Finesse: Fine-Grained Feature Locality based Fast Resemblance Detection for Post-Deduplication Delta Compression

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Background

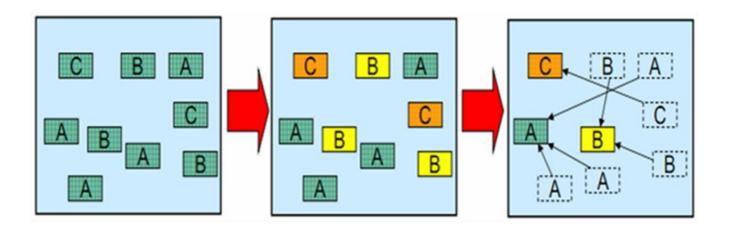


• Big data era

- Amount of digital data in the world will reach 44 ZB by 2020
- Redundant data in backup systems
 - About 88-90% of the data in EMC and Symantec's backup systems are duplicate (FAST'12, USENIX ATC'15)

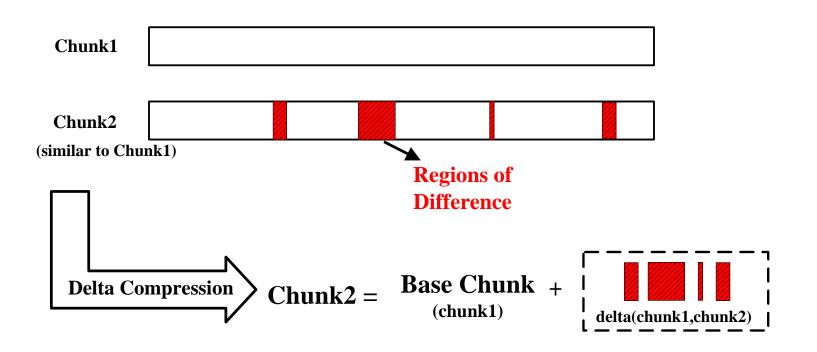
Data Reduction Technologies

- Data deduplication
 - Remove duplicate chunks according to their fingerprints, only store one unique chunks
 - Drawback: cannot remove redundant data among nonduplicate but very similar chunks



Data Reduction Technologies

- Delta compression
 - Achieve 2X more compression ratio beyond deduplication (FAST'12, Performance'14, Sigmod'17)



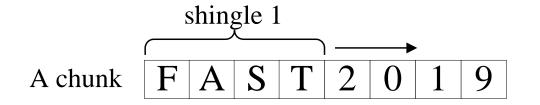
Resemblance Detection

• Detecting delta compression candidates

Traditional N-transform Super-Feature

- Generally, It extracts a fixed number of features from a chunk and grouping N features (N=12) into M SFs (e.g., M=3) for matching. One SF matching means the two chunks are very similar
- Feature extraction is time-consuming: Requiring N linear transformations for each fingerprint to generate N-dimensional hash value sets (features)

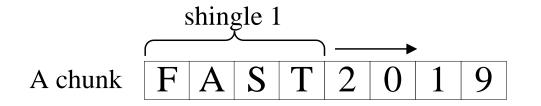
A simple example: extracting 4 features from a string



shingle 1 (S1): FAST shingle 2 (S2): AST2 shingle 3 (S3): ST20 shingle 4 (S4): T201 shingle 5 (S5): 2019 **Rabin fingerprinting:**

 $S1 \longrightarrow R1$ $S2 \longrightarrow R2$ $S3 \longrightarrow R3$ $S4 \longrightarrow R4$ $S5 \longrightarrow R5$

A simple example: extracting 4 features from a string



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Rabin fingerprinting:

 $S1 \longrightarrow R1$ $S2 \longrightarrow R2$ $S3 \longrightarrow R3$ $S4 \longrightarrow R4$ $S5 \longrightarrow R5$

4 Linear transformations:

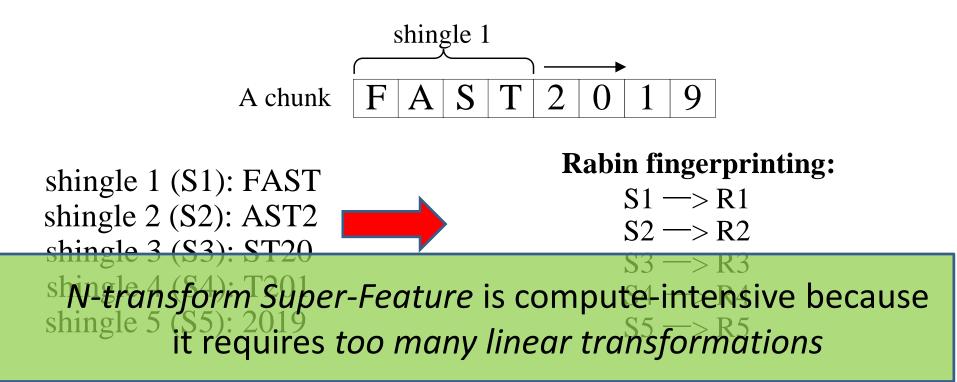
 $\begin{array}{l} \text{R1} \longrightarrow \text{R11}, \ \text{R12}, \ \text{R13}, \ \text{R14} \\ \text{R2} \longrightarrow \text{R21}, \ \text{R22}, \ \text{R23}, \ \text{R24} \\ \text{R3} \longrightarrow \text{R31}, \ \text{R32}, \ \text{R33}, \ \text{R34} \\ \text{R4} \longrightarrow \text{R41}, \ \text{R42}, \ \text{R43}, \ \text{R44} \\ \text{R5} \longrightarrow \text{R51}, \ \text{R52}, \ \text{R53}, \ \text{R54} \end{array}$



Feature extraction

Feature 1:max{R11,R21,R31,R41}
Feature 2:max{R12,R22,R32,R42}
Feature 3:max{R13,R23,R33,R43}
Feature 4:max{R14,R24,R34,R44}

A simple example: extracting 4 features from a string



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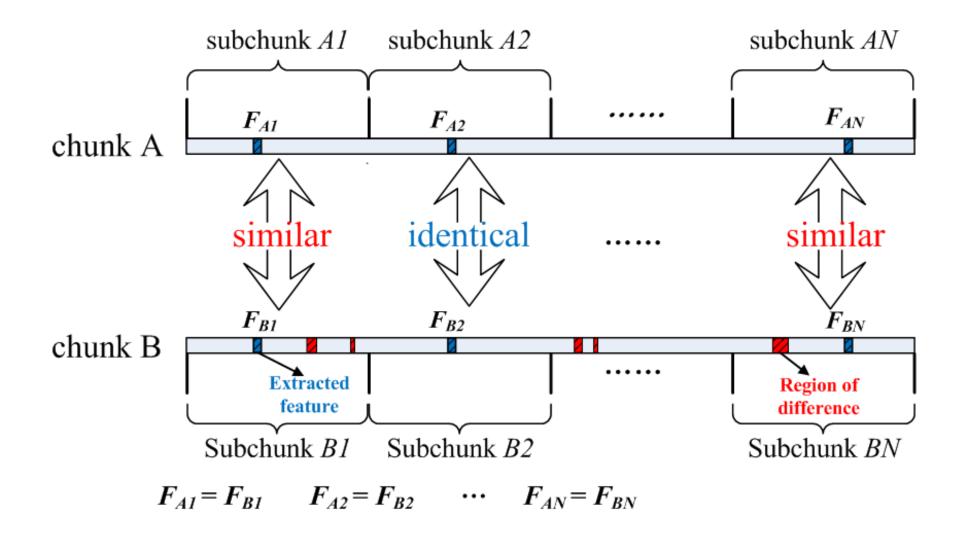
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Observation: Fine-grained Feature Locality



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Datasets	WEB	TAR	RDB	SYN	VMA	VMB
Avg. # of identical subchunks	8.27	9.19	6.86	5.78	5.99	6.34
Avg. # of subchunks owning the same features	10.82	10.97	10.23	10.10	10.04	10.64

All identified chunks are all divided into 12 equal-sized subchunks

Most of the corresponding subchunk pairs in the detected similar chunks have the same features

Design of Finesse

Finesse, a fast resemblance detection approach that exploits the fine-grained feature locality

Step1: Feature extraction

- Dividing a chunk into N equal-sized subchunks, computing Rabin fingerprints for all shingles in the chunk, and selecting the maximum fingerprints in each subchunk as features to obtain N features
- Advantages: Do not require the time-consuming linear transformations for extracting more features, only need to divide the chunk into more subchunks

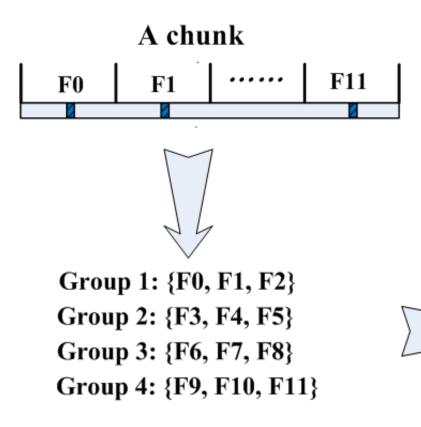
Design of Finesse

Step2: Feature grouping

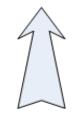
Principle: features in an SF should be extracted from the subchunks distributed uniformly across the chunk.

Design of Finesse

Step2: Feature grouping



SF0: hashing{F1, F5, F7, F9} SF1: hashing{F2, F3, F8, F10} SF2: hashing{F0, F4, F6, F11}



F0 < F2 < F1 F4 < F3 < F5 F6 < F8 < F7 F11 < F10 < F9

Computational Overheads

Approaches	Operations		
N-transform SF	Rabin fingerprinting		
	N linear transformations		
	N conditional branches for feature selection		
Finesse	Rabin fingerprinting		
	1 conditional branch for feature selection		

Computational overhead required to process one shingle.

$DR = \frac{total \ data \ size \ before \ deduplication}{total \ data \ size \ after \ deduplication}$

Name Size DR Workload descriptions WEB 4.21 367 GB 135 days' snapshots of the website: news.sina.com 258 versions of Linux kernel source code. Each version is 1.70 TAR 112 GB packaged as a tar file 100 backups of the redis key-value store database 540 GB 12.25 RDB 176 synthetic backups by simulating file 13.07 330 GB SYN create/delete/modify operations 78 virtual machine images of different OS release 1.61 VMA 117 GB versions, including Fedora, CentOS, Debian, etc 20 backups of an Ubuntu 12.04 VM image in use by a 10.45 VMB 321 GB research group

Datasets

Resemblance Detection Efficiency

Dataset	Approaches	DCR	DCE
WEB	N-transform SF	7.60	0.8749
	Finesse	7.52 (-1.05%)	0.8795 (+0.53%)
TAR	N-transform SF	15.00	0.9516
	Finesse	15.34 (+2.27%)	0.9846 (+3.47%)
RDB	N-transform SF	3.67	0.9129
	Finesse	3.94 (+7.36%)	0.9448 (+3.49%)
SYN	N-transform SF	1.75	0.9326
	Finesse	1.70 (-2.86%)	0.9640 (+3.37%)
VMA	N-transform SF	1.56	0.9088
	Finesse	1.51 (-3.21%)	0.9161 (+0.80%)
VMB	N-transform SF	1.30	0.9093
	Finesse	1.28 (-1.54%)	0.9193 (+1.10%)

 $DCR = rac{total\ size\ before\ delta\ compression}{total\ size\ after\ delta\ compression}$

 $DCE = \frac{chunk \ data \ size \ after \ delta \ compression}{chunk \ data \ size \ before \ delta \ compression}$

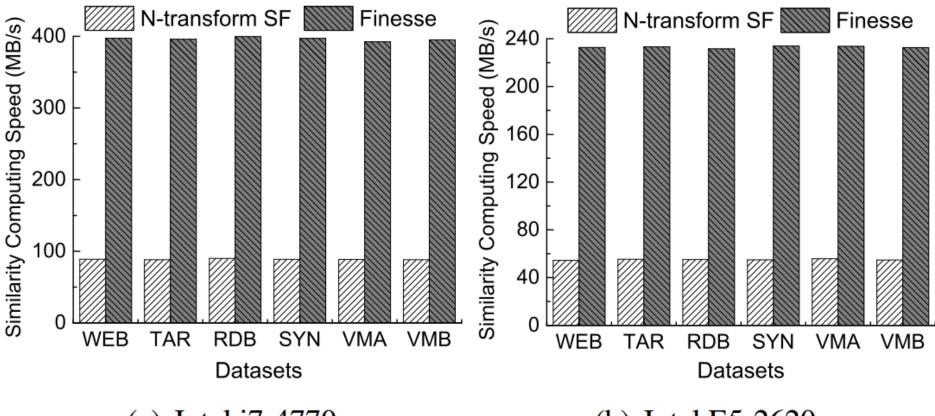
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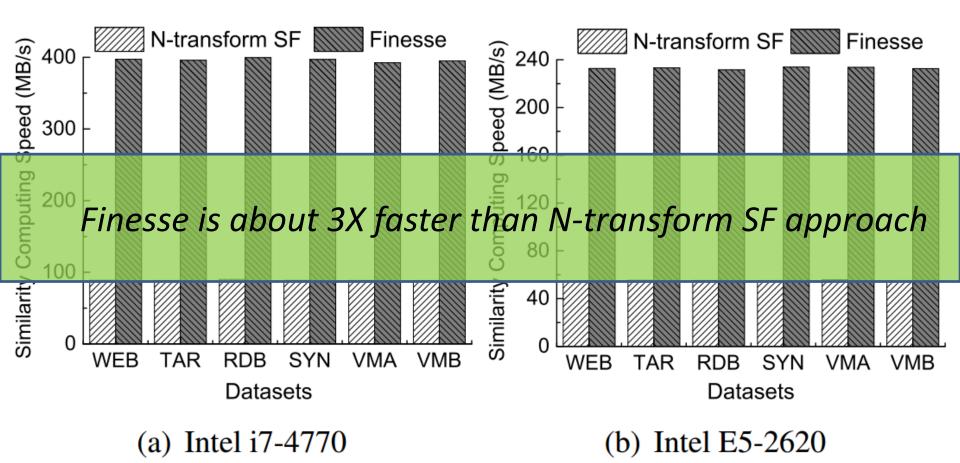
Similarity Computing Speed



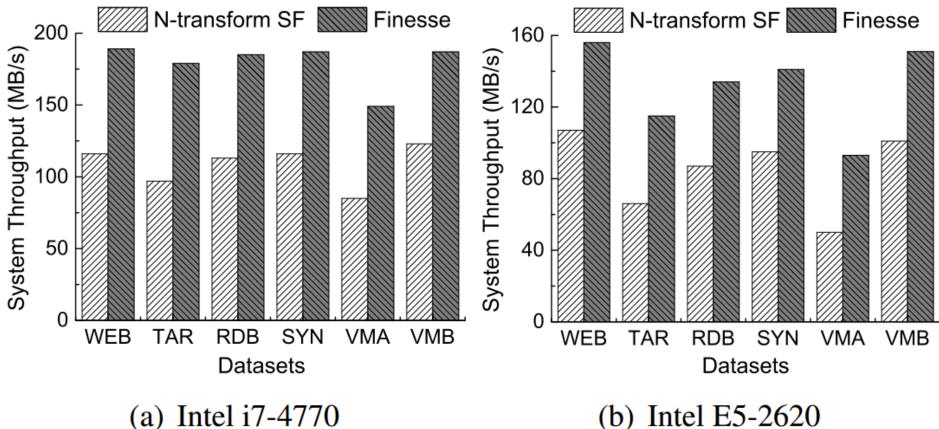
(a) Intel i7-4770

(b) Intel E5-2620

Similarity Computing Speed

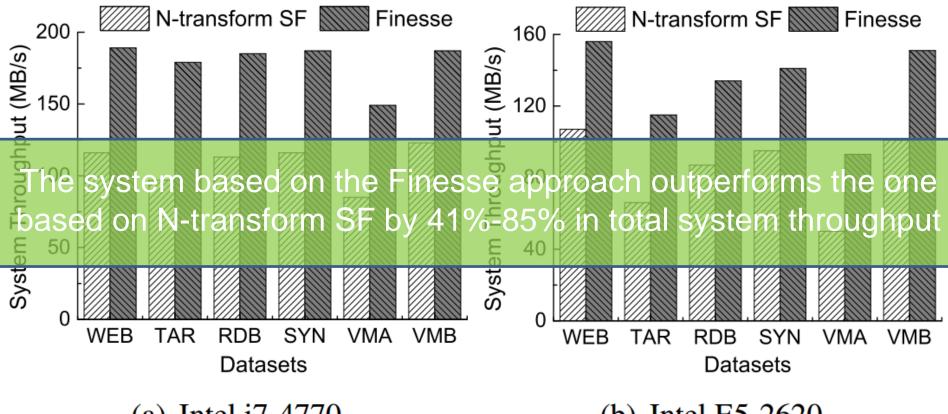


System Throughput



(b) Intel E5-2620

System Throughput



(a) Intel i7-4770

(b) Intel E5-2620

Conclusion

- We observed fine-grained feature locality among similar chunks in backup workloads
- We proposed Finesse, a fast resemblance detection based on fine-grained feature locality
- Our experimental results suggest Finesse runs 3X faster than N-transform SF for resemblance detection

Thank you!

Questions?