



Enhanced Gap Sequencing Shell Sort

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Abstract:Sorting algorithms are used for arranging a list of numbers or characters in an ascending or descending order. Many sorting algorithms have been proposed for sorting a given sequence, some important algorithms of them are Bubble Sort, Insertion Sort, Selection Sort, Shell sort and many more. Sorting algorithms put elements in a certain order such as numerical or lexicographical order. Shellsort is the fastest algorithm in comparison to bubble, insertion and selection. Shell sort is an enhanced version of insertion sort. It reduces the number of swaps of the elements being sorted to minimize the complexity and time as compared to insertion sort. Shell sorting algorithm sorts the elements according to the gap sequences. The operations depend on the used gap sequences (many gap sequences have been proposed). In this paper we analyze the algorithm by using the following gap sequence: $N_1; N_2; N_3; \dots; 1$, where

$$N_1 = \text{floor}(3n/4);$$

$$N_2 = \text{floor}(3N_1/4);$$

$$N_3 = \text{floor}(3N_2/4);$$

.....;

.....;

$N_k = \text{floor}(3N_{k-1}/4)$ and so on until the value of N_k becomes one. Here 'n' is the number of elements to be sorted.

Keywords: Algorithm, Shell, Sorting, Comparison.

I. INTRODUCTION

Shell sort is the algorithm that has been the focus of research. Shell sort is introduced by Donald L. Shell in 1959 [1]. Shell sorting algorithm is the generalization of insertion sort algorithm [2]. It is easy to implement, and thus a practical choice for sorting moderate-sized lists. Shell sort has been proposed to improve the average running time of the insertion sort, which is $\Theta(n^2)$ [2]. It has been observed that Shell sort is a non-stable in-place sort. Shell sort improves on the efficiency of insertion sort by quickly shifting values to their destination [2]. Many different gap sequences may be used to implement the shellsort [1]. Typically, the array is sorted with large gap sequencing, then the gap sequencing is reduced, and the array is sorted again. On the final sort, gap sequencing is one.

In this paper work, I use the tools developed in some previous published papers on shell sort to analyze the gap sequence: $N_1; N_2; N_3; \dots; 1$, where

$$N_1 = \text{floor}(3n/4);$$

$$N_2 = \text{floor}(3N_1/4);$$

$$N_3 = \text{floor}(3N_2/4);$$

.....;

$N_k = \text{floor}(3N_{k-1}/4)$ and so on until the value of N_k becomes one. Here 'n' is the number of elements to be sorted. In analyzing Enhanced Gap Sequencing Shell Sort, it turns out that the value of the gap sequence depends on the number of elements to be sorted. First, we find $N_1 = \text{floor}(3n/4)$, after this all other gaps $N_2; N_3; \dots; 1$; can be found by putting N_1 in place of 'n' to get N_2 and N_2 in place of 'n' to get N_3 and so on until we get the value 1. For example, if we want to sort 125 numbers of elements, then $N_1 = \text{floor}(3(125)/4)$ because here $n=125$, so $N_1=93$. Now find $N_2 = \text{floor}(3N_1/4)$ which is 69, and

$N_3=51, N_4=38, N_5=28, N_6=21, N_7=15, N_8=11, N_9=8, N_{10}=6, N_{11}=4, N_{12}=3, N_{13}=2, N_{14}=1$. So the gap sequence for sorting 125 elements is 93, 69, 51, 38, 28, 21, 15, 11, 8, 6, 4, 3, 2, 1.

In this paper, I made the list of comparison of "Enhanced Gap Sequencing Shell sort" with Insertion sort and Shell sort. In Shellsort the numbers of swaps are reduced as compared to Insertion sort and in "Enhanced Gap Sequencing Shell Sort" the numbers of swaps are further reduced as compared to Shellsort. [3]

A. Insertion Sort:

Insertion Sort algorithm sorts the elements by inserting them into their proper position in the final sorted list. Insertion sort keeps making the left side of the array sorted until the whole array is sorted. It sorts the values seen far away and repeatedly inserts unseen values into the array into the left sorted array. It is the simplest of all sorting algorithms. Although it has the same complexity as Bubble Sort ($\Theta(n^2)$), the insertion sort is a little over twice as efficient as the bubble sort. The advantage of Insertion Sort is that it is relatively simple and easy to implement.

In the following example, the number of swaps calculated to sort the elements using Insertion Sort algorithm. In this example, the list contains 38 elements, which are unsorted, the Insertion sort is applied in order to find the total number of swaps required to sort the elements in the increasing order.

a. Unsorted list:

22, 12, 53, 94, 27, 59, 50, 39, 14, 88, 35, 3, 115, 3, 4, 230, 29, 84, 6, 2, 102, 14, 54, 5, 3, 87, 67, 16, 43, 73, 19, 27, 64, 16, 4, 2, 85, 51, 34.

After applying the Insertion sort on this array, the number of swaps calculated for sorting it, are 367.

B. Shell Sort:

Shell sort sorting algorithm is introduced by the Donald L. shell in 1959[1,2]. Shell sort works by comparing elements that are distant rather than adjacent elements in an array or list where adjacent elements are compared. Shellsort uses a gap sequence $g_1; g_2; \dots; g_n$, called the increment sequence. Any increment sequence is fine as long as $g_1 = 1$ and some other choices are better than others. Shellsort makes multiple phases through a list and sorts a number of equally sized sub lists using the insertion sort. Shellsort is also known as diminishing increment sort[2].

The distance between comparisons decreases as the sorting algorithm runs until the last phase in which adjacent elements are compared. After each phase and some increment g_h , for every i , we have $a[i] \leq a[i+g_h]$ all elements spaced g_h apart are sorted. The file is said to be g_h -sorted. The size of the sub-lists, which are to be sorted gets larger with each phase through the list, until the sub-list consists of the entire list. (Note that as the size of the sub list increases, the number of sub-lists to be sorted decreases.) This arrangement makes the insertion sort to run for an almost-best case with a complexity that approaches $O(n)$.

The elements contained in each sub-list are not contiguous, rather, if there are i sub-lists then a sub-list is composed of every i -th element. For example, if there are 4 sub lists then the first sub-list would contain the elements located at positions 1,5,9 and so on. The second sub-list would contain the elements located at positions 2,6,10, and so on; while the fourth sub-list would contain the items located at positions 4,8,12, and so on.[3]

The efficiency of the algorithm is depends on the size of the sub-lists used for each iteration. Along with the benefit of being robust, Shellsort is a complex algorithm and not nearly as efficient as the merge, heap, and quicksorts. The shell sort is still significantly slower than the merge, heap, and quick sorts, but its relatively simple algorithm makes it a good choice for sorting lists of less than 5000 items unless speed is important. It is also an excellent choice for repetitive sorting of smaller lists. It has been observed that Shell sort is a non-stable in-place sort. Shell sort improves the efficiency of insertion sort by quickly shifting values to their appropriate position. Average sort time is $O(n^{1.25})$, while worst-case time is $O(n^{1.5})$ [3].

Knuth has experimented with several values and recommends that gapping 'h' for an array of size N be based on the following formula: Let $h_1 = 1; h_{s+1} = 3h_s + 1$, and stop with h_t when $h_{t+2} \geq N$ [3]. Thus, values of h are computed as follows:

$$\begin{aligned} h_1 &= 1 \\ h_2 &= (3 \cdot 1) + 1 = 4 \\ h_3 &= (3 \cdot 4) + 1 = 13 \\ h_4 &= (3 \cdot 13) + 1 = 40 \\ h_5 &= (3 \cdot 40) + 1 = 121 \\ h_6 &= (3 \cdot 121) + 1 = 364 \end{aligned}$$

To sort 125 items we first find an 'h_s' such that $h_s \geq 125$. For 125 items, h_6 is selected. The final value h_6 is two steps lower, or h_4 . Therefore sequence for the values of 'h' will be 40,13,4, 1. Once the initial 'h' value has been determined, subsequent values may be calculated using the formula

$$h_{s-1} = \text{floor}(h_s/3).$$

Now we apply the shell sort for calculating the number of swaps to sort the same problem (as discussed in Insertion sort) by using Knuth's gap sequences.

a. Unsorted list:

22,12,53,94,27,59,50,39,14,88,35,3,115,3,4,230,29,84,6,2,102,14,54,5,3,87,67,16,43,73,19,27,64,16,4,2,85,51,34.

After applying the shell sort using Knuth's gap sequences on this array of numbers, the number of swaps calculated for sorting it, are **170**.

C. Enhanced Gap Sequencing Shell Sort:

Enhanced Gap Sequencing Shell Sort introduced a new way to find the gap sequences to sort the large list of elements rapidly. Enhanced Gap Sequencing Shell Sort algorithm also works in same fashion as the previous existing versions of the Shell sort algorithm. The only difference is the way to choose the more efficient gap sequence, which is a key step for the algorithm to be more effective and better. Calculating the value of the gap sequence h_i in conventional Shell sort is a key step in execution of the algorithm.

In conventional Shell sort, given by Knuth, the value of h_i is found by the following formula:

$$h_1 = 1; h_{s+1} = 3h_s + 1, \text{ and stop with } h_t \text{ when } h_{t+2} \geq N.$$

By using this existing formula Shell sort reduces the number of swaps up-to 50% as compared to that of Insertion sort.

Enhanced Gap Sequencing Shell sort mainly focuses to improve the efficiency of the previous existing shell sort algorithms. It can be achieved by choosing the appropriate values of gap sequences, which can reduce the number of swaps. In conventional Shell sort, the gap sequences are small, so they divide the list into large number of sub-steps, for example, if $n=100$ then the gap sequence is (13,4,1), which means to sort the elements, it first divide the list into 8-sublists of size 13, then 25-sublists of size 4. It makes algorithm to swap many elements many times to place them into right place. So, if we increase the size of gap sequence, it divides the list into less number of sub-lists of larger size and elements are not needed to swap many numbers of times.

Enhanced Gap Sequencing Shell Sort introduces a new mechanism to calculate the value of gap sequence. The formula for calculating gap sequence is as follows:

The gap sequence is of the form $N_1, N_2, N_3, \dots, 1$, where $N_1 = \text{floor}(3n/4)$;
 $N_2 = \text{floor}(3N_1/4)$;
 $N_3 = \text{floor}(3N_2/4)$;
;

$N_k = \text{floor}(3N_{k-1}/4)$ and so on until the value of N_k becomes one. Here 'n' is the number of elements to be sorted. Thus to sort 125 numbers of elements, the gap sequence is calculated as follows: here $n=125$

$$\begin{aligned} N_1 &= \text{floor}(3 \cdot 125/4) = 93 \\ N_2 &= \text{floor}(3 \cdot 93/4) = 69 \\ N_3 &= \text{floor}(3 \cdot 69/4) = 51 \\ N_4 &= \text{floor}(3 \cdot 51/4) = 38 \\ N_5 &= \text{floor}(3 \cdot 38/4) = 28 \\ N_6 &= \text{floor}(3 \cdot 28/4) = 21 \\ N_7 &= \text{floor}(3 \cdot 21/4) = 15 \\ N_8 &= \text{floor}(3 \cdot 15/4) = 11 \\ N_9 &= \text{floor}(3 \cdot 11/4) = 8 \\ N_{10} &= \text{floor}(3 \cdot 8/4) = 6 \end{aligned}$$

$$N_{11} = \text{floor}(3*6/4) = 4$$

$$N_{12} = \text{floor}(3*4/4) = 3$$

$$N_{13} = \text{floor}(3*3/4) = 2$$

$$N_{14} = \text{floor}(3*2/4) = 1$$

So the gap sequence for sorting 125 elements is (93,69,51,38,28,21,15,11,8,6,4,3,2,1). But unconventional Shell sort the gap sequence for 125 elements is (40, 13, 4, 1).

Now we apply the Enhanced Gap Sequencing shell sort for calculating the number of swaps to sort the same problem as discussed in Insertion sort and Shell Sort).

a. Unsorted list:

22,12,53,94,27,59,50,39,14,88,35,3,115,3,4,230,29,84,6,2,102,14,54,5,3,87,67,16,43,73,19,27,64,16,4,2,85,51,34.

Here n=38,so

$$N_1 = \text{floor}(3*38/4) = 28$$

$$N_2 = \text{floor}(3*28/4) = 21$$

$$N_3 = \text{floor}(3*21/4) = 15$$

$$N_4 = \text{floor}(3*15/4) = 11$$

$$N_5 = \text{floor}(3*11/4) = 8$$

$$N_6 = \text{floor}(3*8/4) = 6$$

$$N_7 = \text{floor}(3*6/4) = 4$$

$$N_8 = \text{floor}(3*4/4) = 3$$

$$N_9 = \text{floor}(3*3/4) = 2$$

$$N_{10} = \text{floor}(3*2/4) = 1$$

So the gap sequence for 38 elements is (28,21,15,11,8,6,4,3,2,1). After applying the Enhanced Gap Sequencing shell sort on this array of numbers, the number of swaps calculated for sorting it, are 67.

II. COMPARISON OF ABOVE DISCUSSED THREE TECHNIQUES

Now the comparison for the three techniques is made here for the same problem.

Table 1: Comparison

Sr. No.	Insertion Sort	Shell Sort	E.G.S. Shell Sort
01	367	170	67

It is apparent that Shell sort reduces the number of swaps up to half as compared to the number of swaps in Insertion sort and Enhanced Gap Sequencing Shell Sort reduces the number of swaps further up to less than half as compared to the number of swaps in Shell Sort, thus improving the efficiency of the algorithm.

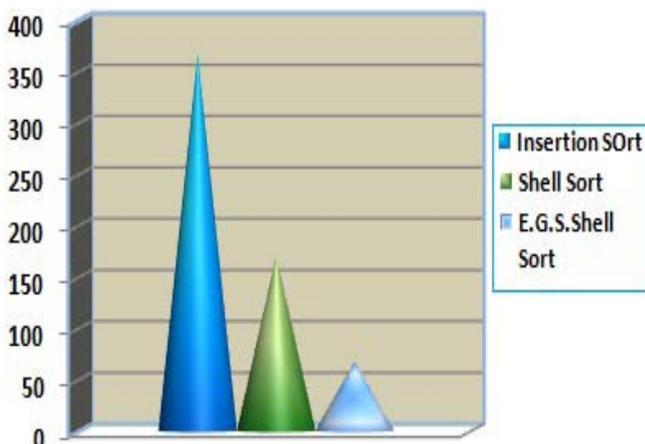


Figure 1: Comparison of Sorting Techniques

III. DETAIL DISCUSSION OF RESULTS FOR MORE PROBLEMS

It is necessary to execute a detailed comparison of all three algorithms by applying the algorithms on a wider variety of data, in order to ensure and establish results concretely.

A. For 25 Elements:

0,1,13,4,14,10,19,11,22,27,14,17,13,20,19,15,16,19,18,19,18,17,21,20,14.

Number of swaps in Insertion Sort:83

Number of swaps in Shell Sort:47

Number of swaps in E.G.S.Shell Sort:22

B. For 50 Elements:

0,1,21,4,23,16,32,16,40,50,21,28,16,38,33,16,20,39,35,37,38,40,25,29,37,35,35,33,30,31,31,31,31,31,34,32,35,29,3,2,32,30,35,23,37,40,34,26,42,30,26.

Number of swaps in Insertion Sort:464

Number of swaps in Shell Sort:127

Number of swaps in E.G.S.Shell Sort:63

C. For 100 Elements:

1,1,38,6,40,27,59,26,75,98,35,50,21,73,61,18,29,80,70,75,79,88,36,51,84,77,81,74,53,61,64,60,58,66,39,52,40,62,5,5,55,58,51,66,49,48,54,59,49,56,58,57,51,58,54,55,55,56,56,56,58,59,53,59,59,58,61,58,66,57,44,70,59,70,71,65,38,62,60,70,70,70,69,43,35,34,43,59,55,86,81,59,58,50,75,86,41,4,5,25,25,96.

Number of swaps in Insertion Sort:2154

Number of swaps in Shell Sort:419

Number of swaps in E.G.S.Shell Sort:179

D. For 200 Elements:

2,1,71,9,76,48,113,45,145,193,62,95,32,142,117,22,45,162,138,152,162,184,58,93,180,161,173,155,98,122,130,12,0,113,138,50,92,53,129,102,104,114,85,153,76,66,96,129,6,110,122,119,51,137,81,101,94,93,113,115,85,75,133,87,9,1,98,84,99,71,105,141,70,99,76,77,90,135,97,100,89,89,91,93,118,123,123,150,104,107,89,93,104,105,108,99,96,110,1,08,112,112,99,106,106,106,107,105,105,106,107,106,109,1,06,111,101,111,102,104,102,101,105,105,116,112,95,95,10,2,89,87,86,96,106,116,96,118,99,107,125,103,114,89,114,9,6,133,132,119,75,94,95,89,93,93,87,62,114,101,118,59,155,61,111,134,101,107,92,83,97,72,115,167,65,79,44,112,126,48,87,98,79,146,134,48,170,38,89,171,52,71,79,38,108,142,76,139,131,62,22,32,152,123,30,121.

Number of swaps in Insertion Sort:10180

Number of swaps in Shell Sort:886

Number of swaps in E.G.S.Shell Sort:544

E. For 350 Elements:

3,2,121,14,129,81,193,75,250,335,103,162,48,247,202,28,70,284,241,266,286,328,91,157,323,286,312,276,166,21,2,229,210,195,246,67,153,71,230,173,178,199,137,282,115,93,160,233,90,191,219,213,51,255,121,171,152,150,201,20,6,127,101,254,132,141,159,120,161,79,179,289,70,161,87,8,8,128,281,150,161,118,119,122,130,231,257,259,226,173,1,86,94,113,173,178,198,137,113,215,206,245,243,106,189,1,82,200,223,143,137,141,156,190,128,197,115,242,141,222,198,208,210,187,189,140,160,224,218,194,231,235,233,206,182,160,202,160,194,180,151,186,170,204,171,193,151,15,3,168,212,192,191,195,191,191,194,210,176,184,174,206,1,56,202,178,169,182,180,185,188,183,190,178,167,189,186,

191,180,178,188,183,181,182,178,179,182,180,181,181,183,179,180,180,176,182,185,178,185,185,175,169,170,189,184,167,184,188,166,196,197,190,198,175,158,194,204,154,197,199,208,180,156,153,198,173,152,169,125,180,154,152,161,171,177,137,135,224,195,158,223,162,199,224,195,210,181,238,122,242,233,213,176,192,218,187,230,222,193,251,187,188,229,239,194,111,250,200,258,154,170,252,251,236,232,234,223,208,267,137,245,112,272,233,113,102,137,126,107,160,223,85,163,150,166,147,222,162,246,73,279,256,173,99,270,209,255,79,184,205,281,145,219,243,68,276,159,243,106,168,189,276,147,84,141,70,53,214,196,256,315,262,126,134,165,50,37,276,325,152,103,168,318,234,61,38,323,77,237,214,152,68,179,250,88,45,57.
 Number of swaps in Insertion Sort:29068
 Number of swaps in Shell Sort:2232
 Number of swaps in E.G.S.Shell Sort:1079

F. For 500 Elements:

5,3,172,19,183,114,274,104,355,478,145,228,65,352,287,35,95,406,344,381,410,471,124,221,465,412,450,398,233,303,329,300,278,354,84,214,89,331,245,252,283,189,413,155,120,223,337,115,217,316,306,52,373,162,240,210,206,288,296,169,126,375,176,191,220,156,223,87,253,436,70,222,2,97,100,166,427,203,221,148,154,168,345,390,395,336,242,266,99,132,242,250,287,175,131,320,303,378,374,112,272,259,293,338,181,170,176,206,275,148,289,120,385,173,343,292,316,320,270,275,165,210,355,342,286,374,383,380,316,258,204,310,202,290,254,177,270,228,320,229,290,170,175,219,349,290,288,301,289,290,302,360,239,268,230,353,158,343,246,205,264,254,279,294,271,308,243,167,315,293,339,248,231,328,279,266,287,212,226,317,190,324,272,194,306,288,280,314,255,227,279,231,237,288,315,307,225,245,303,246,237,298,217,218,237,220,268,300,233,218,300,233,231,220,258,286,288,237,264,285,267,226,257,277,278,270,262,258,282,282,233,249,267,237,263,249,240,251,247,256,240,270,242,245,249,256,254,250,255,251,252,255,252,255,255,255,256,257,255,259,257,261,254,256,263,264,263,263,264,263,261,271,249,268,243,275,267,241,238,246,243,238,251,267,230,251,248,252,246,269,251,277,221,289,282,254,228,288,267,283,219,257,265,295,242,271,281,210,295,247,282,225,251,260,298,241,213,238,206,198,271,263,291,319,295,230,234,249,192,185,303,328,242,217,249,326,284,194,182,331,202,286,274,241,195,255,294,205,181,187,220,332,238,275,199,190,277,254,174,216,228,355,226,210,283,189,242,353,187,234,155,192,348,211,160,354,337,358,212,280,318,255,351,251,382,312,306,327,188,349,309,174,225,366,126,239,298,281,137,390,255,151,301,148,211,303,399,398,289,161,379,279,123,149,103,247,175,187,101,393,272,325,116,251,93,225,245,382,248,301,184,198,347,232,276,315,413,89,205,99,130,124,257,418,113,371,178,364,312,115,383,235,247,206,261,359,355,285,114,346,138,249,176,279,360,402,184,263,110,359,337,431,378,79,50,117,49,73,312,272,146,286,303,444,62,78,313,406,335,461,205,426,42,154,354,192,412,221,196,65.
 Number of swaps in Insertion Sort:62679
 Number of swaps in Shell Sort:3456
 Number of swaps in E.G.S.Shell Sort:1613

G. For 1000 Elements:

10,5,339,36,361,222,542,202,705,953,282,450,119,701,569,55,177,814,687,763,824,951,234,435,942,832,912,803,459,605,659,600,552,714,141,417,150,666,482,497,565,361,848,287,210,435,685,197,541,638,618,53,766,296,470,403,395,578,598,309,212,777,323,358,425,277,430,114,500,928

,71,427,132,137,293,915,381,424,246,247,260,292,724,835,848,704,473,532,117,197,471,492,586,302,188,671,628,821,812,133,549,514,603,724,307,279,294,374,558,214,596,135,859,280,747,606,673,686,546,561,249,376,789,754,593,851,879,871,685,513,352,669,345,610,500,264,549,420,708,424,616,233,247,387,809,617,612,656,617,621,663,859,450,549,419,845,166,814,472,323,538,502,592,648,562,703,460,169,732,651,833,477,407,798,601,548,637,324,381,766,227,803,579,235,729,651,617,777,502,370,620,386,414,665,803,767,347,450,760,456,407,741,290,291,398,296,580,774,368,268,789,355,339,263,522,725,740,370,572,732,594,268,515,688,702,639,569,532,769,773,265,432,631,286,604,418,298,440,373,508,250,768,242,286,373,528,465,361,485,324,354,463,256,487,483,346,314,466,733,288,448,270,588,540,301,306,351,364,361,395,436,283,620,344,679,284,380,669,693,608,633,672,554,416,714,545,571,538,576,423,543,380,715,320,366,521,655,345,456,377,683,501,467,340,565,445,408,686,359,541,412,618,525,494,369,555,643,562,658,679,462,487,408,334,403,575,565,526,665,678,394,338,539,595,521,354,448,636,659,356,614,449,473,535,618,508,439,595,635,622,565,383,534,477,595,608,474,508,625,563,545,369,547,567,470,592,525,385,591,533,627,581,400,557,613,398,418,397,552,481,443,507,413,511,386,453,459,441,567,425,460,575,532,415,611,519,473,486,596,406,506,582,474,581,537,474,410,412,485,566,429,492,586,569,595,511,551,544,590,433,497,470,577,508,585,521,511,447,509,486,537,530,467,516,498,482,445,569,525,563,551,552,505,451,553,468,530,472,489,547,469,511,508,519,504,479,481,499,539,485,532,507,523,501,485,477,519,504,532,488,492,477,532,536,525,534,530,498,504,518,502,501,486,524,522,501,493,499,491,509,495,518,511,501,508,499,507,508,511,504,510,503,506,506,505,506,507,505,507,505,502,509,508,506,499,503,512,506,507,502,519,498,493,503,516,526,515,528,510,530,484,518,516,499,511,517,537,507,541,528,541,493,526,483,507,542,526,469,519,499,535,465,462,482,468,528,537,523,514,466,538,449,463,448,455,558,451,542,453,551,544,464,438,559,498,467,482,540,580,488,445,488,545,484,493,505,468,455,586,438,516,575,469,528,474,507,449,528,504,575,472,517,504,547,440,575,533,525,514,469,417,460,564,579,564,604,556,515,529,583,594,394,516,620,483,570,577,467,584,571,506,380,570,449,538,632,558,434,627,641,440,429,403,402,514,596,574,459,480,377,385,631,450,579,631,362,395,505,543,365,644,408,418,407,405,578,553,664,345,571,513,673,462,344,405,538,511,677,345,661,566,616,572,366,453,393,514,510,401,684,650,494,620,452,409,394,381,576,694,324,617,483,678,698,493,479,448,591,323,547,502,704,459,564,424,457,403,669,585,609,330,486,559,642,660,419,567,603,717,305,300,643,681,409,383,358,673,388,558,669,605,496,453,513,355,502,671,299,512,716,354,690,726,390,670,561,628,589,721,353,527,679,351,390,456,583,742,600,543,429,747,624,372,270,505,490,448,455,307,706,683,246,394,732,586,706,493,653,430,601,277,220,555,277,673,690,602,514,252,444,418,648,406,440,611,431,236,567,239,515,216,668,261,504,652,620,699,658,513,498,714,272,598,378,513,211,256,251,307,498,215,242,617,489,627,346,300,197,678,510,820,194,368,263,509,311,638,779,686,544,740,415,468,492,497,235,470,803,474,283,681,692,747,429,654,674,793,194,310,332,710,606,573,356,662,659,420,635,756,187,540,136,382,372,704,674,149,366,314,199,164,455,340,202,748,692,401,723,767,838,463,788,711,327,223,671,905,510,363,871,337,585,854,850,798,597,531,467,225,604,490,129,202,566,92,859,794,443,313,901,354,280,681,262,768,559,558,

738,811,885,172,699,654,661,402,228,246,806,714,844,247,568,440,338,844,657,341,193,293,597,750,731,202,780,129,359,502,476,485,118,921,726,774,272,65,391,142,799,200,938,571,782,979,576,115,546,430,147,836,394,886,256,905,672,152,730,913.

Number of swaps in Insertion Sort:243764

Number of swaps in Shell Sort:8744

Number of swaps in E.G.S.Shell Sort:3934

Table 2: Comparison of all three sorts

Cases	Insertion Sort	Shell Sort	E. G. S. Shell Sort
3.1	83	47	22
3.2	464	127	63
3.3	2154	419	179
3.4	10180	886	544
3.5	29068	2232	1079
3.6	62679	3456	1613
3.7	243764	8744	3934

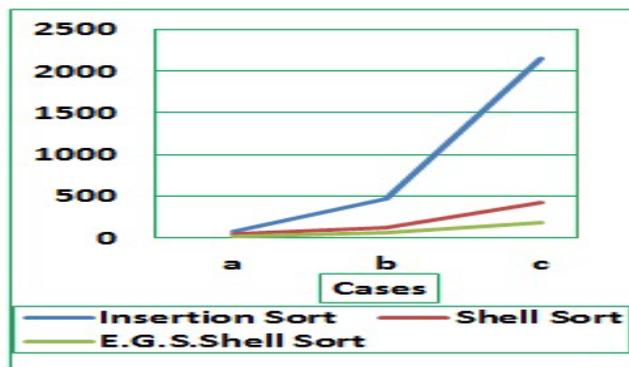


Figure 2: Comparison graph based on cases

The Enhanced Gap Sequencing Shell Sorting algorithm is a good approach towards achieving the excellence in the algorithms to provide the more efficient solutions. This has been achieved by decreasing the number of swaps required to sort the list of elements.

The results of above solved problems show that the new algorithm provides a much efficient way to sort the elements and hence causes to save the computational resources. It has been observed that the Enhanced Gap Sequencing Shell Sorting algorithm can solve the problem in almost 60 times less swaps as compared to insertion sort and in almost half swaps as compared to the traditional shell sorting algorithm. Figure 3 shows the detailed overview of the number of swaps required to sort different number of elements.

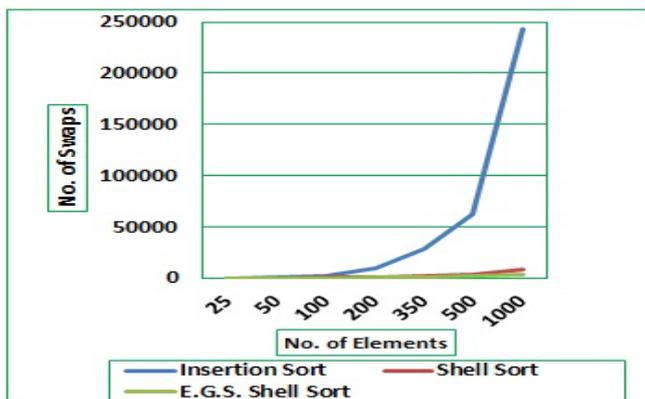


Figure 3: Graph showing elements-swap ratio

IV. ANALYSIS

In the paper “Analysis of Shellsort and Related Algorithms”, Robert Sedgewick [4] has described an open problem, “Are there increment sequences that perform better than known ones in practice?”, for performance issues and claims that finding a sequence that leads to running times 25% lower than the best known certainly would be of practical interest, we can reduce the running time by reducing the number of comparisons for the algorithm. In our proposed algorithm, we have reduced the number of comparison up-to 40% to 50% in some ideal cases but in many cases up-to 20% to 30%. We compare our algorithm with insertion sort and shell sort and get some interesting results as can be seen in the above given graphs and tables. Marcin Ciura [5] in his paper “Best Increments for the Average Case of Shell sort”, shows the result for 128 elements where data get sorted in 535 (approx.) swaps but in our case 200 elements takes 544 swaps to get sorted. Thus the proposed algorithm is better for sorting.

V. CONCLUSION

This research paper focuses on an enhancement and improvement in existing sorting algorithms. The traditional Shell sort algorithm results an average number of comparisons of elements but it does not give minimum number of swaps. The older approaches of the Shell Sort algorithm have stated that the number of swaps produced by Shell Sort can be further reduced by choosing an efficient gap sequence.

The main purpose of reducing the number of swaps is to use the computational resources that are available in terms of processor speed, memory and storage.

Enhanced Gap Sequencing shell sort algorithm provides an efficient and better approach to decrease the number of comparisons as well as number of swaps. Enhanced Gap Sequencing shell sort algorithm results least number of swaps on any size of data. This algorithm works more efficiently as the size of data grows.

This algorithm has described a simple and easy formula that calculates the values of N_1 by the formula $N_1 = \text{floor}(3n/4)$, where ‘n’ is the number of elements to be sorted, and then find values of $N_2; N_3; \dots; N_k$ by placing values of $N_1; N_2; N_3; \dots; N_{k-1}$ in place of ‘n’ in the formula. This algorithm improves the performance of the existing algorithms up to 60% in some cases.

VI. ACKNOWLEDGMENT

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