructs the user items set P
 - α Probability of missing a user item
 - η Probability of including at least one wrong item

$$\log \eta \geq \log N - \frac{I(\tilde{\mathbf{B}}_P; \mathbf{X}_j)}{1 - \alpha}$$

Shift of paradigm

- We are now interested by testing if a given subset of c items is the true user items set P .
 - We call it a *joint* decoder
- By testing all c -items subsets, the adversary finds the true P with probability

$$\log \eta \geq \log N - \frac{I(\tilde{\mathbf{B}}_P; (\mathbf{X}_{j_1}, \dots, \mathbf{X}_{j_c}))}{c(1 - \alpha)}$$

Theoretical performances of a joint decoder

- From [4]: $I(\tilde{\mathbf{B}}_P; (\mathbf{X}_{j_1}, \dots, \mathbf{X}_{j_c})) / c \geq I(\tilde{\mathbf{B}}_P; \mathbf{X}_j)$
- This shows that:
Joint decoding is always more efficient than single decoding
- Our paper gives these theoretical performances for
 - Single and Joint decoding, applied to
 - BLIP and JLT approach
- Depending on the setup, there is a substantial difference between performances of single and joint decoding

[4] Universal Fingerprinting: Capacity and Random-coding exponents, P. Moulin, ISIT 2008

A nice theoretical result but...

It does NOT work in practice because

- For $c \ll N$, the number of c -items subset is

$$\binom{N}{c} = O(N^c)$$

It is not tractable to test all c -items subsets!!!

Our Contribution

PRACTICAL DECODERS

Markov chain

- Instead of testing all subsets:
 - We do a guided random walk in the space of c -items subsets
 - This random walk leads to the most likely c -items subsets
- Input: The adversary observes one BLIP $\tilde{\mathbf{b}}$
- Starting point: a random subset $P^{(0)}$
- New state $P^{(t+1)}$ sampled with transition probability

$$\mathbb{P}[P^{(t+1)} = P | P^{(t)}] = \frac{\mathbb{P}[\tilde{\mathbf{B}}_P = \tilde{\mathbf{b}} | P] \mathbb{P}[P]}{\sum_{P' \in \mathcal{V}(P^{(t)}, i)} \mathbb{P}[\tilde{\mathbf{B}}_{P'} = \tilde{\mathbf{b}} | P'] \mathbb{P}[P']}$$

- Converges to $\mathbb{P}[P | \tilde{\mathbf{b}}]$ as $t \rightarrow \infty$, in practice at $t > T$ (burn-in period)

Monte Carlo

- Once the Markov chain has converged,
 - We sample subsets according to posteriori probability $\mathbb{P}[P|\tilde{\mathbf{b}}]$
 - We let the chain running for M more iterations.
- Possible outputs of the Monte Carlo Markov Chain:
 - Marginal a posteriori probability per item estimated by empirical frequency

$$\hat{\mathbb{P}}[j \in P|\tilde{\mathbf{b}}] = |\{t \in [T + 1, T + M] | j \in P^{(t)}\}|/M$$

- Maximum a posteriori estimator of the profile

$$\hat{P} = \arg \max_{T+1 \leq t \leq T+M} \mathbb{P}[P^{(t)}|\tilde{\mathbf{b}}]$$

EXPERIMENTS

Setup

- Datasets
 - Digg: Social bookmarking dataset
 - MovieLens: Movie rating dataset

	Nb of users	Training set size	Testing set size	N	c_{avg}	Sparsity %
Digg	531	331	200	1237	317	25.63%
MovieLens	943	600	343	1682	106	6.30%

- Attack algorithms:
 - Single decoder
 - Joint decoder with uniform prior
 - Joint decoder with prior estimated from the training set
 - Popularity-based attack (baseline)
 - The c most popular items

Privacy measures

- Profile reconstruction

- Cosine between original and reconstructed profiles

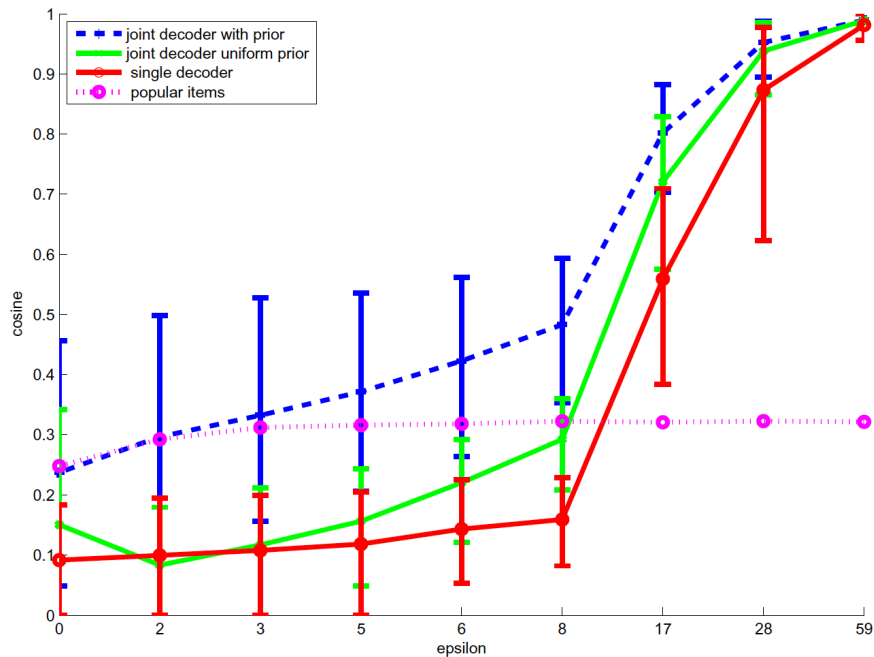
$$\cos(P, \hat{P}) = \frac{|P \cdot \hat{P}|}{|P| |\hat{P}|}$$

- Presence of individual item

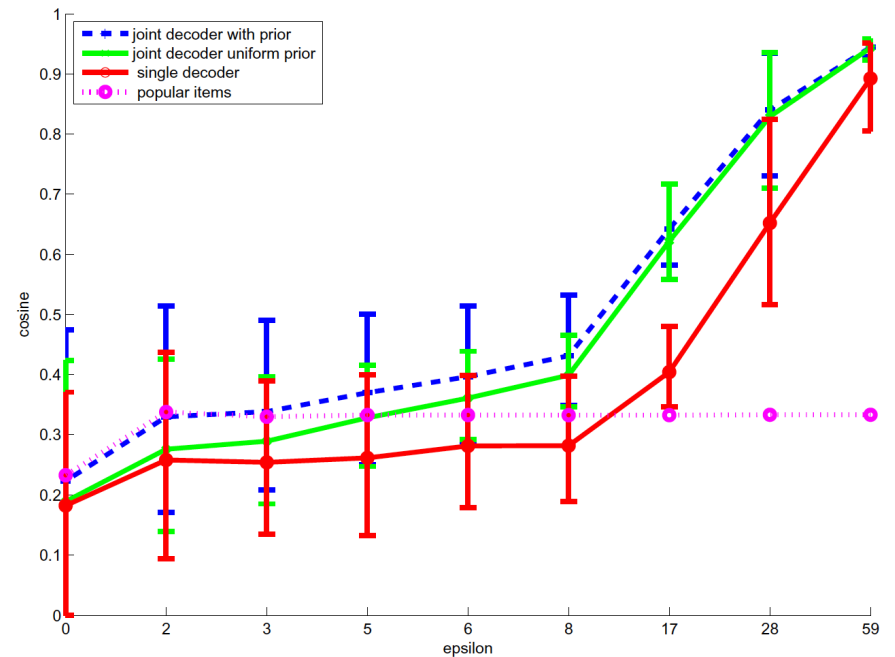
- Mean average precision of top- R ranked items

$$\text{mAP}@K = \frac{1}{Q} \sum_{q=1}^Q \left(\frac{1}{R} \sum_{r=1}^R \text{precision}_q(r) \right)$$

Profile reconstruction

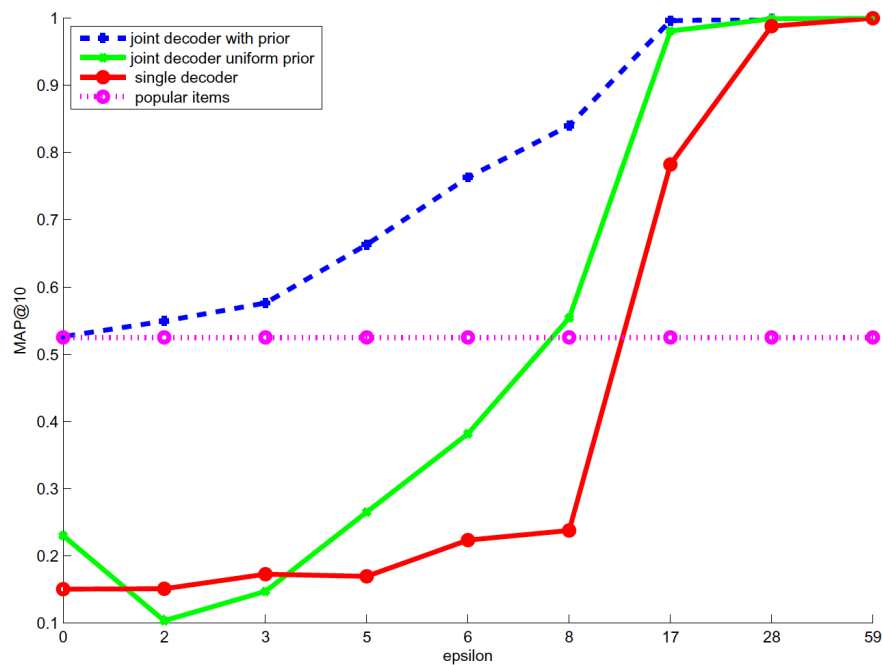


MovieLens

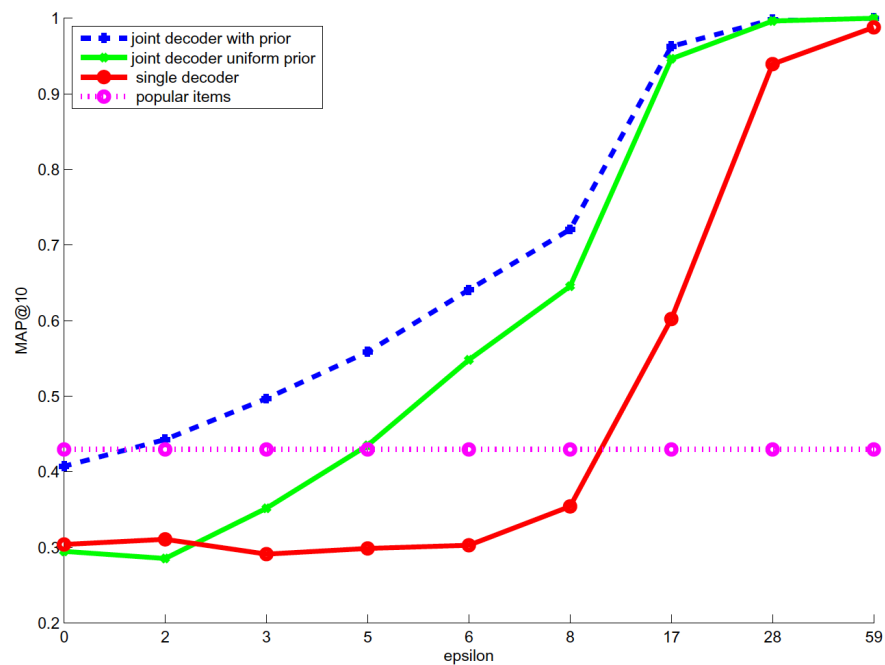


Digg

Presence of individual item



MovieLens

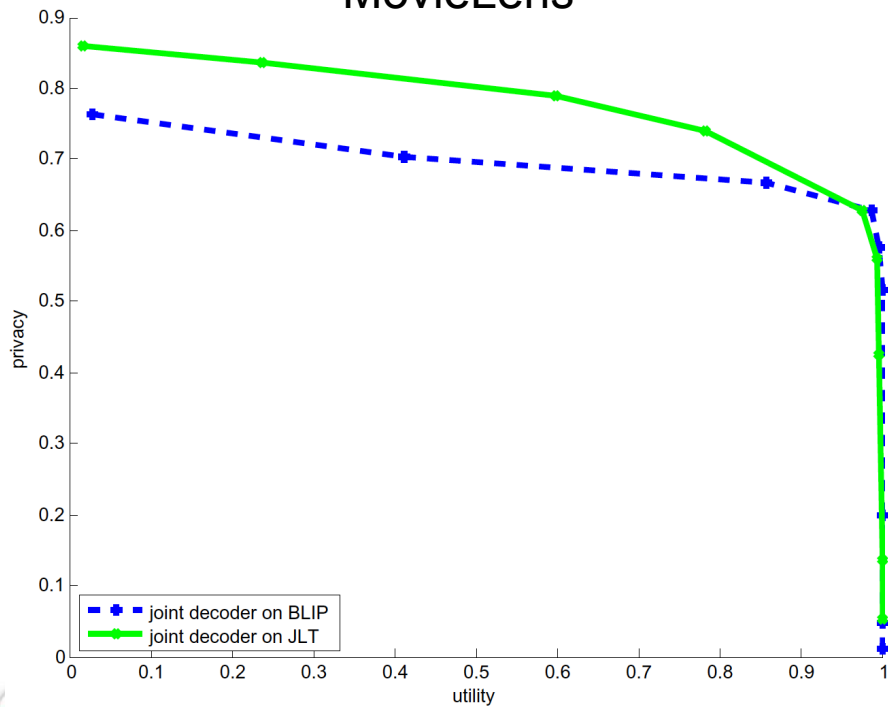


Digg

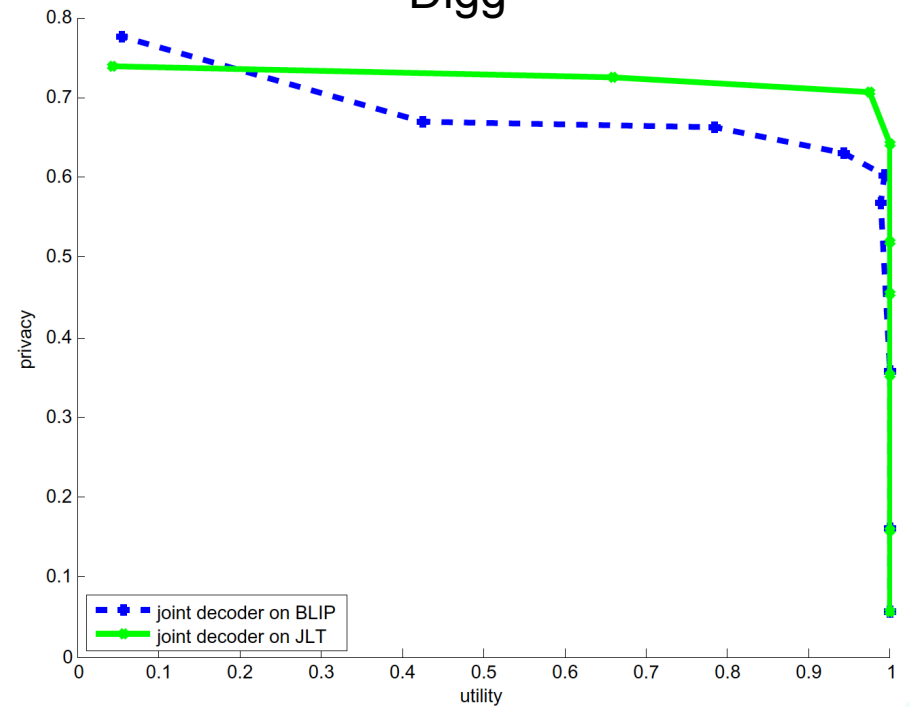
Utility vs privacy

- Privacy-Utility tradeoff
 - Privacy: $1 - \cos(P, \hat{P})$
 - Utility: $recall@10$
- $recall@k$: probability that the most similar profile is among the top- k ranked profiles

MovieLens



Digg



Conclusion

- Two attacks
 - Single and joint decoding
- Evaluated against two differentially private schemes:
 - BLIP and JLT
- Theoretical analysis and experimental results shows:
 - Joint decoding is more powerful than single decoding**
- Practical implementations of a joint decoder
 - We use a Monte Carlo Markov Chain (MCMC)
 - There are alternatives (Belief propagation, Iterative joint decoders)

Conclusion

- Our attacks help:
 - Understand the privacy guarantees of differentially-private mechanisms
 - Experimentally tune the parameter ϵ
 - Compare different non-interactive mechanisms
- Open Question
 - Towards a new definition of differential privacy?

$$\frac{I(\mathcal{F}(\mathbf{X}); \mathbf{X})}{|\mathbf{X}|} < \epsilon$$

where \mathbf{X} is a collection of $|\mathbf{X}|$ items.

Questions?

THANK YOU!