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A Dynamic Repository Approach for Small File Management With Fast Access Time on Hadoop Cluster: Hash Based Extended Hadoop Archive

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ABSTRACT Small file processing in Hadoop is one of the challenging task. The performance of the Hadoop is quite good when dealing with large files because they require lesser metadata and consume less memory. But while dealing with enormous amount of small files, metadata grows linearly and Name Node memory gets overloaded hence overall performance of the Hadoop degrades. This paper presents a dual merge technique HB-EHA (Hash Based-Extended Hadoop Archive), that will resolve the small file issue of Hadoop and provide an excellent solution for massive small files that are generated in the health care management applications. The proposed technique merges the small files using two-level compaction, therefore, the size of metadata at the name node gets reduced and less memory will be used. The indexing will be carried out over the archives and files can be accessed after merging in real-time. Index files in the proposed approach can read partially that improves the name node memory usage and also offers the file appending capability in the existing archive. The proposed technique first creates Hadoop archive from the small files and then uses two special hash functions i.e. SSHF (Scalable-Splittable Hash Function) and HT-MMPHF (Hollow Trie Monotone Minimal Perfect Hash Function), SSHF is used to dynamically distribute the archives meta-data to the associated slave index files, and these slave index files will be further written to the final index files, the order of the meta-data in final index file will be preserved by the HT-MMPHF. The evaluation outcome exhibit that the proposed technique is 13% & 17% faster than HDFS with caching enabled and disabled respectively, and 38% & 47% faster than the HAR with caching and without caching, respectively. While comparing with the map file, the proposed technique is 28 $\&$ 35 times faster with caching and without caching, respectively. HB-EHA is a maximum of 40% & 28% faster than the HBAF with and without caching, respectively.

INDEX TERMS Extended Hadoop archive, HAR archive, healthcare small files, HB-EHA, HDFS, HT-MMPHF, map file archive, sequential file, SSHF.

I. INTRODUCTION

Hadoop is a free platform for processing enormous volumes of unstructured and big data. It has a wide range of features when compared to relational databases. Using a master-slave architecture, Hadoop's file system is termed

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Hadoop Distributed File System (HDFS) [1]. Name node is responsible for managing the metadata of the files and have processing capabilities, it acts as a master and coordinates with slave nodes in the HDFS architecture. No processing is done on any of the data nodes that act as slaves, which means that they are just used to store data. In HDFS, files are stored in 128 MB size blocks. HDFS blocks may be whatever size the user desires; by default, it is 128 MB in size, but this

can be changed by the user. To make data availability all the time, HDFS replicates data blocks across several data nodes. To ensure the data availability data blocks are replicated on several data nodes, the value of the replication factor indicate that a single data block is written to many data nodes in the cluster and by default, the value of the replication factor is three. HDFS is well-suited to handle large files since it was designed with this in mind. Small files are generated from various application domains i.e. social media, e-commerce, online business, research and analysis, climate forecasting and educational sites, log records, health care devices and applications, etc. a file is call a small file in HDFS if it's size is less than the default block size of the HDFS. The scope of small files in technology and analytics is important and vast. Due to the excessive metadata at the name node, the overall performance of the HDFS degrades.

There are not only the processing issues with small files in HDFS but also some other issues that will significantly affect the overall performance of HDFS i.e. in case of the several file access request on the Name Node will increase the memory overheads and in case of a massive number of small files, Map Reduce Algorithm [2] requires excessive read and write operation that will take more time to process the small files in comparison with the large files. One more problem with small files is their upload time to the HDFS, the upload time of the small files is comparatively more significant than the large files. There are several feasible alternatives that Hadoop provides to solve the small file problem and its associated disadvantages i.e., HAR Archive [3], Sequential File Format [4], Combine File Input Format [5], and Map File Archive [6]. These solutions will merge the small files and merged files will be saved to the HDFS. By merging these small files, the memory overhead of the Name Node can be minimized but the access performance of the small files will degrade as the side effect of the merging. The meta-data of the small files are stored in the index files and these index files are also stored with the merged file in the HDFS. There are two major categories of the solution provided by the HDFS to solve the small file issue of Hadoop, one is the index-based archive i.e. HAR, Map File etc, and another is without indexing or based on the sequential search approach i.e. Sequential File. To access the content of a small file from the merged file, without considering the caching effect, the entire index file is imported to the memory and then the content of the small file will be recovered from the merged file with the help of meta-data written in the index file. Here meta-data will provide all the information about the small file i.e. the location of the small file in the merged file, length and size of the file, etc. Reading the entire index file in the memory will lead to excess input/output operations that result in increased access time for the small files. In case of the massive number of small files the size of the index file will be large and each time when a small file is accessed, the entire large index file will be loaded into the memory. This process will become very expensive and degrades the overall performance of the HDFS. While considering the caching effect once the index file is imported in the memory its copy will be kept in the client's memory that will improve the access performance of the archives, but the dependency of the access performance on the memory of the client is not a good practice, because client's memory is limited and not sufficient in the case of the vast number of small files.

Most of the healthcare applications/devices generate text or picture files; in our research work, we focused on text files only. Because the size of these text files is smaller than the HDFS block size, they are referred to as ''small files.'' The proposed technique can be interfaced with the various health care devices that are generating the text files. Merging of small files is carried out to utilize the HDFS block completely. These merged small text files can be stored at the Hadoop cluster and can be accessed in real time by using the proposed approach. In this paper, a novel technique, ''Hash Based-Extended Hadoop Archive (HB-EHA)'' is being proposed to manage thousands of small files that are generated in the health care sector. The proposed technique's most significant feature is that it can take Hadoop archives as input and treat them as input files. The proposed technique may also retrieve the archive's meta-data directly without the usage of any caching mechanisms. HT-MMPHF (Hollow Trie Monotone Minimal Perfect Hash Function) [7]–[9] is a recommended method for locating meta-data about a file in an index file and figuring out the index file's size and location in memory. The Scalable-Splittable Hash Function (SSHF) [10]–[12] will dynamically disseminate the meta-data of small files to several index files.

II. RELATED WORK

Jude Tchaye-Kondi *et al.* [13] advocated using hash functions to create a perfect file. To obtain the metadata of a specific file, this approach eliminates the requirement to parse the whole index file. Hadoop's dynamic hashing and orderpreserving operations can efficiently handle thousands of small files. Weipeng Jing *et al.* [14] suggested a dynamic queue method (DQSF) based on the analytical hierarchical process, in which small files are classified and processed according to their size. Jian-Feng Peng *et al.* [15] tackle the HDFS small file issue by merging related small files and caching frequently accessed data to reduce small file access time. By giving the notion of distribution and correlation when merging the files, Xun Cai *et al.* [16] increased the access and storage efficiency of small files. Yanfeng Lyu *et al.* [18] describe an efficient merging approach that considerably reduces the access time for small files by using the concepts of caching and prefetching. X. Fu *et al.* [19] suggested a block replica placement technique for effectively processing small files, in which files are merged according to pre-determined parameters. Qi Mu *et al.* [20] suggested an approach for dealing with small files that is both efficient and effective. The introduction of secondary indexes will increase storage efficiency and minimize Name-Node memory utilization in this system. Tao Wang *et al.* [21] suggested a method based on the behavior of small files access. To build

the association between numerous small files, the probability model is employed. Hui He *et al.* [22] provided a fantastic technique for dealing with the small file issue of the HDFS. Small files are combined in a homogenous manner by balancing the data blocks. This method improves the speed of small files in general. Songling Fu *et al.* [23] presented a novel storage system that makes use of the metadata of small files to effectively manage small files. In compared to existing competing algorithms, the experimental findings suggest that the proposed method would increase the performance of small files by a significant margin. Sharma V.S. *et al.* [24] presented a survey paper on the small file problem of the HDFS. This paper provides quick insight into the possible solution to the small file problem. Sharma V.S. *et al.* [25] evaluated various existing techniques in Hadoop that will provide the solution to the small file problem of the HDFS. N. Alange *et al.* [26] compared various existing techniques for small file problem based on the performance throughput. The CSFC technique, developed by R. Rathidevi *et al.* [27], involves grouping together small files that are connected in some way. HDFS is used for additional processing of the merged small files after clustering. Using the name node, one may store information about all of the bundled files. An innovative strategy for archiving small files was suggested by Y. Chen *et al.* [28], the logical file naming notion is used to identify the files created by the pairing at the name node. W. Wu *et al.* [29] presented a pile structure for sequential file storing that is more efficient. To combine the small files, the worst-fit approach is applied. To retrieve the merged small files from the pile structure, a global index approach is employed. R. Aggarwal *et al.* [30] explored the Hadoop small file issue in-depth and provided systematic literature as well as a Hadoop ecosystem classification. To tackle small file issue, Y. Zhai *et al.* [31] suggested a new archive file. The suggested method retrieves the meta data directly from the index file, reducing access overhead. P. Sobia *et al.* [32] undertook a thorough evaluation of the literature on particle swarm optimization (PSO) techniques for medical illness diagnosis. PSO techniques may also be utilized with a Hadoop cluster to improve response times. Bangyal W.H. *et al.* [33]–[35] suggested three unique pseudorandom initialization techniques, as well as the bat algorithm for population initialization in the PSO algorithm. The concept of pseudorandom initialization can be used to build large index files. Sharma V.S. *et al.* [36] proposed a HBAF technique that solves the small file problem by provding faster access time for small files and reducing the name node memory usage at name node, the technique proposed in this paper is the extension of the HBAF in which Hadoop archives are processed along with dual merge technique.

Table 1 summarize various existing solutions for small file problem of Hadoop. In this table, important research papers from the literature are identified and analyzed. Table 1 address the following issues: what approach is employed, what is the uniqueness of the study that is being done,

and what parameters are being assessed. Table 2 shows a comparison of existing small file approaches based on a number of key parameters, including the proposed approach category, name node memory usage, whether the proposed approach can append new files or not, whether the proposed approach modifies the HDFS or not, whether an extra system is required or not, what is the amount of overheads, and what is the file access complexity of the proposed approach.

III. PROPOSED ARCHITECTURE

A. OVERVIEW OF FAST ACCESS REPOSITORY

In the present scenario, small files have applications in several domains, the application of the small files can be seen almost in every field. There are a number of applications in health care management that will generate massive small files. There is crucial requirement to store and process these small files efficiently [47], [48]. One should be able to store the small files in such a database that will provide reliable storage and faster access to the stored small files. Hadoop is the latest database technology that will provide the distributed reliable storage for the large files. Hadoop is based on the master slave technology; there is one name node and number of data nodes. Data nodes are responsible to store the data in the distributed fashion and name nodes store the metadata information of the data that is stored on the data nodes. In case of the health care management applications, there is a requirement of frequent access to the small files, if Hadoop is chosen as the database technology for health care applications, then the performance of the Hadoop degrades because there is frequent access to small files therefore name node memory usage is higher that will slow down the overall performance of the Hadoop [49]. There is a requirement of a fast access repository for Hadoop that can handle the frequent operations of the small files in health care applications. HB-EHA is a technique that will handle the frequent operations of the small files in health care management applications by merging the small files and then these merged small files are stored in the Hadoop. Hadoop treat these merged files as the larger files and processing of these merged files can be done efficiently in Hadoop. To access the small files from the merged files, HB-EHA uses two special hash functions that will locate the particular small file in the merged file and also maintain the order of the small file while appending more files to the existing archive. The whole process is depicted in figure 1.

B. PROPOSED HB-EHA APPROACH

The proposed solution HB-EHA will increase performance by providing an entry-level archive capability for small files; all small files will be archived, resulting in better memory usage and overall name node performance. Figure 2 shows how the proposed approach builds index files for small files archive's meta-data using two-level hash functions and allowing us for quick access of small files. Parallel numerous archives are formed when combining the small files. In the beginning, a temporary_master_index_file

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TABLE 1. (Continued.) A quick review to the existing approaches for handling massive small files [24].

and a master_name_file are created as a backup. After the slave_index_files have been constructed, this temporary_master_index_file will be deleted. The names of all the small files to be processed are retained in the HB-EHA archive in a permanent file called the master name file. Prior to being appended to a part file, small files may be compressed on the client-side, allowing for quicker processing on the client-side than on the HDFS. During the process of adding files to the part file, a capacity threshold on the size of the part file is defined and regularly checked. Once the threshold limit is reached, a new part file will be created and all other small files will be merged into that new part file. There is need to create new part files when the threshold limit gets over and the remaining small files are merged in the newly created part file. A limit on the index file's size is also

required because in case of a random seek operation, a new connection is established each time to access a file from the various data nodes. A fundamental issue in our strategy is the dynamic distribution of data across many slave_index_files. This dynamic dissemination may be achieved using SSHF, and theses slave index files will be turned into final index files afterward. A small file is appended to a part file and its metadata and name are added to the temporary master index and master name files at the same time. The process of building final index files is accomplished in two phases, the first phase starts with the merging of small files. SSHF will be used to add the data to the corresponding slave index file. This whole process is executed on the client side. In the second stage, the slave index file is sorted. Sorting is maintained using another hash function named HT-MMPHF.

TABLE 2. Comparative study of the small file handling approaches [41].

HT-MMPHF is an order-preserving hash function that will be responsible for sorting metadata at slave_index_files. Finally, slave_index_files are written to the final_index_file.

SSHF is a hash function that belongs to the extendible hashing class [12] that allocates meta-data from small files to slave index files using a dynamic hashing approach. This arrangement involves a hash as the piece string and an arranged tree information structure for the query [50]. There are many slave index files that may be used for storing metadata about a given file. The last two bits of a file name hash value's bit string define the hash. All items with the identical pattern in the last bits will be included in the slave index file. Dynamic additions and deletions on slave index files are made possible by Scalable-Splittable Hashing. It is necessary to partition and generate new slave index files when an existing one has reached its limit. Because of the splitted hash procedure, it is possible to create new slave indexes dynamically. Slave_index_files might be gotten to straightforwardly during the query activity. During the production of slave_index_files and their related final_index_files, the two files are made at the same time. To ensure that the newly generated slave_index_file is in sync with the old and newly splitted slave_index_files during the split hash method, the meta data components are reorganized. It is unimaginable for an assortment of static sort 'p' keys to crash into a number kind 'q' number

utilizing the HT-MMPHF hash function. q ought to be more prominent all the time than or equivalent to P, the worth of the static key. Whenever the upsides of 'q' and 'p' are equivalent, the hash function satisfies the 'minimal' condition and is alluded known as the minimal_perfect_hash_function. A minimal_ perfect_hash_function ought to be utilized to protect the request for keys; this function returns whole number qualities in the request for the static key. This approach guarantees that the lexicographic request of metadata parts in conclusive file documents is safeguarded. Metadata in slave index files is coordinated lexicographically utilizing record name hash esteems. In the final index file, the minimal _perfect_hash_function is developed and put toward the start of the hash esteems. All slave index files will be kept in touch with their final index files before the temporary master index file is erased. In comparison to other hash algorithms of similar complexity, this one's primary advantage is that it has a much lower time and space complexity (logarithmic). The time it takes to access to a given file from the final index files is very less since the meta-data elements are sorted and may be accessed instantly with (Big Oh (1)) complexity [9].

C. ACCESS TIME FOR PROPOSED HB-EHA

The access time for a particular file is the sum of time for accessing metadata and the corresponding content of the file.

FIGURE 1. Overview of the fast access repository for health care applications.

FIGURE 2. Proposed hash based extended hadoop archive (HB-EHA).

First, a client sends the file access request to the Name Node, Name Node finds out the metadata of the requested file and sends results to the client. Based on the information received by the Name Node, the client sends the request to the Data Nodes for content access.

$$
T_{Access\text{-}HDFS} = T_{Access\text{-}Meta} + T_{Access\text{-}Content}
$$
 (1)

$$
T_{Access-Meta} = T_{Req-Meta-NN} + T_{NN-Response}
$$
 (2)

 $T_{\text{Access-Content}} = T_{\text{Req-Read-DN}} + T_{\text{DN-Response}}$ (3)

Where:

- $T_{\text{Req-Read-DN}} =$ Time to read the desired file from the data node
- $T_{DN\text{-}Response}$ = Data Node response time to the client file access request

Since HDFS provides the normal access to the files stored and does not maintain special index files for the lookup of the metadata, whereas HAR and map file maintain the special index files for the lookup of the metadata of the files and require reading and processing whole index file in the memory.

$$
T_{Access\text{-}HDFS} < T_{Access\text{-}HAR\mid Map\text{-}File} \tag{4}
$$

HAR provides two levels of indexing that maintain the metadata of the files, which results in high T_{Access}–_{Meta} in comparison to the map file.

$$
T_{Access-Meta\ (Map-File)} < T_{Access-Meta\ (HAR)} \tag{5}
$$

Eq. 4 states that the access time of native HDFS is lesser than the HAR and map file because these approaches process entire index files while accessing a particular small file. Eq. 5 state that the metadata access time of the map file is lesser than the HAR, because in HAR, multilevel indexing is done therefore the size of index files is more than the map file hence HAR access time is more than the map file. The content in the map file is stored sequentially therefore, to access a random file in the map file requires a very high access time in comparison to the HDFS and HAR.

$$
T_{\text{Access-Content (Map-File)}} > T_{\text{Access-Content (HAR)}}
$$
\n
$$
> T_{\text{Access-Content (HDFS)}}
$$
\n(6)

$$
T_{\text{Access (Map-File)}} > T_{\text{Access (HAR)}} > T_{\text{Access (HDFS)}}
$$
 (7)

Eq. 6 states that a map file requires the highest time to access the content of a small file in comparison to the HAR and HDFS, the reason for this is the sequential access nature of the map file approach. Based on Eq. 4, 5 and 6, Eq. 7 concludes that the map file evolves the highest access time in comparison to the HAR and HDFS. The calculation of the access time of the proposed HB-EHA can be done as follows: In the proposed approach, small files are processed at two levels first, several small files are converted in the archive, and then these archives are processed using the SSHF, this two-level compaction and partial access of the index files from the archives will improve the access time of the proposed HB-EHA at a great extent. Suppose N small files are archived and stored using the Hashing based technique, then the access time for these files can be expressed as

FIGURE 3. Accessing 10 files with cache enabled.

follows:

$$
T_{\text{Access (HB-EHA)}} = N^*(T_{\text{Req-Read-DN}} + T_{\text{DN-Response}}) + \sum_{\text{+} \text{Theq-Meta-NN}} (T_{\text{Req-Meta-NN}} + T_{\text{NN-Response}}) \tag{8}
$$

The proposed HB-EHA reduces the access time in the following aspects: As the small files are merged, archived, and can be dynamically (partially) accessed, it will lead to reduce the communication time between the Name Node and client, ensure fast metadata lookup and decrease the disk overall I/O time & network latency time.

IV. EXPERIMENTAL SETUP AND RESULT ANALYSIS

A cluster of five nodes is being built up to test the proposed HB-EHA technique and other competing archives. The Name Node and Data Node have identical configurations, including an Intel ${}^{\circ}$ coreTM i5-7500 CPU running at 3.40GHz, a 64-bit operating system, and 4 GB of RAM. The operating system in the cluster is Ubuntu 18.04.1 LTS, with open JDK-11.0.4 installed. The most recent version of Hadoop (3.1.3) is installed on all PCs connected to the Hundred MBPS (Backbone) network. The HDFS block size and RF (Replication Factor) are both set to its default values, that is 128 MB and 3, respectively. Five data sets with a variable number of text files were utilized in the testing i.e., 20000, 30000, 60000, 120000. A large number of small text files were selected for processing because, when compared to other file formats, text files provide the best results and take the least amount of time to merge and access. These file ranges between 1 KB to 1 MB in size. ten, fifty, and hundred files are randomly accessed from various archives with and without caching to assess the performance of the different archives. Without caching the performance of the Hadoop archive evolves linear and performs worst in several cases. As shown in the figures, the proposed HB-EHA performs faster than the competitive approaches. With caching, Map File performs the worst, the reason for this is the larger number of the small files and all metadata is loaded in the memory of the client. The reason for the excellent performance of the proposed HB-EHA is as follows:

FIGURE 4. Accessing 50 files with cache enabled.

- 1) To access a particular file from map file and HAR, whole index file is processed therefore, more time is required to access the file, while in the case of the proposed technique, partial index files are processed.
- 2) In map files, small files are de-compressed at two levels, whereas in the case of the proposed technique, only one level of decompression is required.
- 3) For metadata storage, the proposed technique utilizes the data node's memory and relax the Hadoop main memory to facilitate the faster access of the small files.
- 4) In the proposed technique, files are read and written using socket communication means a client can directly interact with the data node's.

A. EVALUATION OF ACCESS TIME FOR RANDOMLY READING 10 FILES FROM VARIOUS ARCHIVES WITH CACHE ENABLED

The proposed HB-EHA is 15 percent to 23 percent (results according to different data sets) quicker in comparison to the original HDFS and 48 percent to 71 percent quicker than the HAR_Archive. As shown in Figure 3 The map file archive will have the highest access time. The access time for HB-EHA is linear as the number of files in the health care data set grows, and it is 17 to 29 times quicker than the map_file_archive. When compared to the HBAF, HB-EHA is 23 percent to 40 percent faster with exceptional case for 120000 files. Due to the dual merge process, the performance of the HB-EHA degrades as the number of files increases in the data set.

B. EVALUATION OF ACCESS TIME FOR RANDOMLY READING 50 FILES FROM VARIOUS ARCHIVES WITH CACHE ENABLED

The proposed HB-EHA is 1 percent to 28 percent quicker than the native HDFS and 32 percent to 46 percent quicker than the HAR_Archive. As shown in Figure 4, again map file archive will have the highest access time. HB-EHA is 24 to 31 times quicker than the map_file_archive, and access time is linear with respect to the number of files in the archive. HB-EHA is 18 percent to 35 percent faster than HBAF. HB-EHA

FIGURE 5. Accessing 100 files with cache enabled.

performs relatively poor for the larger datasets in comparison to the HBAF.

C. EVALUATION OF ACCESS TIME FOR RANDOMLY READING 100 FILES FROM VARIOUS ARCHIVES WITH CACHE ENABLED

As Figure 5 depicts, that proposed HB-EHA is 2 to 13 percent faster than native HDFS and 22 to 38 percent quicker than the HAR_Archive. HB-EHA is 24 to 28 times quicker than the map_file_archive. The access time for the health care data is linear in fashion up to some extent but when the number of files is more than lac then the slight degradation in the performance can be seen, this degradation is due to the high number of read/write operations. HB-EHA is 22 percent to 24 percent faster than HBAF, when accessing more number of files HB-EHF moves towards the stable results in comparison to the HBAF.

D. EVALUATION OF ACCESS TIME FOR RANDOMLY READING 10 FILES FROM VARIOUS ARCHIVES WITH CACHE DISABLED

The proposed HB-EHA is 3 percent to 12 percent quicker than the native HDFS and 22 percent to 38 percent quicker than the HAR_Archive. There is an exceptional case when the number of files is 60,000, in this case, native HDFS performs 6% better than the proposed technique, and the reason for this is the real-time delay in archiving files from the health care data set. The experimental results demonstrate that map file archives take the highest time to access (except in the rare instance of HAR), whereas HB-EHA is 14 to 37 times quicker than map file. As seen in Figure 6, HAR performs the poorest with 1,20,000 small files. This unique behavior is due to HAR's multilayer indexing. While the number of files in the health care data set is in the millions, the notion of multilayer indexing will cause needless delays. HB-EHA is 10 percent to 28 percent faster than the HBAF. In case of the 20,000 dataset the performance of the HBAF and HB-EHA is almost equal and in case of the 1,20,000 data set HB-EHA is 17 percent slower than the HBAF, these are the side effects of the dual merge.

FIGURE 6. Accessing 10 files with cache disabled.

FIGURE 7. Accessing 50 files with cache disabled.

E. EVALUATION OF ACCESS TIME FOR RANDOMLY READING 50 FILES FROM VARIOUS ARCHIVES WITH CACHE DISABLED

The proposed HB-EHA is 3 percent to 22 percent quicker than the native HDFS and 7 percent to 48 percent quicker than the HAR_Archive. The experimental results demonstrate that map_file_archive takes the highest time to access (unless in the rare instance of HAR), whereas HB-EHA is 30 to 36 times quicker than map file. Figure 7 shows that HAR performs the poorest when dealing with 120,000 small files. This unique behavior of HAR is due to a multiple indexing strategies that cause needless delays when dealing with a large number of files in a health care data set. HB-EHA is 5 percent to 24 percent faster than HBAF, HB-EHA performs well in the case of the smaller data sets.

F. EVALUATION OF ACCESS TIME FOR RANDOMLY READING 100 FILES FROM VARIOUS ARCHIVES WITH CACHE DISABLED

The proposed HB-EHA is 5 percent to 17 percent quicker than the native HDFS and 82 percent to 98 percent faster than the HAR archive. The experimental results demonstrate that the map_file_archive has the slowest access time, with HB-EHA being 31 to 35 times quicker. Figure 8 shows that the HAR has a similar uncommon situation owing to the multi-level indexing of the archive's small files. The suggested technique

FIGURE 8. Accessing 100 files with cache disabled.

outperforms the native HDFS by a good margin. HB-EHA is 10 percent to 19 percent quicker than the HBAF.

V. CONCLUSION AND FUTURE WORK

The main motive of HDFS is to govern the ever-increasing volume of big data and ensure that the data is available and accessible at the same time. The HDFS architecture is incompatible with the small files produced by numerous application domains, such as health care, IoT devices, and sensors. In terms of retrieving small files and managing their meta-data, the HDFS is complicated. Managing small files efficiently while simultaneously lowering access time and memory usage at the name node is a pressing issue. Several scholars have researched in this topic and provided several ways for handling small files effectively. Most of the methods presented minimize name node meta-data use by transferring the indexing process to the client-side, although these techniques lag when assessed in terms of access time. There is a need for a technique that can receive small file archives as input, process them as files, and manage the index files in such a manner that the name node memory use is reduced while the small files are accessed quickly. The proposed HB-EHA works better than the HAR, Map File and HBAF. When caching is removed, the HAR's access time increases dramatically due to its multi-level indexing, but the proposed solution is unaffected by whether or not caching is enabled or disabled. The proposed solution would allow relatively quick meta-data access for small files, as well as the ability to append files after the archive has been created. For the goal of caching, data nodes are employed; this notion reduces memory load on the client-side, resulting in faster access times for small files. It will be feasible to get to the substance of small files all the more rapidly on account of the utilization of SSHF for meta-data inclusion and an order-preserving hash function (HT-MMPHF) for keeping everything under the control of meta-data inside final index files. It was demonstrated that the proposed approach is lot quicker than the HAR and map file. While caching is enabled, HB-EHA is 38 and 28 time quicker than HAR and map files, respectively. While caching is disabled, HB-EHA is 47 and 35 times quicker than the HAR and map files, respectively. When compared to the HBAF, HB-EHA is a

maximum of 40% to 28 % faster, but HB-EHA show the good performance for the smaller data sets. Due to the dual merge of the HB-EHA, the process of indexing becomes complex and HB-EHA performs poorly. The proposed approach can be more effective if a few additional features are added as future work i.e. other hash function combinations might be utilized to improve the proposed approach, client-side memory use might be diminished and it is also important to recognize the numerous parts of our approach that are yet utilizing client memory. Deletion of small files and accessing small files by the file magnitude is still a future work in our approach.

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