

Algorithm of Adaptive Genetic Hybrid proposed Approach

Input: Test Suite T
Output: T_{greatest} (A test case which has largest fitness value in population of final generation).
 APSC (Measure Adaptive Percentage of Statement Coverage)

Declaration:

Ts: represent the latest selected test case .
 N : number of test cases
 M : statements
 P: population size .
 G: number of generation.
 Cp: Crossover Point.
 Mp : Mutation Point.
 Ltc : Left Test cases after Adaptive Approach ordering .

Adaptive Process :

1. **Begin**
2. for each test case t in T.
3. calculate initial priority(t).
4. **End for** .
5. Select the test case (ts) with the largest priority in T.
6. Add ts to T'
7. $T \leftarrow T - \{ts\}$.
8. Run ts.
9. **While** T is not empty **do**.
10. For each test case t in T.
11. Change priority(t) based on the output of ts.
12. **End for**
13. Select the test case(ts) with largest priority which cover statement.
14. Add ts to T'.
15. $T \leftarrow T - \{ts\}$.
16. Run ts.
17. **End while**
18. **Return** Ltc : left test case from T those not cover statement.

Genetic Algorithm Process :

19. **Begin :**
20. **Input:** Ltc
21. $P_1 \leftarrow$ generate population (Ltc,P,fl,fsl).
22. For i=1 to g.
23. $F_1 \leftarrow$ evaluateFitness (P_i, t_c, fl, fsl)
24. $P_{i+1} \leftarrow$ addTwoBest(F_i, P_i)
25. For j=3 to P.
26. $Parent_1 \leftarrow$ RandomSelectParent(P_i)
27. $Parent_2 \leftarrow$ RandomSelectParent(P_i)
28. $Child_1, child_2 \leftarrow$ CrossOver($Parent_1, Parent_2, C_p$)
29. $Child_1 \leftarrow$ Mutation($Child_1, mp$)
30. $Child_2 \leftarrow$ Mutation($Child_2, mp$)
31. $P_{i+1} \leftarrow$ addChildren($Child_1, child_2$)
32. $F_{g+1} \leftarrow$ EvaluateFitness(P_{g+1}, t_c, fl, fsl)
33. $T_{\text{greatest}} \leftarrow$ SelectBest Child(F_{g+1}, P_{g+1})
34. **Return** T_{greatest} .
35. **Measure APSC :**
36. $C \leftarrow n * m$ ($n \leftarrow Lts$)
37. $N_2 \leftarrow 2 * n$
38. $S_1 \leftarrow$ sum/c (sum=0)
39. $S_2 \leftarrow 1 / (2 * n)$
40. $Ap_{sc} \leftarrow 1 - (S_1 + S_2)$
41. $Ap_{sc} \leftarrow Ap_{sc} * 100$
42. **Return** Ap_{sc}

Figure 3: Algorithm1 (Adaptive Genetic Hybrid Algorithm)

In figure 4 represent the flow chart of our proposed algorithm the gray shaded area in flow chart represent the adaptive approach and rest part applied by us that is genetic algorithm .

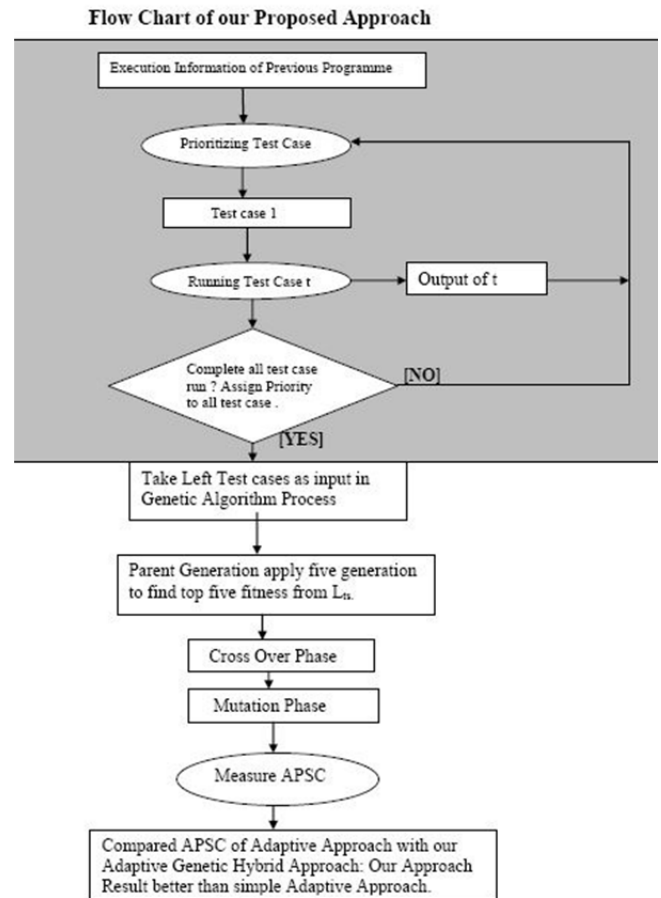


Figure 4 : Flow Chart of our Proposed Approach(Research Methodology)

Adaptive Genetic Algorithm Hybrid proposed test-case prioritization approach in this approach we ordering the test case and find the average percentage of statement coverage for hundred test cases in java . First we measure the APSC of adaptive approach and ordering the test case. In adaptive approach we order the test case like until our statement not cover if test cases left or we can say failure test cases those are unable to cover any statement its means the statement coverage is not done perfectly . We take that Left test cases after applying adaptive approach and perform genetic algorithm on these test case. In Genetic algorithm we apply three main techniques to order the test case like this our APSC improved as compared to adaptive approach. We apply these techniques in genetic algorithm to giving the order to each test case

- 3.1 Adaptive Approach
- 3.2 Parent Generation
- 3.3 Cross Over
- 3.4 Mutation
- 3.5 Measure APSC
- 3.6. Execution time

3.1 Adaptive Approach

In this Research Methodology, we first present the adaptive process of the existing test-case prioritization approach by howing its basic difference with our proposed approach adaptive genetic hybrid approach and then give the details of the adaptive genetic hybrid approach in below sections. For ease of exhibition, we present the adaptive genetic hybrid test case ordering approach in terms of statement coverage, which can also be implemented on other adaptive approach also. In figure 8: the dark area of flow chart represent the adaptive approach methodology the rest for flow chart is further methodology of Genetic algorithm. The overall flowchart figure 3. Represent the our adaptive genetic hybrid approach methodology.

We take hundred apache server test cases Antloader package of test cases in java IDE Eclipse.

First we set each test case priority 1.

$$Priority(t) = \sum Potential(S) \text{ ----- (1)}$$

Where potential(s) represent how likely statement covered by the existing selected test case. Potential(S) of any statement S in which scope [0,1].

$$Potential(S) = \begin{cases} \text{If test case}(t') \text{ passed then,} \\ Potential(s), s \text{ is not executed by } t'. \\ Potential(s)*q, s \text{ is executed by } t' \\ \text{If test case}(t') \text{ failed then,} \\ Potential(s)*p, s \text{ is executed by } t' \end{cases}$$

P and q are two non-negative constants whose values are between 0 and 1. In our implementation process while all test case priority set 1 in initial than, we run all test case those test case cover statement we provide “G” to that test case. Those test case gain maximum number of G their priority must be high. so according to this process we found that in this approach by running hundred test cases few test cases cover the statements on that bases we calculate APSC of this approach.

The effect of passed/failed output on the *Potent(s)* of any statement *s* is measured by *q/p* in the earlier equation. Moreover, when *p=q=0*, the adaptive approach becomes the additional statement-coverage based test-case prioritization approach, whereas when *p=q=1*, the adaptive approach becomes the total statement-coverage based test case ordering approach. That is, the total or additional statement-coverage based test-case ordering approach can be viewed as an instance of the adaptive approach. The existing research on test-case prioritization has fully evaluated the effectiveness of the total approach and the additional approach. Although *p* and *q* in the preceding equation are two independent variables, to facilitate evaluation of the proposed adaptive approach, currently we assume *p+q = 1* in this research and evaluate the effectiveness of the adaptive approach by setting *q=0, 0.2, 0.4, 0.6, 0.8, or 1*,

Than we calculate APSC for adaptive approach by applying APSC formula .

$$APSC = 1 - \frac{T_{s_1} + T_{s_2} + \dots + T_{s_m}}{n * m} + 1/2 * n$$

3.2 Parent Generation : This is the first step of genetic algorithm of parent selection we apply this process only on the remaining test cases after adaptive approach for the selection of five top parents we set priority to each test case according to the statement coverage we calculate fitness In our proposed algorithm 1 Pg + 1, is produced, the fitness value of each chromosome is determined on line 33. In Algorithm1 , and the chromosome whose fitness value is the greatest is selected to be the test order. We select parent randomly Algorithm 2 show that how parent selection process going on.

Algorithm 2: Random Parent Selection Algorithm

- Input : Pi the population of the ith generation.
 output : Parent chromosome selection
1. FitnessSum← calculate fitnessSum of chromosome(Pi)
 2. r← generate random number(FitnessSum)
 3. for K=1 to P
 4. r← r-fitof Chromosomek
 5. if r < 0
 6. Break
 7. Parent← chromosomek
 8. Return Parent

Figure 5: Random Parent Selection Algorithm.

As the above Figure 5 Algorithm 2 states that first we input the population of the rest of test cases after adaptive approach in1st generation we apply five generation in our experiment. According to above algorithm first we take randomly chromosomes. We take two highest priority test cases from previous adaptive approach as Parent1 and Parent 2. While we calculate the fitness and the highest fitness test case become the next parents of nest generation. Like this process we got five highest parents with high fitness value. After performing first generation we not consider that highest parent fitness in second generation. Same like this after getting second highest fitness value we don't consider that test case in third so on until we not complete all process for each test case.

$$Fitness = 1 - \frac{T_{s_1} + T_{s_2} + \dots + T_{s_m}}{n * m} + 1/2 * n$$

3.3 Cross Over : After completion of first step of Genetic algorithm we get two Parents of high fitness value now we will perform cross over operation in our proposed approach. Crossover is ordinarily a recombination transform that consolidates the portions of one chromosome with the sections of another. The new chromosomes framed by hybrid acquire a few qualities from both folks. The calculation of the hybrid administrator is given in Fig. 3.5. The calculation is the single point hybrid. In the first place, an arbitrary number, *r*, which extends from 0 to 100, is created on line 1. On the off chance that *r* is not exactly the hybrid likelihood, *cp*, the recombination procedure will start on line 3. Something else, the kid is the copy of the guardian. At the point when hybrid is connected, the calculation chooses hybrid focuses, *p1* and *p2*, for parent1 and parent2, separately, on lines 3 and 4. On lines 5 and 6, the subsequences before the hybrid point are then duplicated

from both folks. The joined capacity on lines 7 and 8 creates a tyke by consolidating the duplicated subsequence of one guardian with the qualities of another guardian that are not in the replicated subsequence.

Input: Parent₁ : selective Chromosome from population
 Parent₂ : Another Selective chromosome from population
 C_p : CrossOver Point

Output: Children₁, Children₂ (two new chromosomes produced by algorithm)

1. n ← generateRandomNumber(100)
2. if n < C_p
3. P₁ ← select Crosspoint(Parent₁)
4. P₂ ← select Crosspoint(Parent₂)
5. Segment₁ ← fragment(P₁, Parent₁)
6. Segment₂ ← fragment(P₂, Parent₂)
7. children₁ ← join(Segment₁, Parent₂)
8. children₂ ← join(Segment₂, Parent₁)
9. else
10. children₁ ← Parent₁
11. children₂ ← Parent₂
12. **Return** children₁ children₂

Figure 6: Algorithm 3 Cross over

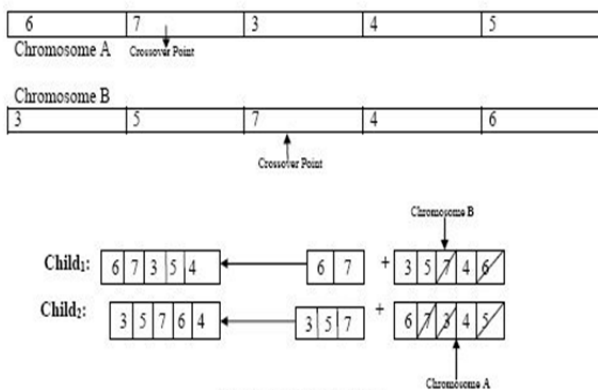


Figure 7: Example of Crossover

For example, considering the chromosomes in Fig 7 demonstrates their hybrid process. The hybrid purposes of An and B are at positions 2 and 3, individually. Child1 gets the subsequence before the hybrid point from An, and the rest from B. Since 6 and 7, which are qualities of B, are additionally in the subsequence duplicated from A, they are not added to the child1. So also, child2 acquires the subsequence before the hybrid point from B, and the qualities that are not in that subsequence from A.

3.3.4 Mutation: Mutation is performed on the chromosomes got by the hybrid process First, the transformation (children_c, mp) creates a number, n, which goes from 0 to 100 on line 1. On the off chance that n is not exactly the change likelihood, mp, the calculation chooses two qualities of children_c arbitrarily and swap their positions, as demonstrated in Fig.8. Something else, the transformation administrator would not be connected.

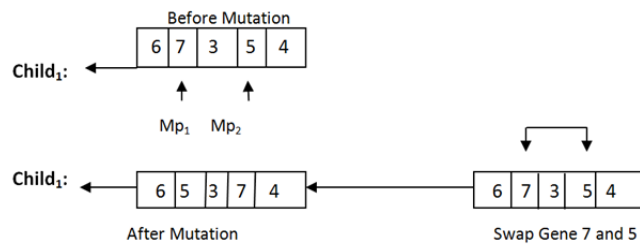


Figure 8 : Example of Mutation

Algorithm 4: Mutation

Input: Children_c chromosome produced by crossover .
 M_p Mutation Point

Output: Children_m, chromosome produced by algorithm

1. N ← GenerateRandomNumber(100)
2. if N < M_p
3. Mp₁, Mp₂ ← select Mutation Points
4. Children_m ← SwapPosition(Mp₁, Mp₂, Children_c)
5. else
6. Children_m ← Children_c
7. **Return** Children_m

Figure 9: Algorithm 4 Mutation Algorithm

In Mutation phase of genetic algorithm paper we take both children chromosome generated by the crossover operator. We show in figure 3.8, how the children change after applying mutation operator. First we take children₁ and randomly generate number for two different mutation points . as in example Mp₁ and Mp₂ indicate gene 7 and 5 in above example. We simply swap these genes and got children_{m1} ,and Children_{m2} . same process going on for each chromosomes we received after cross over operator/phase . The mutation (child_c, mp) also gives those test cases a chance to get a higher priority for test case ordering .

3.5 Measure APSC: The fifth step of our methodology is measuring the average percentage of statement coverage which will show our experimental work, the result of APSC represent how our approach is better than adaptive approach. The general formula to measure APSC.

$$APSC = 1 - \frac{T_{s1} + T_{s2} + \dots + T_{sm} + 1/2 * n}{n * m}$$

but in our experimental coding we apply this formula like .

$$APSC = 1 - \frac{\text{sum}/c + 1/2 * n}{n * m}$$

Where , n= number of test case (L_{tc} left test cases after adaptive approach)

- M= statements
- C= n*m
- S₁ = sum/n*m , sum/C .
- S₂ = 1/(2*n)

$$APSC = 1 - S_1 + S_2 .$$

We take all high order test cases to evaluate the apsc for our proposed approach, we tak summation of each test-case priority calculate by our algorithm 1. On the basis of that proritry reading we measure APSC and our results shows that our proposed approach is better than the previous adaptive approach. We are able to increase the efficiency of average percentage of statement coverage. Our results show in graphical form in the chapter Result and Analysis.

3.6 Execution time

The last parameter measures in this research is execution time .we calculate how much execution time should be taken by existing approach and proposed approach .and we found that execution time is high in our proposed approach because this is generally clear as well it should take more time than adaptive approach because we execute adaptive approach as well genetic algorithm process in which five parent generation , crossover and mutation operators processing in our proposed approach .we also found that execution time depend on the system configuration also . while we process this approach on high configuration system it take less time while we process on low configuration system it take a lot of time . so we conclude this parameter in our future scope we can improve APSC as well time execution if we apply any other approach/technique further.

4. RESULT AND GRAPHS

In this chapter we will discuss about the result obtained by us for both existing approach as well our proposed approach. In existing approach of adaptive test-case prioritization we calculate APSC (average percentage of statement coverage) and execution time also by vary the q and p value .

The existing research on test-case prioritization has fully evaluated the effectiveness of the total approach and the additional approach. Although p and q in the preceding equation are two independent variables, to facilitate evaluation of the proposed adaptive approach, currently we assume $p+q = 1$ in this research we evaluate the effectiveness of the adaptive approach and proposed approach by setting $q=0, 0.2, 0.4, 0.6, 0.8, \text{ and } 1$. We focus on Q factor value just because the q factor value multiply only when test case is pass. Same like that we calculate the execution time for the adaptive approach and proposed approach by setting $q=0, 0.2, 0.4, 0.6, 0.8, \text{ and } 1$.

Table 4.1 Adaptive Approach by different p, q factor value.

Q	p	APSC	Execution time
0	1	97.6097052	1343 ms
0.2	0.8	98.6940925	1047 ms
0.4	0.6	98.6645584	859 ms
0.6	0.4	98.6645584	969 ms
0.8	0.2	98.6645584	1214 ms
1	0	98.61392425	1191 ms

As it shown in table number 4.1 while we take different p, q factor values we get different average percentage of statement coverage and execution time. We found that the highest APSC is at $q=0.2, p=0.8$ value while the minimum time taken at $q=0.4, p=0.6$. so from this table we analyses we can change the factor value according to our need in which we have need to focus. If our focus on to statement coverage we take best value of p, q in which we get highest APSC. While we have needed to focus on execution time we will select p, q value according to the low executions time value.

Figure 10, represent the value of APSC on Y-axis while there is different Q factor value on x-axis. While in figure 11 the graph resent the execution time take by adaptive approach while we measure APSC.

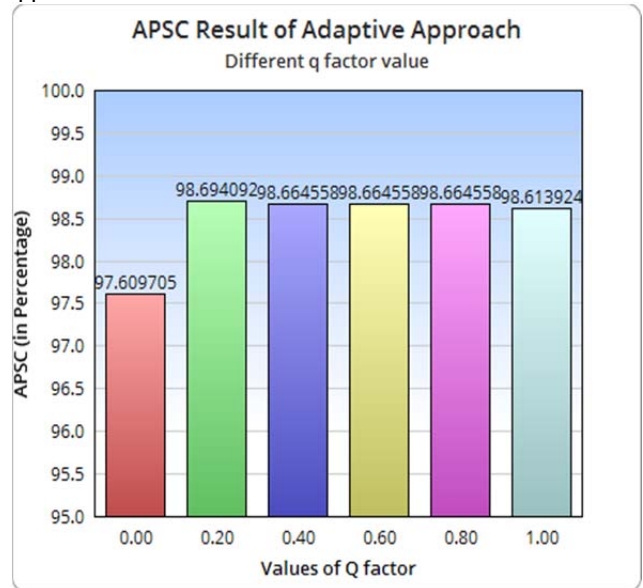


Figure 10: Graph of APSC according to Different Q values in Adaptive Approach.

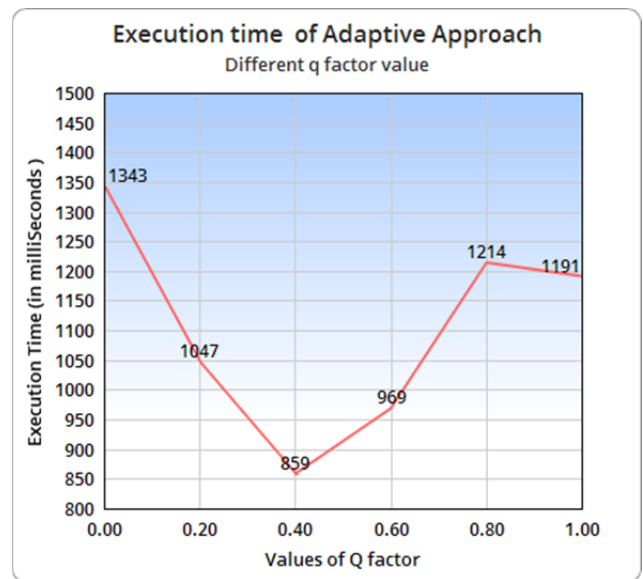


Figure 11: Graph of Execution Time according to Different Q values in Adaptive Approach

After measuring the APSC and execution time for adaptive approach. Than we measure the APSC and execution time for our proposed approach adaptive genetic hybrid approach. In Table number 4.2 the values of APSC and Execution time date at different Q and P factors. On the basis of this data set figure 12 represent the graph of APSC values on y axis and different Q factor values on x-axis. Similarly figure 13 represent the execution time value on y-axis and different q factor value on x-axis.

Table 4.2 Adaptive Genetic hybrid Approach by different p, q factor value.

Q	p	APSC	Execution time
0	1	99.34755322	41691 ms
0.2	0.8	99.54693411	52706 ms
0.4	0.6	99.67835639	50404 ms
0.6	0.4	99.65760801	48956 ms
0.8	0.2	99.67144225	39178 ms
1	0	99.58897335	51171 ms

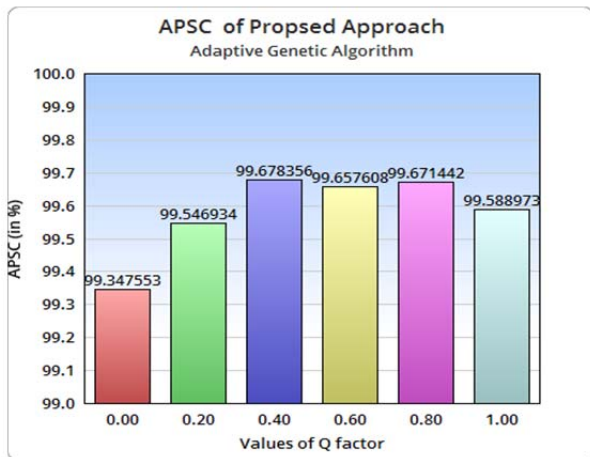


Figure 12: Graph of APSC according to Different Q values in proposed Approach

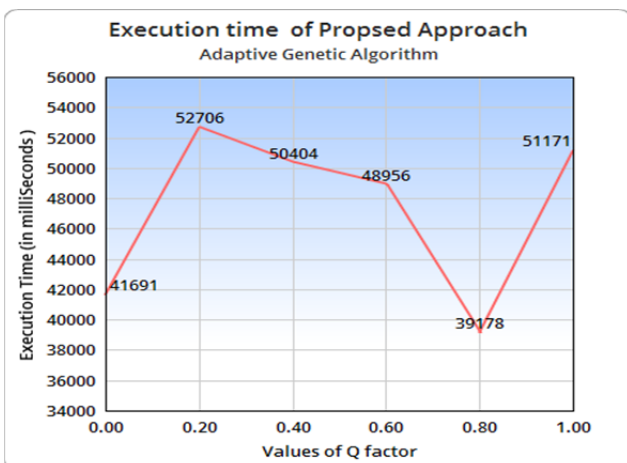


Figure 13: Graph of Execution Time according to Different Q values in Proposed Approach.

In the above figure and tables we represent the separate outcomes from both approach the existing adaptive approach as well our proposed approach. After that we compare the both approach on the basis of APSC as well time execution . In Figure 14 it shows that the comparison between the adaptive and our proposed approach. The red color bar indicate the adaptive approach while green bar represent our proposed approach and it is clear represented by graph our approach is better than existing adaptive technique in this research basically we focus only on APSC rather than execution time. While in figure 15 the line graph represent the time taken comparison between the existing

approach as well our proposed approach red line in graph represent the time taken by adaptive approach and green line represent the time taken by our proposed approach.

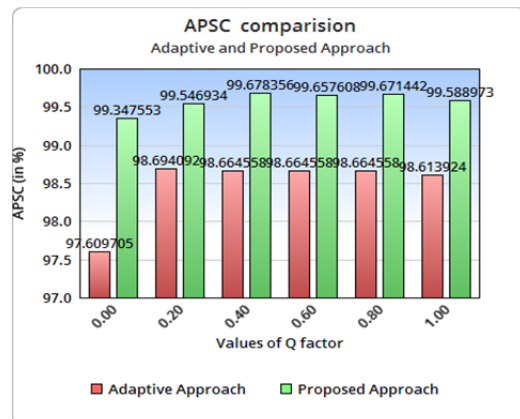


Figure 14: APSC Comparison of Adaptive and proposed Approach.

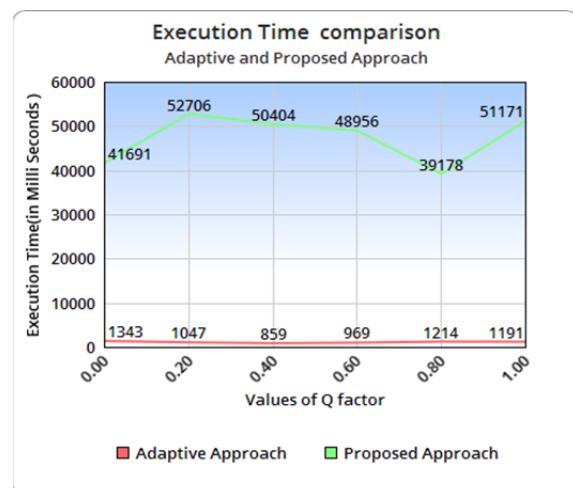


Figure 15 : Comparison of Execution Time among Adaptive and Proposed Approach.

5. CONCLUSION AND FUTURE SCOPE

In this Research we proposed an approach that improves APSC (average percentage of statement coverage). Our work is extension into the adaptive approach for APFD (average percentage of fault detection) into adaptive genetic algorithm hybrid approach from which we conclude that our proposed approach improve the APSC. We take hundred java test cases package of apache server to evaluate our approach. First we apply adaptive approach and calculate APSC. Then we apply our proposed algorithm adaptive genetic algorithm hybrid approach than we calculate APSC than we found that our approach gives better results than adaptive approach for APSC only. Basically in this research we focused on APSC only but while we calculate Execution time for both approach we found that our proposed approach take large time to execute as compare to adaptive approach. But as the tester view our main aim to cover all statements of the code for better quality. So, we considering this work as our next future work and we believe that if we apply any other technique we can improve execution time as well APSC together. And we take small data set in our research while in future we take large data set of test cases for efficient results.

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