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References 3



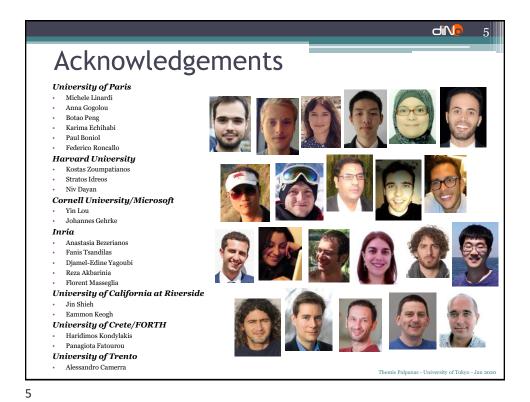
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data series toolbox

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- demos
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- DPiSAX: http://imitates.gforge.inria.fr/
- RINSE: http://daslab.seas.harvard.edu/rinse/

nestor project

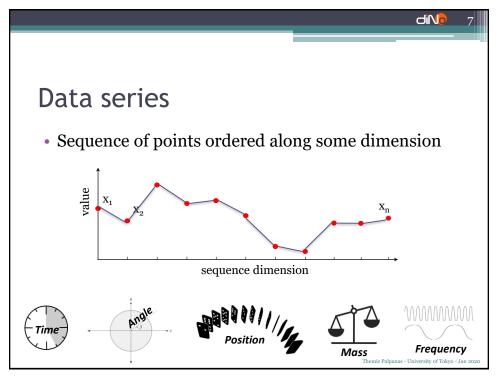
http://nestordb.com

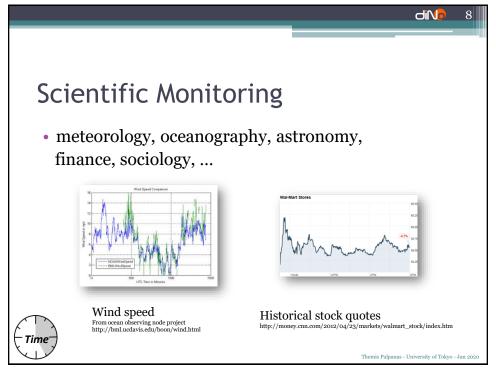


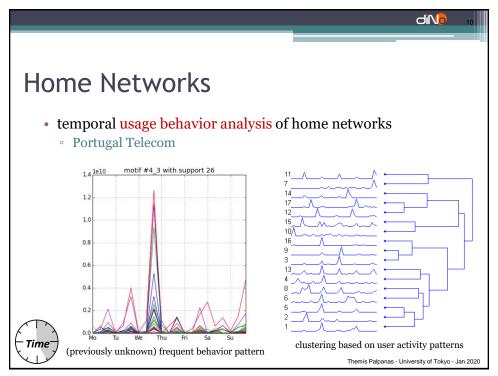
Executive Summary
 data collected at unprecedented rates
 they enable data-driven scientific discovery
 lots of these data are sequences

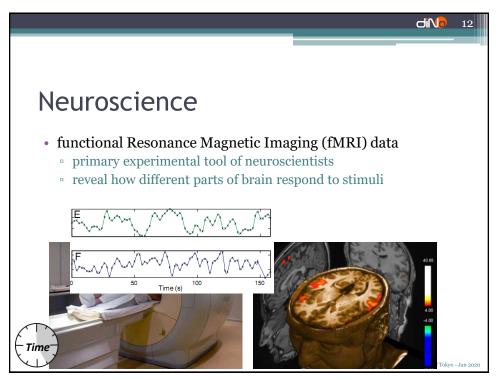
 takes days-weeks to analyze big sequence collections

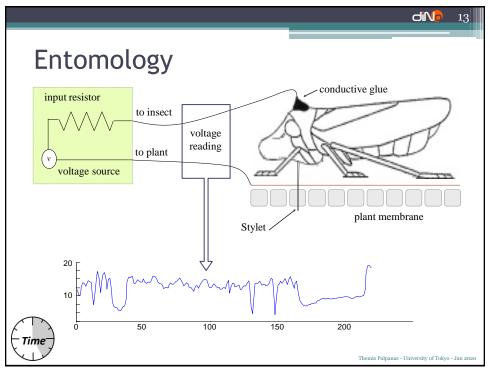
 goal: analyze big sequences in minutes/seconds

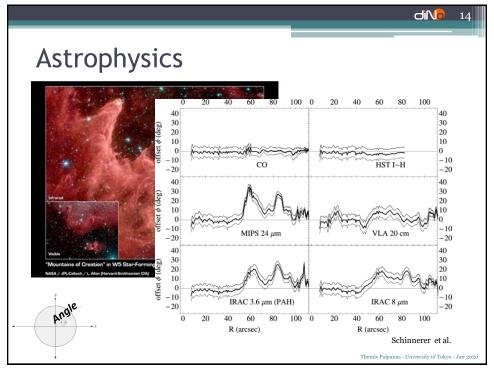


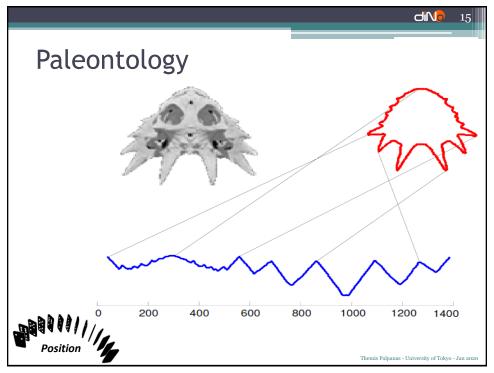


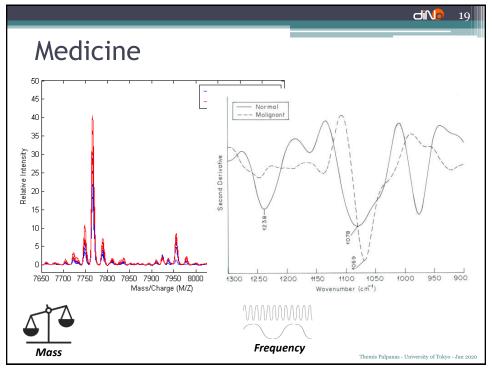


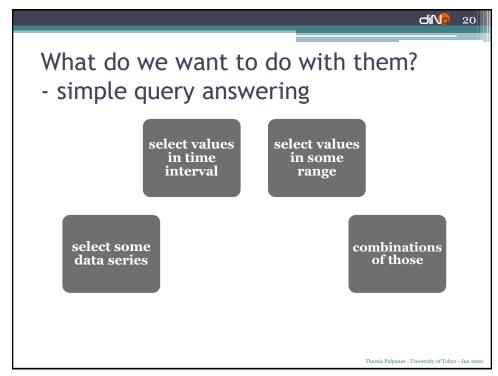


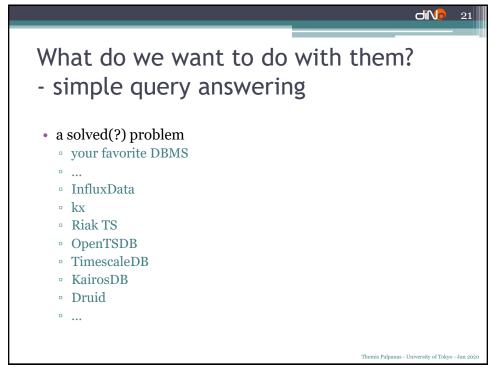


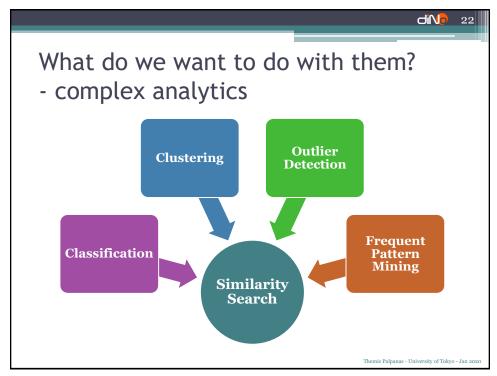


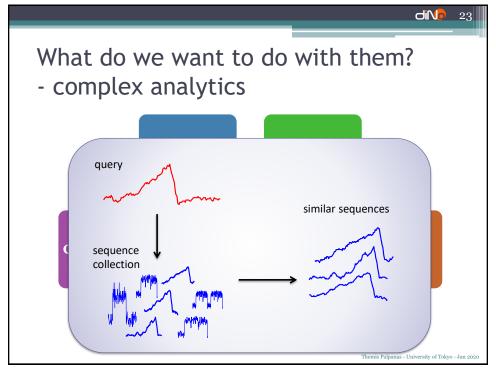


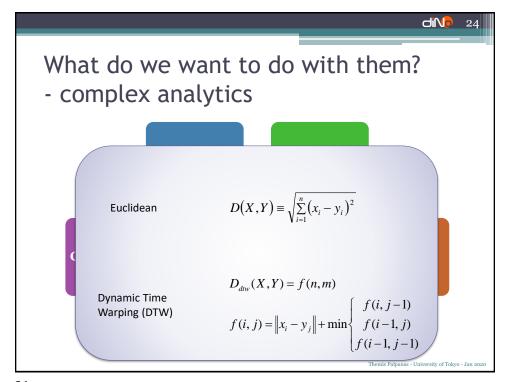


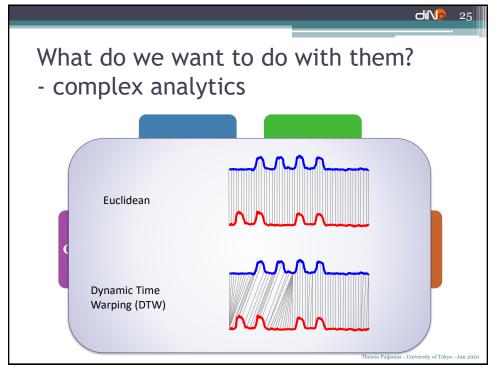


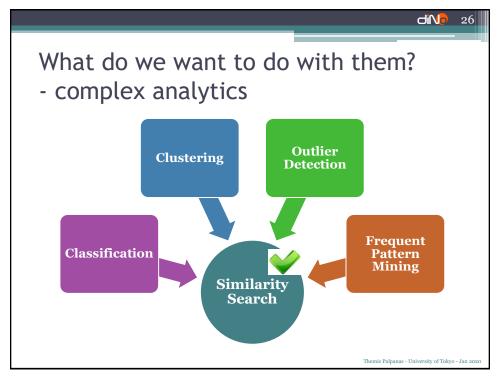


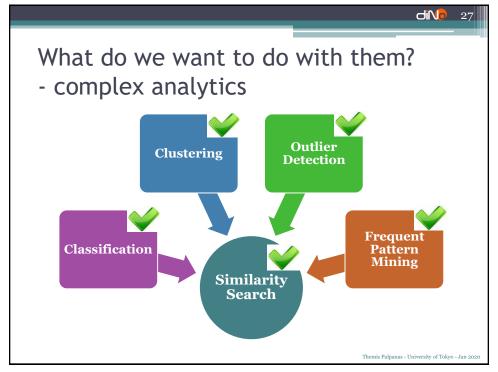


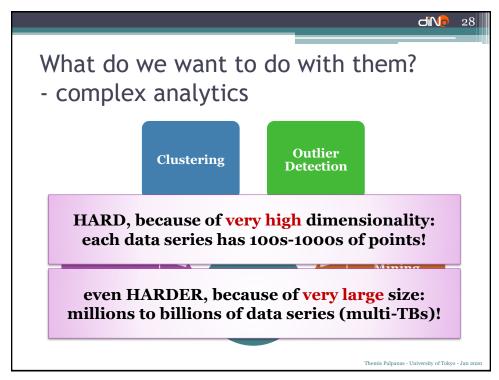


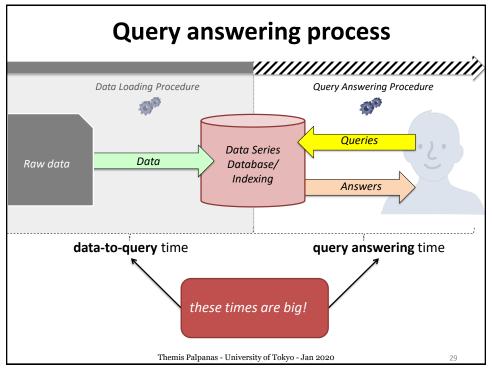


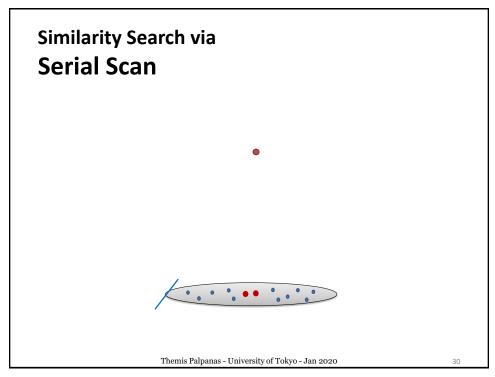


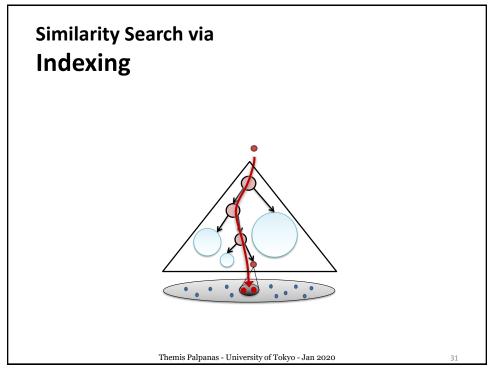


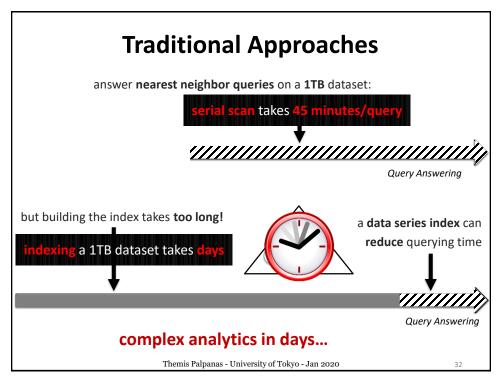


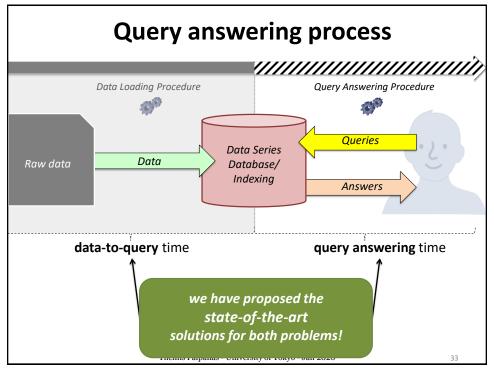


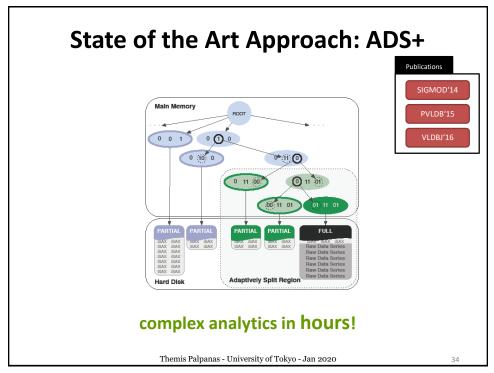


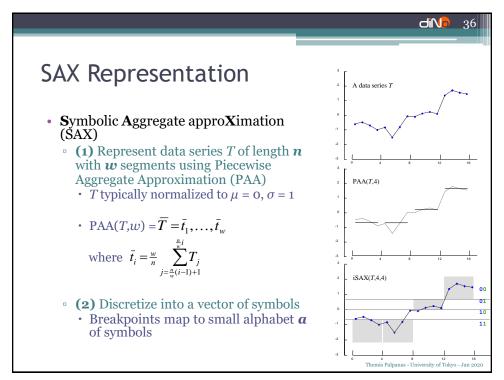


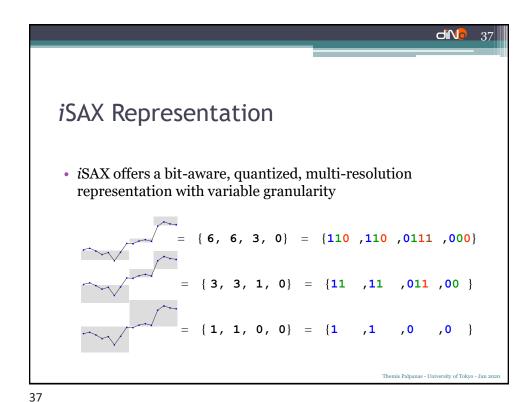


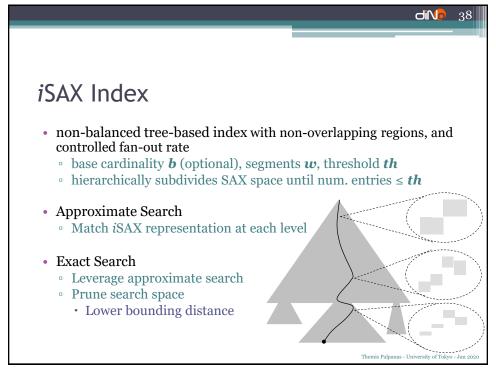


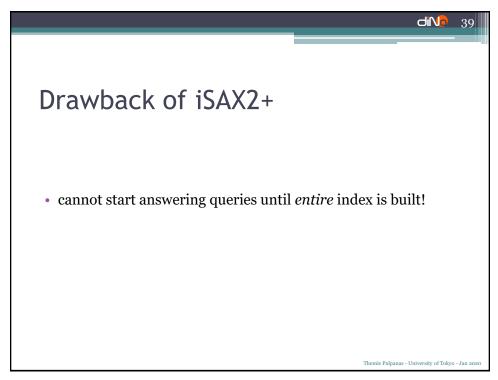


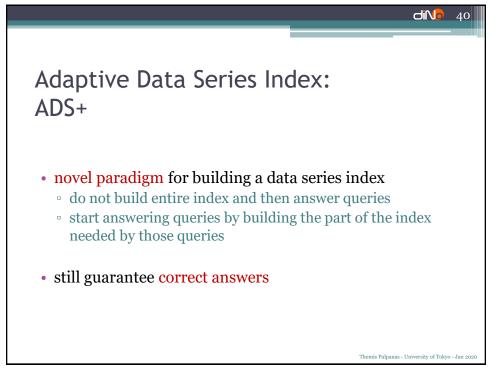


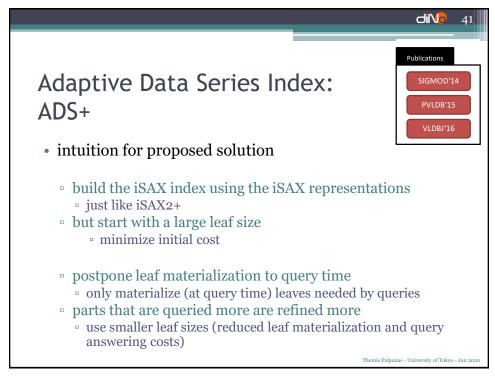


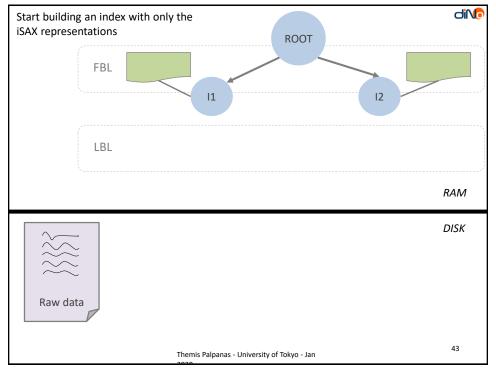


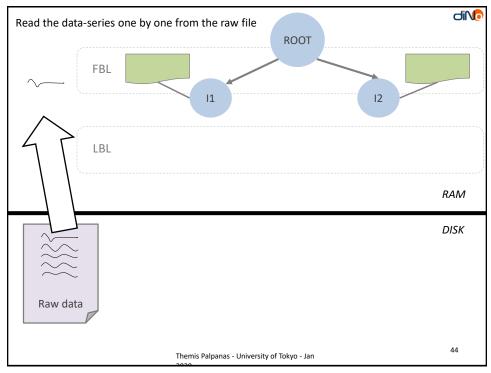


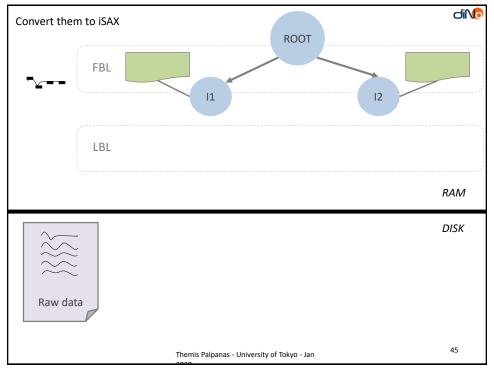


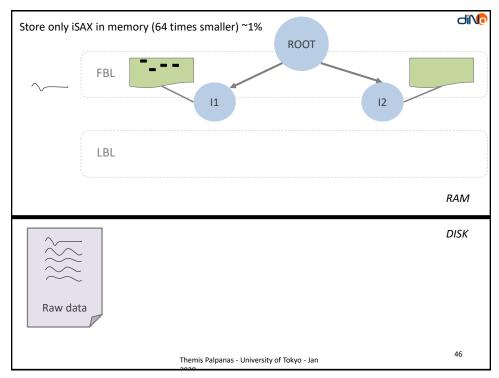


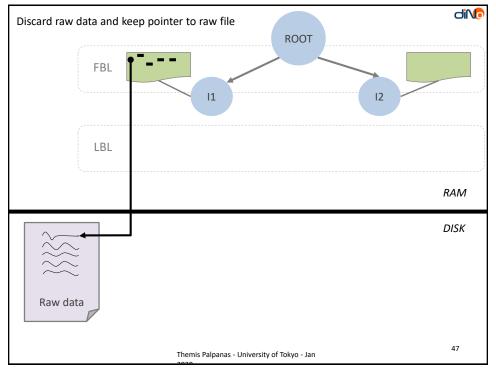


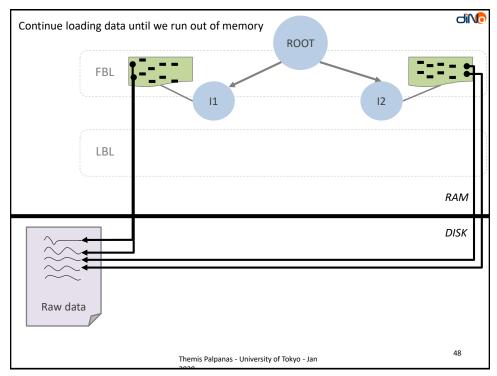


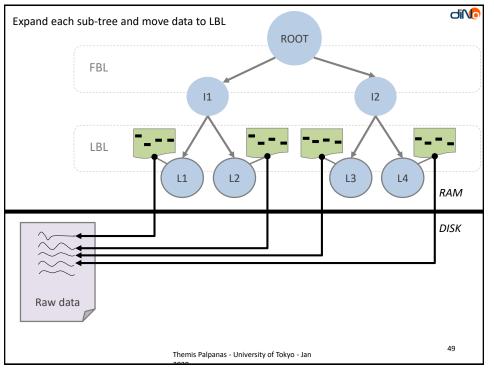


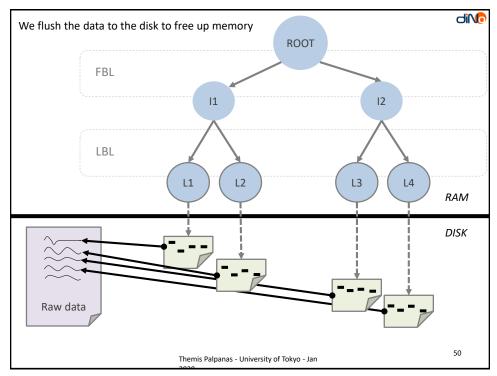


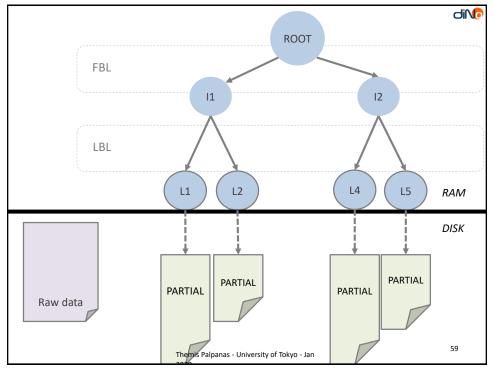


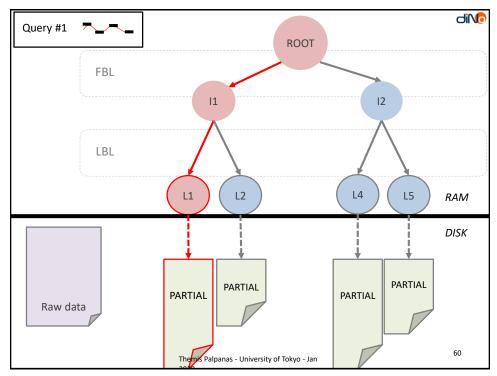


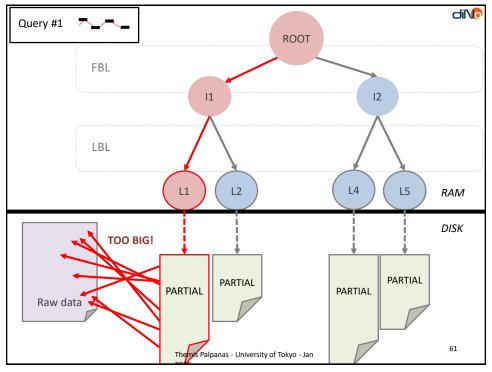


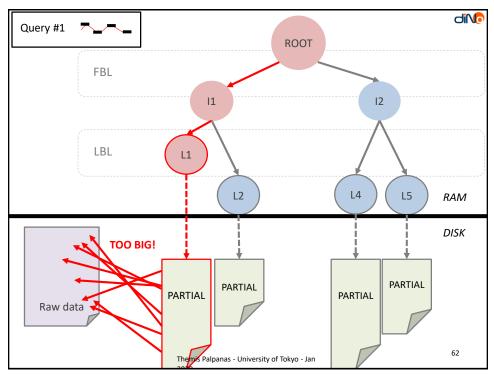


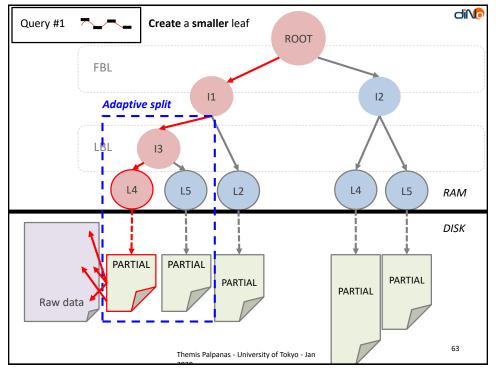


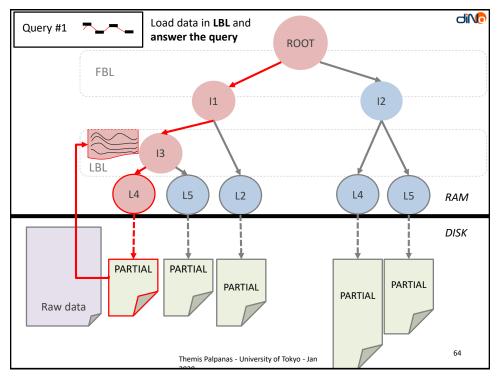


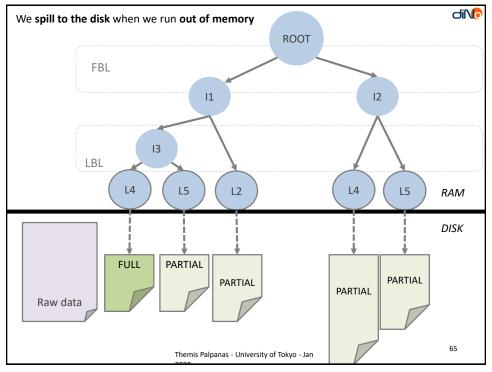


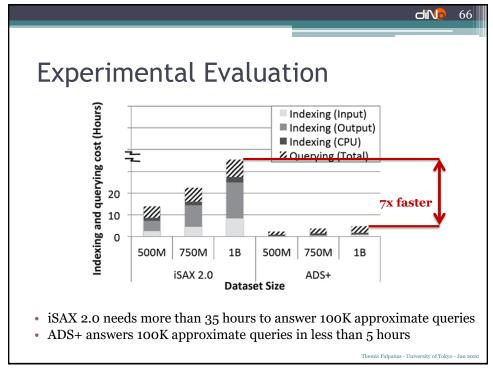


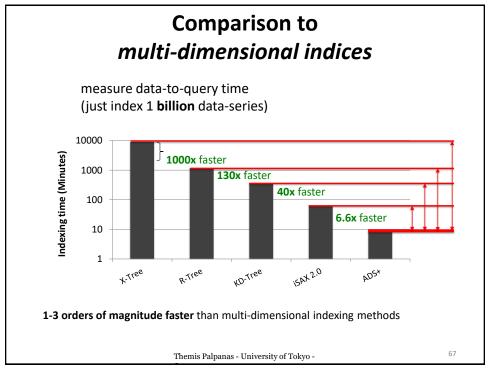


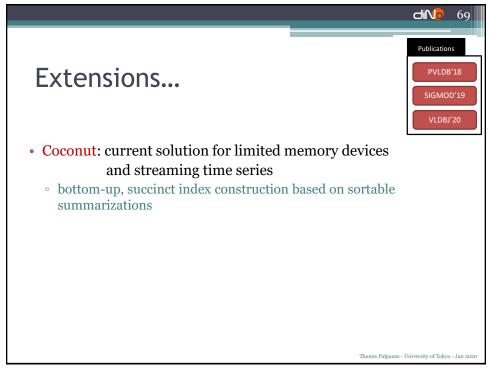


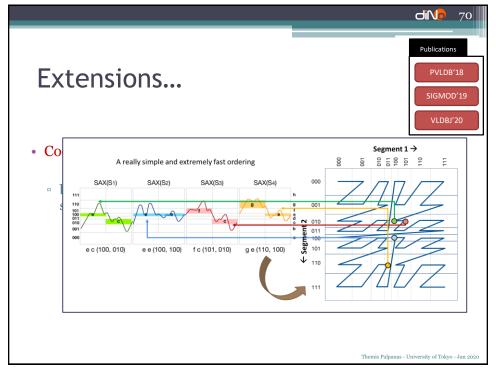


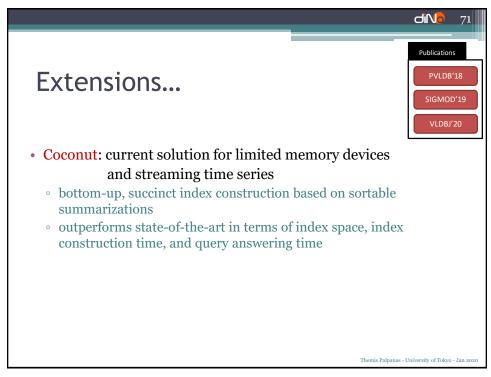


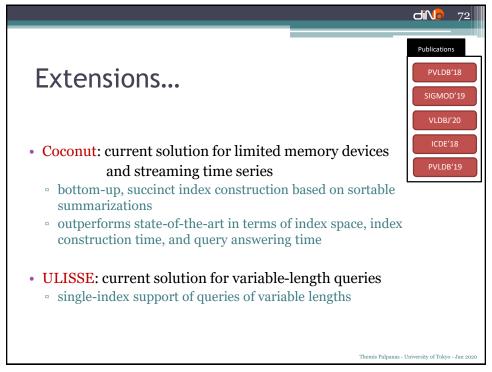


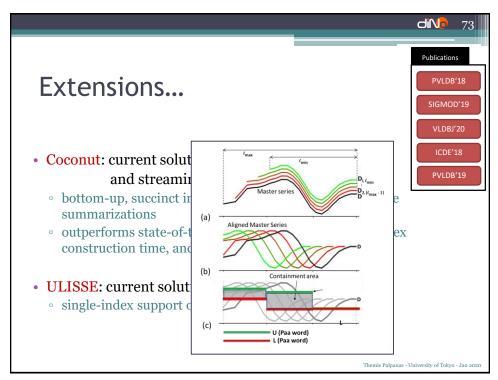


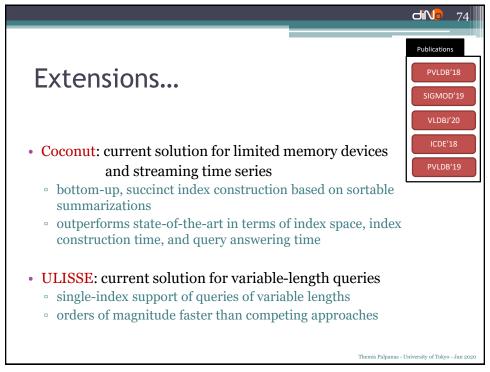


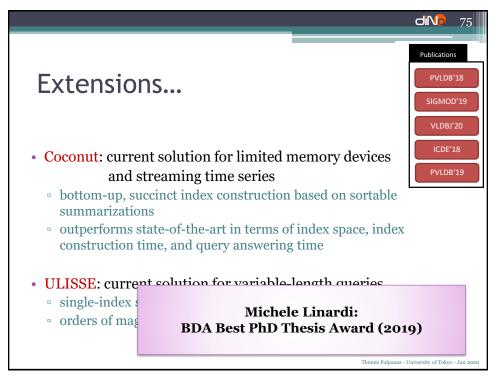


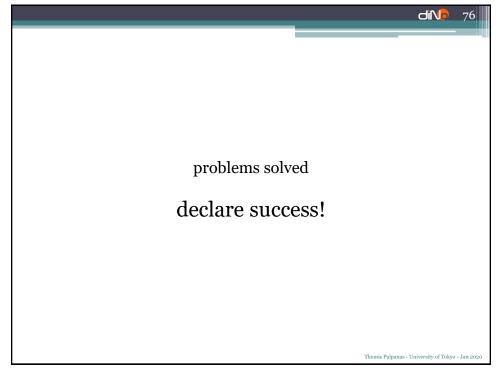


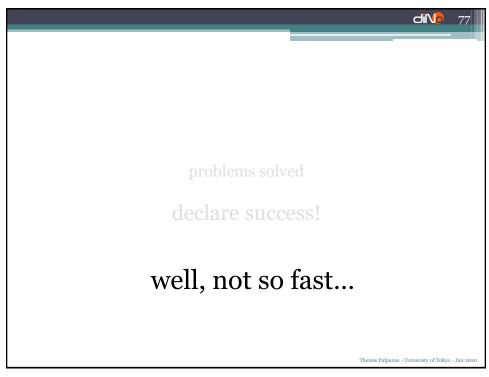


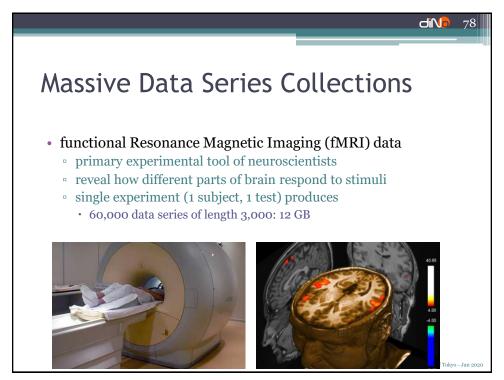




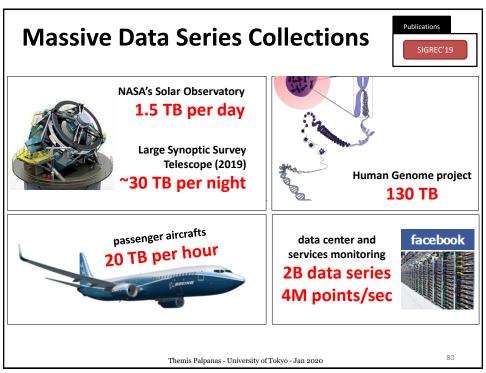


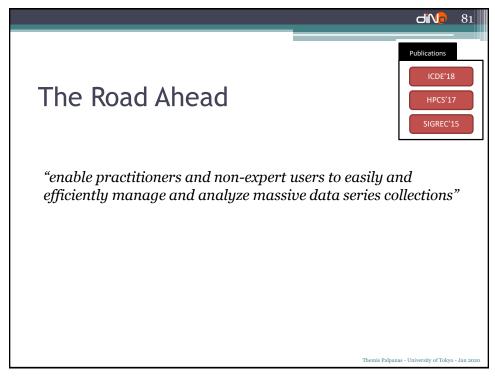


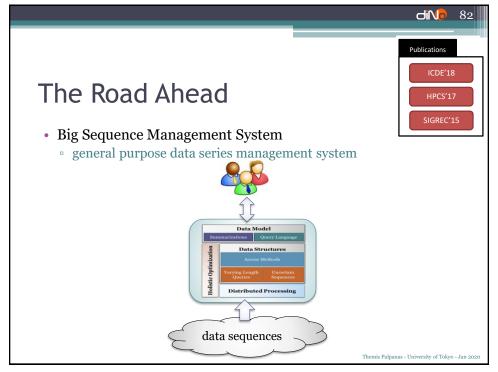


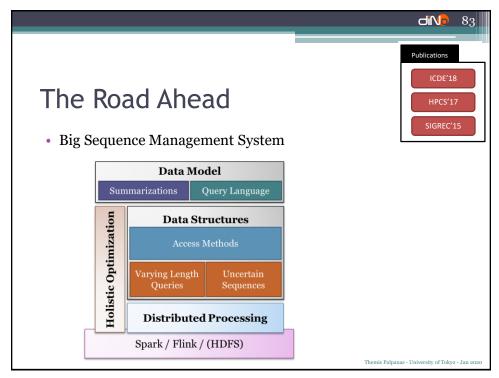


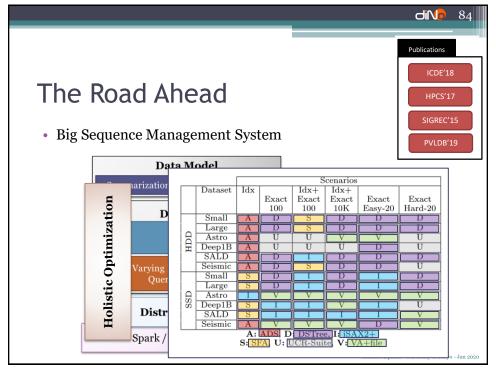


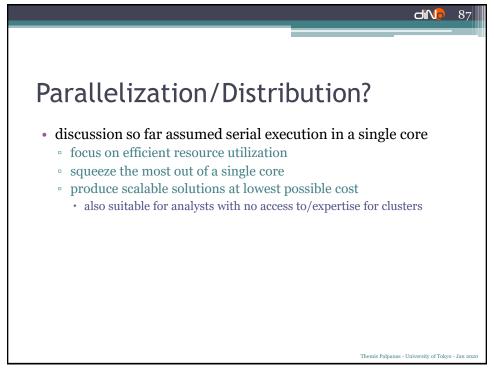


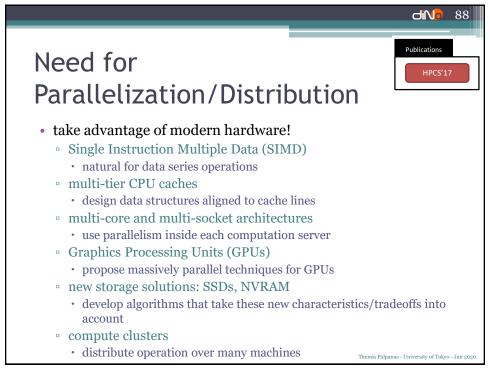


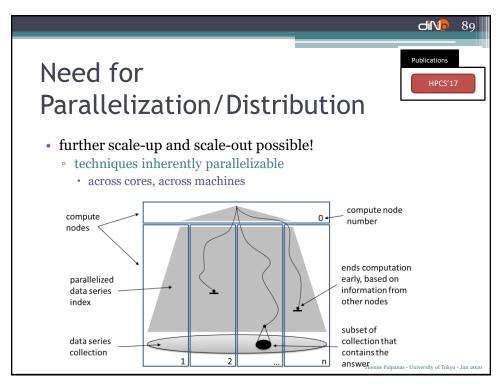


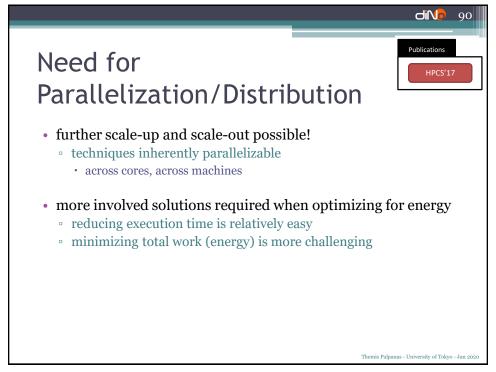


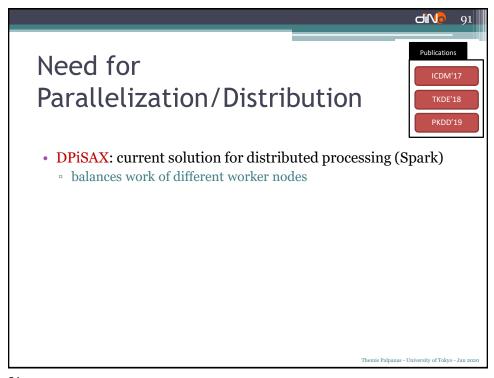


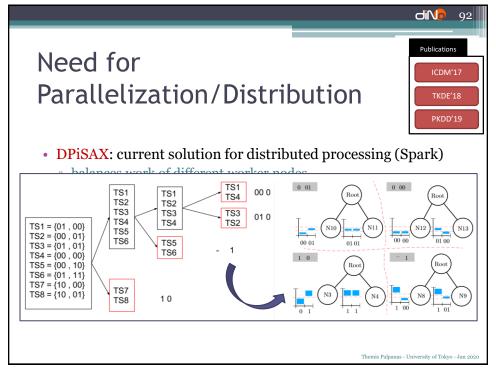


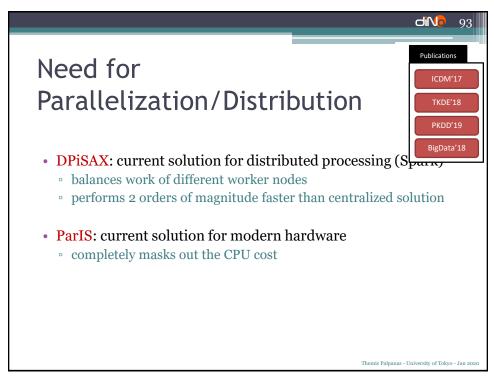


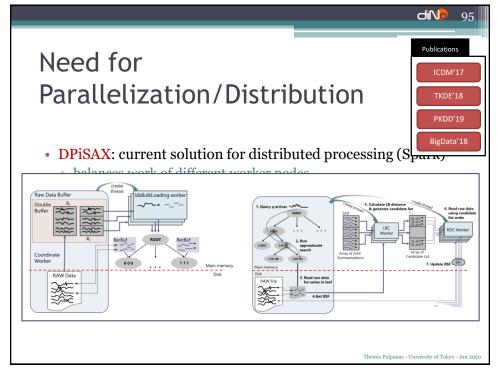


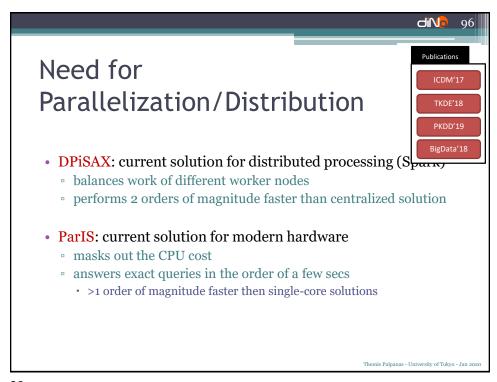


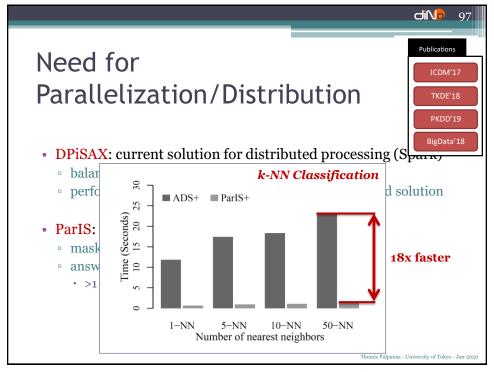


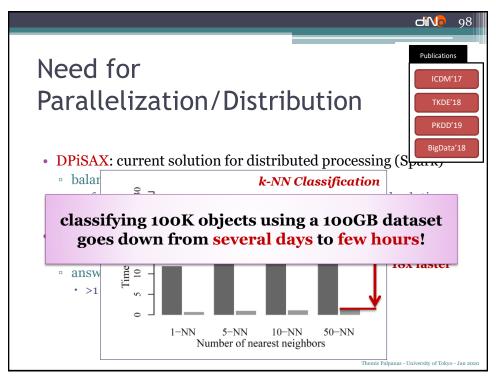


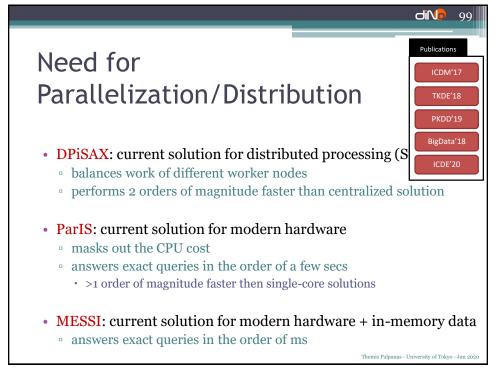


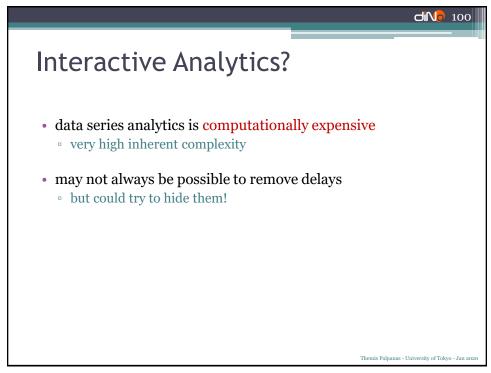


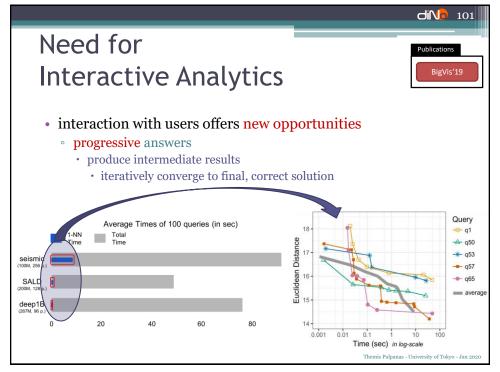












Need for Interactive Analytics

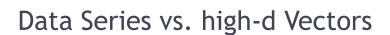


- interaction with users offers new opportunities
 - progressive answers
 - · produce intermediate results
 - · iteratively converge to final, correct solution
 - provide bounds on the errors (of the intermediate results) along the way
 - imprecise queries
 - enable user to specify varying accuracy requirements for different parts of the same query
- several exciting research problems in intersection of visualization and data management
 - *frontend*: HCI/visualizations for querying/results display
 - backend: efficiently supporting these operations

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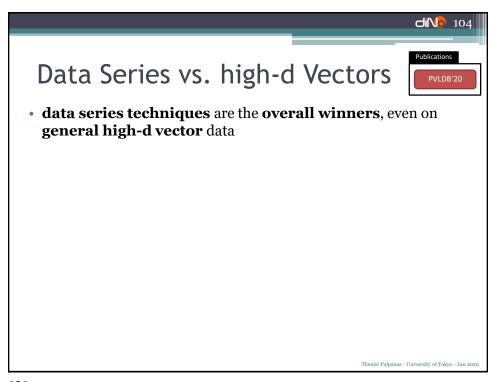
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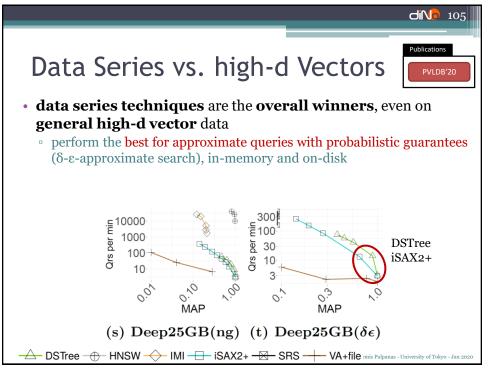
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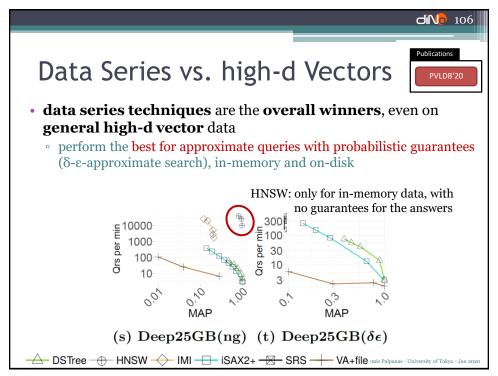


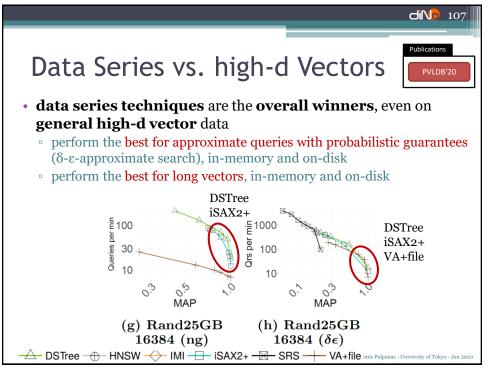
- two sides of the same(?) coin
 - data series as multidimensional points
 - for a specific ordering of the dimensions
- several techniques for similarity search in high-d vectors
 - using LSH (SRS), space quantization (IMI), k-NN graphs (HNSW)
- how do these high-d vector techniques compare to data series techniques?
 - currently conducting extensive experimental comparison

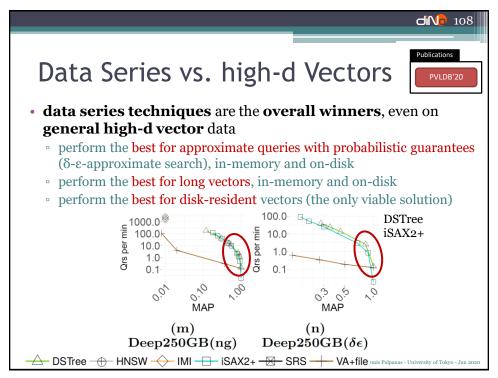
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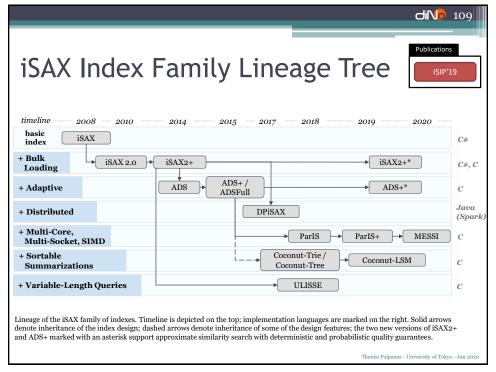


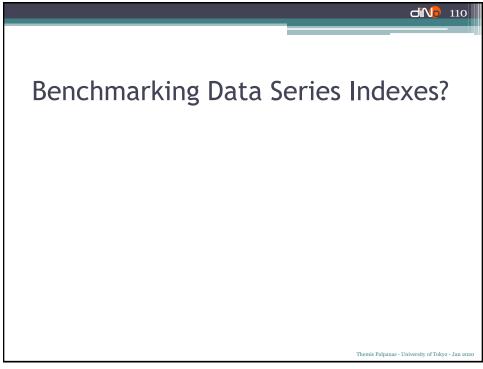












Previous Studies

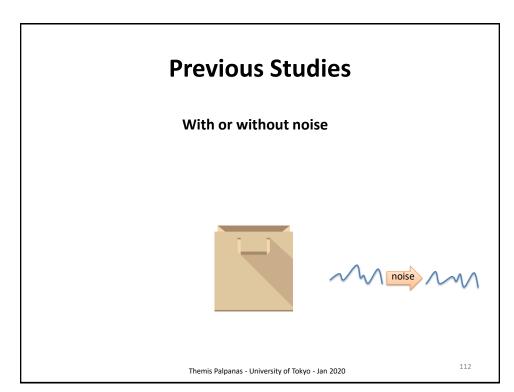
evaluate performance of indexing methods using random queries

• chosen from the data (with/without noise)

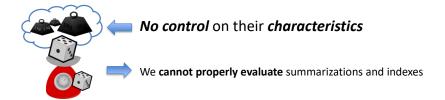


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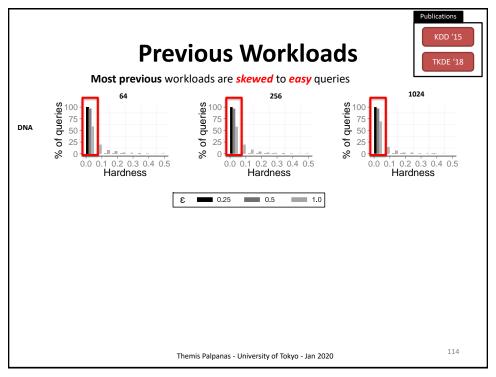


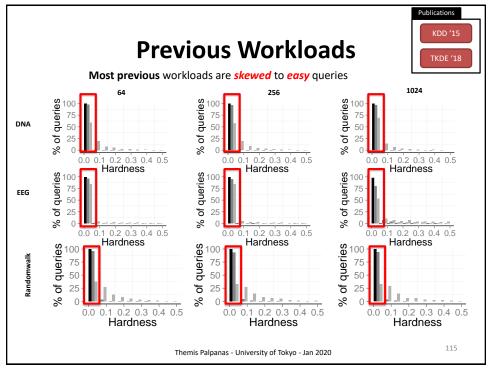


We need queries that cover the entire range from easy to hard

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Benchmark Workloads



If all queries are **easy** all indexes look **good**





If all queries are **hard** all indexes look **bad**





need methods for generating queries of varying hardness



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Contributions





Theoretical background

Methodology for characterizing NN queries for data series indexes



Nearest neighbor query workload generator Designed to stress-test data series indexes at varying levels of difficulty

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Subsequence Anomaly Detection

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Data Series Anomalies Problem

 develop anomaly detection techniques based on sequences (data series), not on individual values

O individual values can be normal, but their sequence can be abnormal!

Data Series Anomalies Problem 150 points in a sequence S Minimal critical value • develop anomaly detection techniques based on sequences (data series), not on individual values O individual values can be normal, but their sequence can be abnormal! Walues are not outside critical thresholds values are normal

