Journal of Computer Science 4 (5): 393-401, 2008 ISSN 1549-3636 © 2008 Science Publications

A Fast Pattern Matching Algorithm with Two Sliding Windows (TSW)

Amjad Hudaib, Rola Al-Khalid, Dima Suleiman, Mariam Itriq and Aseel Al-Anani Department of Computer Information Systems, University of Jordan, Amman 11942 Jordan

Abstract: In this research, we propose a fast pattern matching algorithm: The Two Sliding Windows (TSW) algorithm. The algorithm makes use of two sliding windows, each window has a size that is equal to the pattern length. Both windows slide in parallel over the text until the first occurrence of the pattern is found or until both windows reach the middle of the text. The experimental results show that TSW algorithm is superior to other algorithms especially when the pattern occurs at the end of the text.

Key words: Pattern matching, string matching, berry-ravindran algorithm, boyer moore

INTRODUCTION

Pattern matching is a pivotal theme in computer research because of its relevance to various applications such as web search engines, computational biology, virus scan software, network security and text $processing^{[1-4]}$.

Pattern matching focuses on finding the occurrences of a particular pattern P of length 'm' in a text 'T' of length 'n'. Both the pattern and the text are built over a finite alphabet set called \sum of size σ .

Generally, pattern matching algorithms make use of a single window whose size is equal to the pattern length^[5]. The searching process starts by aligning the pattern to the left end of the text and then the corresponding characters from the pattern and the text are compared. Character comparisons continue until a whole match is found or a mismatch occurs, in either case the window is shifted to the right in a certain distance^[6-12]. The shift value, the direction of the sliding window and the order in which comparisons are made varies in different pattern matching algorithms.

Some pattern matching algorithms concentrate on the pattern itself^[5]. Other algorithms compare the corresponding characters of the pattern and the text from left to right^[6]. Others perform character comparisons from right to left^[8,11]. The performance of the algorithms can be enhanced when comparisons are done in a specific order^[9,13]. In some algorithms the order of comparisons is irrelevant such as Brute Force and Horspool algorithms^[7].

In this study, we propose a new pattern matching algorithm: The Two Sliding Windows algorithm (TSW). The algorithm concentrates on both the pattern and the text. It makes use of two windows of size that is equal to the size of the pattern. The first window is aligned with the left end of the text while, the second window is aligned with the right end of the text. Both windows slide at the same time (in parallel) over the text in the searching phase to locate the pattern. The windows slide towards each other until the first occurrence of the pattern from either side in the text is found or they reach the middle of the text. If required, all the occurrences of the pattern in the text can be found.

Related works: Several pattern matching algorithms have been developed with a view to enhance the searching processes by minimizing the number of comparisons performed^[14-16]. To reduce the number of comparisons, the matching process is usually divided into two phases. The pre-processing phase and the searching phase. The pre-processing phase determines the distance (shift value) that the pattern window will move. The searching phase uses this shift value while searching for the pattern in the text with as minimum character comparisons as possible.

In Brute Force algorithm (BF), no pre-processing phase is performed. It compares the pattern with the text from left to right. After each attempt, it shifts the pattern by exactly one position to the right. The time complexity of the searching phase is O (mn) in the worst case and the expected number of text character comparisons is (2n).

New ways to reduce the number of comparisons performed by moving the pattern more than one position are proposed by many algorithms such as Boyer-Moore $(BM)^{[11,17]}$ and Knuth-Morris-Pratt algorithms (KMP)^[6,18].

Corresponding Author: Amjad Hudaib, Department of Computer Information Systems, University of Jordan, Amman 11942, Jordan Tel.: +962-5355000/ext: 22610 Fax: +962-5354070

KMP algorithm compares the pattern with the text from left to right. If a mismatch occurs it uses the failure function f(j)that indicates the proper shift of the pattern^[6,18]. The failure function f (j) is defined as the length of the longest prefix of P that is the suffix of P[1..j]. Thus, KMP reduces the number of times it compares each character in P with a character in the text T. KMP performs (2n) text character comparisons and the complexity of the pre-processing phase is O (m). KMP achieves a running time of O (n+m), which is optimal in the worst case^[6].

BM algorithm improves the performance by preprocessing the pattern using two shift functions: the bad-character shift and the good-suffix shift. During the searching phase, the pattern is aligned with the text and it is scanned from right to left. If a mismatch occurs, the BM algorithm shifts the pattern with the maximum value taken between the two shift functions. The worst case time complexity when searching all occurrences of the pattern is O (mn) and O (nm⁻¹) for best performance^[11,17].

A simplification of BM algorithm is the Horspool algorithm^[7]. It does not use the good suffix function, instead it uses the bad-character shift with the rightmost character. Its pre-processing time complexity is $O(m+\sigma)$ and the searching time complexity is $O(mn)^{[7]}$.

The Berry-Ravindran algorithm (BR) calculates the shift value based on the bad character shift for two consecutive text characters in the text immediately to the right of the window. This will reduce the number of comparisons in the searching phase. The pre-processing and searching time complexities of BR algorithm are $O(\sigma^2)$ and O(nm) respectively^[7]. In this research, the proposed algorithm makes use of the pre-processing phase of BR algorithm.

The Two Sliding Windows (TSW) algorithm: The Two Sliding Windows algorithm (TSW) scans the text from both sides simultaneously. It uses two sliding windows, the size of each window is m which is the same size as the pattern. The two windows search the text in parallel. The text is divided into two parts: the left and the right parts, each part is of size r n/2r. The left part is scanned from left to right using the left window and the right part is scanned from right to left using the right window. Both windows slide in parallel which makes the TSW algorithm suitable for parallel processors structures. TSW algorithm stops when one of the two sliding windows finds the pattern or the pattern is not found within the text string at all. The TSW algorithm finds either the first occurrence of the pattern in the text through the left window or the last occurrence of the pattern through the right window. If necessary, the algorithm can be modified easily to find all the occurrences of the pattern. Also if the pattern is exactly in the middle of the text, TSW can find it easily.

The TSW algorithm utilizes the idea of BR bad character shift function^[8] to get better shift values during the searching phase. BR algorithm provides a maximum shift value in most cases without losing any characters. The main differences between TSW algorithm and BR algorithm are:

- TSW uses two sliding windows rather than using one sliding window to scan all text characters as in BR algorithm
- The TSW uses two arrays, each array is a one dimensional array of size (m-1). The arrays are used to store the calculated shift values for the two sliding windows. The shift values are calculated only for the pattern characters. While the original BR algorithm uses a two-dimensional array to store the shift values for all the alphabets^[8]. Using one dimensional array reduces the search processing time and at the same time reduces the memory requirements needed to store the shift values

Pre-processing phase: The pre-processing phase is used to generate two arrays *nextl* and *nextr*, each array is a one-dimensional array. The values of the nextl array are calculated according to Berry-Ravindran bad character algorithm (BR). nextl contains the shift values needed to search the text from the left side. To calculate the shift values, the algorithm considers two consecutive text characters a and b which are aligned immediately after the sliding window. Initially, the indexes of the two consecutive characters in the text string from the left are (m+1) and (m+2) for a and b respectively as in Eq. 1.

Bad Char shiftl [a,b] = min $\begin{cases}
1 & \text{if } p[m-1] = a \\
m-i & \text{if } p[i]p[i+1] = ab \\
m+1 & \text{if } p[0] = b \\
m+2 & \text{Otherwise}
\end{cases}$ (1)

On the other hand, the values of the nextr array are calculated according to our proposed shift function. nextr contains the shift values needed to search the text from the right side, initially the indexes of the two consecutive characters in the text string from the right

```
\begin{array}{l} \text{Begin} \\ \text{shiftl=shiftr=m+2} \\ \text{for (each character } p_i \in P_{i=0 \dots m\cdot 2}) \\ & \quad \left( \text{Next}[i] = m\text{-}i, nextr[i] = m\text{-}((m-2)\text{-}i) \right) \\ \text{if } P[m-1] = a \ (shiftl=1, shiftr=m+1) \\ \text{else if } P[0] = b \ (shiftl=m+1, shiftr=1) \\ \text{else if } p[i] \ [i+1] = ab \ (shiftl=nextl[i], shiftr=nextr[i]) \\ \text{End} \end{array}
```

Fig. 1: The pre-processing algorithm

are (n-m-2) and (n-m-1) for *a* and *b* respectively, which are used to calculate the shift values as in Eq. 2.

Bad Char
shiftr [a,b] = min
$$\begin{cases} m+1 & \text{if } p[m-1] = a \\ m - ((m-2)-i) & \text{if } p[i]p[i+1] = ab \\ 1 & \text{if } p[0] = b \\ m+2 & \text{Otherwise} \end{cases}$$
(2)

The two arrays will be invariable during the searching process. Figure 1 shows the steps of the preprocessing algorithm.

Searching phase: In this phase, the text string is scanned from two directions, from left to right and from right to left. In mismatch cases, during the searching process from the left, the left window is shifted to the right, while during the searching process from the right, the right window is shifted to the left. Both windows are shifted until the pattern is found or the windows reach the middle of the text. Figure 2 explains the steps of the TSW algorithm.

Step1: Compare the characters of the two sliding windows with the corresponding text characters from both sides. If there is a mismatch during comparison from both sides, the algorithm goes to step2, otherwise the comparison process continues until a complete match is found. The algorithm stops and displays the corresponding position of the pattern on the text string. If we search for all the pattern occurrences in the text string, the algorithm continues to step2.

Step2: In this step, we use the shift values from the next arrays depending on the two text characters placed immediately after the pattern window. The two characters are placed to the right side of the left window and to the left side of the right window. The corresponding windows are shifted to the correct positions based on the shift values, the left window is shifted to the right and the right window is shifted to

```
L=m-1;//text index used from left
R=n-(m-1)-1; //text index used from right
Tindex=0;/Aext index used to control the scanning
DBOCESS
While (Tindex <= ┌ n/2┐)
Begin
  I=m-1 ; // pattern index used at left side
  r=0; // pattern index used at right side
 temp-lindex=temp-rindex=0;//keep record of the
 text index where the pattern match the text during
 comparison
  if(P[m-1]=T[L])
    begin
      temp-lindex=L
     L=L-1
     while (l \ge 0 \text{ and } P[l]=T[L])
        { L=L-1, l=l-1; } //search from left
   end
 if(P[0]=T[R]
  begin
    temp-rindex=R
    R=R+1
    while( (r<m and P[r]=T[R]) )
      { R=R+1, r=r+1;} //search from right
  end
   if (r>m-1) {display "match at right: "+R-m); exit
from outer bop;}
   if (1<0){ display "match at left: "+L+1); exit from
outer bop;}
   //exit in case if we search for one occurrence the
 first or last one
  R= temp-rindex; //to avoid skipping characters after
partial matching at right
  L=temp-lindex; // to avoid skipping characters after
partial matching at left
  if(L>R){ display ("not found"); exit from outer
 loop;}
  L=L+get(shift);//fiom pre-processing step
  R=R-get(shiftr) //from pre-processing step
  Tindex=Tindex+1:
End
```

Fig. 2: TSW Pattern Matching Algorithm

the left. Both steps are repeated until the first occurrence of the pattern is found from either sides or until both windows are positioned beyond $rn/2_{7}$.

If the first occurrence of the pattern exists in the middle of the text, the TSW algorithm in Fig. 2 continues comparing pattern characters with text characters through the inner loops before the TSW algorithm terminates the searching process through the outer loop.

Working example: In this study we will present an example to clarify the TSW algorithm. Part of nucleotide sequence of a gene (only 47 nucleotides) from Chromosome I (CHR-I) has been used to test the algorithm^[10], this sequence is taken from the gene index 32854-32901^[10]. The plant genome (Arabidopsis thaliana) consists of 27,242 gene sequences distributed over five chromosomes (CHR-I to CHR-V).



Fig. 3: The nextl and nextr arrays

Given:

Pattern(P)="GAATCAAT", *m*=8 *Text(T)*="ATCTAACATCATAACCCTAATTGGCA GAGAGAGAATCAATCGAATCA", *n*=47

Pre-processing phase: Initially, shiftl=shiftr=m+2=10.

The shift values are stored in two arrays nextl and nextr as shown in Fig. 3a and 3b respectively.

To build the two next arrays (nextl and nextr), we take each two consecutive characters of the pattern and give it an index starting from 0. For example for the pattern structure GAATCAAT, the consecutive characters GA,AA,AT,TC,CA,AA and AT are given the indexes 0,1,2,3,4,5 and 6 respectively.

The shift values for the nextl array are calculated according to Eq. 1 while the shift values for the nextr array are calculated according to Eq. 2.

Searching phase: The searching process for the pattern p is illustrated through the working example as shown in Fig. 4.

First attempt: In the first attempt (Fig. 4a), we align the first sliding window with the text from the left. In this case, a mismatch occurs between text character (A) and pattern character (G), therefore we take the two consecutive characters from the text at index 8 and 9 which are (T and C) respectively. To determine the amount of shift (shiftl) we have to do the following two steps:

- We find the index of TC in the pattern which is 3
- Since we search from the left side we use nextl array and shiftl = nextl[3] = 5

Therefore the window is shifted to the right 5 steps.

Second attempt: In the second attempt (Fig. 4b), we align the second sliding window with the text from the right. In this case, a mismatch occurs between text character (A) and pattern character (T), therefore we take the two consecutive characters from the text at index 37 and 38 which are (A and A) respectively. To determine the amount of shift (shiftr), we have to do the following two steps:

• We find the index of AA in the pattern, AA has two indexes 1 and 5

Since we search from the right side we use nextr array for the two indexes nextr[1] = 3, nextr[5] = 7, then we choose the minimum value to determine shiftr. Shiftr = nextr[1] = 3.

Therefore the window is shifted to the left 3 steps.

Third attempt: In the third attempt (Fig. 4c), a mismatch occurs from the left between text character (A) and pattern character (G), therefore we take the two consecutive characters from the text at index 13 and 14 which are (A and C) respectively, since AC is not found in the pattern, so the window is shifted to the right 10 steps.

Fourth attempt: In the fourth attempt (Fig. 4d), a mismatch occurs from the right between text character (A) and pattern character (T), therefore we take the two consecutive characters from the text at index 34 and 35 which are (A and T) respectively. To determine the amount of shift (shiftr) we have to do the following two steps:

- We find the index of AT in the pattern, AA has two indexes 2 and 6
- Since we search from the right side, we use *nextr* array for the two indexes

nextr[2] = 4, nextr[6] = 8, we choose the minimum. Shiftr = nextr[1] = 4.

Therefore the window is shifted to the left 4 steps.

Fifth attempt: We align the left most character of the pattern P[0] with T[32]. A comparison between the pattern and the text characters leads to a complete match at index 32. In this case, the occurrence of the pattern is found using the right window.



Fig. 4: Working Example

Analysis:

Preposition 1: The space complexity is O(2(m-1)) where m is the pattern length.

Preposition 2: The pre-process time complexity is O(2(m-1)).

Lemma 1: The worst case time complexity is O(((n/2-m+1))(m))

Proof: The worst case occurs when at each attempt, all the compared characters of both the pattern and the text are matched except the last character and at the same time the shift value is equal to 1. If the pattern is aligned from the left then shift by one occurs when the first character of the two consecutive characters is matched with the last pattern character, while if the pattern is aligned from the right then shift by one occurs when the second character of the two consecutive characters is matched with the first pattern character.

Lemma 2: The best case time complexity is O(m).

Proof: The best case occurs when the pattern is found at the first index or at the last index (n-m).

Lemma 3: The Average case time complexity is O(n/(2*(m+2))).

Proof: The Average case occurs when the two consecutive characters of the text directly following the sliding window is not found in the pattern. In this case, the shift value will be (m+2) and hence the time complexity is O([n/(2*(m+2))]).

RESULTS AND DISCUSSION

Several experiments have been conducted using TSW algorithm. In each experiment, we consider Book1 from the Calgary corpus to be the text^[19]. Book1 Table 1: The average number of attempts and comparisons of TSW and BR algorithms

		TSW		BR	
Pattern	No. of				
length	words	Attempts	Comparisons	Attempts	Comparisons
4	8103	3904	4213	6409	7039
5	4535	4456	4896	9577	10645
6	2896	7596	8311	10898	12173
7	1988	9341	10263	11953	13345
8	1167	10056	11087	13256	14807
9	681	9538	10538	14149	15892
10	382	9283	10272	14127	15799
11	191	5451	5967	12808	14243
12	69	6384	7168	9598	10923
13	55	7947	8673	10334	11370
14	139	19437	21319	19548	21673
15	32	19682	21739	19817	22384
16	10	20029	21596	26086	28644
17	3	21897	25404	22554	28148

consists of 141,274 words (752,149 characters). Patterns of different lengths are also taken from Book1.

The searching process is performed from both sides of Book1 and the pattern is located from either sides. Table 1 shows the results of comparing the algorithms TSW with BR and Fig. 5a and 5b show the average number of attempts and comparisons respectively.

In Table 1, the length of the pattern is given in column one while the second column is the number of words selected for each pattern length from Book1. For example, as shown in Table 1, 1167 patterns with length 8, were taken. The average number of

comparisons made by TSW algorithm is 11087 and average number of attempts is 10056, while the average number of comparisons made by BR is 14807 and average number of attempts is 13256. TSW made the minimum average number of comparisons and average number of attempts. Although TSW algorithm uses the same shift function of BR algorithm the TSW algorithm searches the text from both sides simultaneously, while



Fig. 5: The average number of attempts and comparisons of TSW and BR algorithms

BR algorithm only searches the text from the left side, so the average number of comparisons and attempts in BR algorithm are more than that of our algorithm.

Table 2 shows the results of comparing TSW algorithm with other algorithms. TSW algorithm has the minimum average number of comparisons and attempts among all other algorithms. The results are reasonable since TSW algorithm searches the text from both sides while all other algorithms search the text from one side. This can be justified by the following two advantages of TSW algorithm. First, it searches the text from both sides simultaneously. Second, the BR shift function shifts the pattern by a value that ranges from 1 up to m+2 positions from both sides when a mismatch occurs. This has a positive effect on the number of comparisons and attempts in most cases.

BF algorithm in Table 2, has the largest number of comparisons and attempts because it shifts the pattern by one position to the right each time a mismatch occurs. Noticeably, KMP algorithm has almost the same number of comparisons and attempts as in BF algorithm. Although, KMP uses a failure function to determine the shift values in case of a mismatch, its searching results are close to BF because multiple occurrence of a substring in a word is not common in a natural language. The performance of BM algorithm is better than KMP and BF, since it uses the good suffix function and the bad-character shift to calculate the shift which depends on the reoccurrence of the substring in a word which reduces the number of comparisons performed.

Table 3-5 show the index, number of comparisons and attempts needed to search for the first appearance of the pattern (with different lengths) in the beginning, middle and at the end of Book1. In Table 3 and 4, TSW algorithm results are reasonably

Table 2: The average number of attempts and comparisons for patterns with different lengths

		TSW		BR		BM		KMP		BF	
Pattern length	No. of words	Attempts	Comparisons								
4	8103	3904	4213	6409	7039	9549	10055	35946	36972	36029	37056
5	4535	4456	4896	9577	10645	13435	14246	61500	63460	61685	63645
6	2896	7596	8311	10898	12173	14793	15749	79064	81663	79353	81952
7	1988	9341	10263	11953	13345	15797	16817	97291	100722	97667	101100
8	1167	10056	11087	13256	14807	17190	18314	117903	122341	118360	122799
9	681	9538	10538	14149	15892	18145	19403	136829	142234	137387	142793
10	382	9283	10272	14127	15799	18048	19254	148359	154279	148997	154917
11	191	5451	5967	12808	14243	16449	17477	144335	149852	145007	150525
12	69	6384	7168	9598	10923	12074	13001	114781	120531	115338	121088
13	55	7947	8673	10334	11370	13422	14176	133469	140255	133952	140739
14	139	19437	21319	19548	21673	25075	26603	265189	275981	266460	277257
15	32	19682	21739	19817	22384	24791	26609	277260	288103	278900	289750
16	10	20029	21596	26086	28644	33423	35146	391604	403333	393580	405313
17	3	21897	25404	22554	28148	26266	30016	334855	347547	336367	349060

J. Computer Sci., 4 (5): 393-401, 2008

		TSW		BR		BM		KMP		BF	
Pattern											
length	Index	Attempts	Comparisons								
4	67	25	29	13	17	18	22	68	72	68	72
5	33	11	15	6	10	8	12	34	38	34	38
6	82	23	28	12	17	16	22	83	89	83	89
7	39	11	17	6	12	7	14	40	46	40	46
8	99	21	28	11	18	16	24	100	109	100	109
9	260	51	65	31	44	35	43	259	278	261	280
10	590	105	120	55	69	74	93	589	607	591	607
11	189	35	47	16	26	19	29	190	202	190	202
12	2401	363	402	198	226	254	269	2398	2537	2402	2541

Table 3: The number of attempts and comparisons performed to search for the first appearance of a selected pattern from the beginning of the text

Table 4: The number of attempts and comparisons performed to search for the first appearance of a selected pattern from the middle of the text

	TSW		BR		BM		KMP		BF		
Pattern											
length	Index	Attempts	Comparisons								
4	370383	129947	137725	65258	72993	92820	98573	370381	370529	370384	370532
5	370382	113693	136161	57118	63667	76853	81719	370310	430611	370383	430684
6	370381	98091	106374	49270	54977	65873	69675	368479	376328	370381	378230
7	370380	87631	99564	44280	49423	57885	61504	365718	390065	370380	394727
8	370379	79241	92310	40408	45146	52106	55303	360918	388684	370379	398145
9	370378	72129	80309	37118	41536	47672	50528	368576	386294	370378	388096
10	370377	66303	74941	34490	38503	44166	46915	368073	393393	370377	395697
11	370376	61585	71378	32152	35861	41471	44021	367999	405422	370376	407799
12	370375	54745	63718	29387	29440	41713	41758	365826	430607	370374	435156

Table 5: The number of attempts a			

		TSW		BR		BM		KMP		BF	
Pattern											
length	Index	Attempts	Comparisons								
4	768606	66	76	136118	151948	195168	207233	768607	816438	768607	816440
5	768468	94	102	112181	114792	158135	160057	768469	816294	768469	816296
6	768189	164	183	99387	102611	132797	134899	768190	805725	768190	805729
7	768510	68	79	93558	104611	116750	124074	768322	791589	768511	791778
8	768486	66	78	83653	93537	101472	107957	768485	795109	768487	795575
9	768595	44	55	77868	87499	93751	99992	768593	780584	768596	780587
10	768222	104	117	70970	79150	84280	89630	768220	780205	768223	780208
11	768240	92	107	65279	72965	77326	82378	768238	780224	768241	780227
12	768306	78	96	60905	67108	70803	74580	768304	780292	768307	780295

Table 6: The average number of attempts and comparisons performed to search for (100) patterns selected from the beginning of the text

		TSW		BR		BM H		KMP		BF	
Pattern	No. of										
length	words	Attempts	Comparisons								
4	100	143	157	76	85	113	121	422	437	423	439
5	100	185	206	99	115	138	151	633	658	635	660
6	100	227	255	121	142	163	179	866	903	869	906
7	100	347	388	195	226	255	281	1550	1621	1555	1627
8	100	504	568	270	310	350	380	2366	2484	2378	2496
9	100	670	750	363	417	469	509	3493	3652	3511	3671
10	100	1160	1290	640	727	812	878	6645	6951	6689	6996
11	100	622	705	331	396	426	474	3698	3822	3710	3834
12	100	865	972	478	557	580	632	5542	5823	5567	5848

acceptable compared with BR and BM algorithms since they search the text only form left to right, while TSW algorithm searches the text from both sides, which normally increases the number of comparisons and attempts.

On the other hand, BF and KMP have the largest number of comparisons. TSW algorithm performance is

observed in Table 5 where a pattern with different lengths is selected from the end of Book1. TSW algorithm finds it with minimum effort by the right to left window.

Table 6-8 show the average number of comparisons and attempts needed to search for the first, middle and last appearance of 100 words selected from

J.	Computer	Sci.,	4	(5):	393-4	401,	2008

		TSW	ГSW		BR		BM			BF	
Pattern	No. of										
length	words	Attempts	Comparisons								
4	100	2726	2959	3645	4070	5451	5794	20263	20982	20324	21043
5	100	13965	15140	11558	12793	16057	16984	74308	76485	74524	76701
6	100	16682	18317	12878	14337	17545	18639	93846	97081	94112	97347
7	100	27267	30095	19547	22006	25732	27537	156518	162057	157247	162786
8	100	27830	30915	20831	23336	27043	28845	184588	191677	185458	192551
9	100	33929	37200	23284	25852	30154	32043	226004	234510	227028	235537
10	100	29676	32817	20546	22989	26300	28049	214942	224036	216065	225160
11	100	23195	24646	20264	22005	26377	27620	231558	237139	232474	238061
12	100	26806	30222	21113	24235	26818	29098	255284	269960	256314	270993

Table 7: The average number of attempts and comparisons performed to search for (100) patterns selected from the middle of the text

Table 8: The average number of attempts and comparisons performed to search for (100) patterns selected from the end of the text

		TSW	TSW		BR		BM			BF	
Pattern	No. of										
length	words	Attempts	s Comparisons	Attempts	Comparisons	Attempts	Comparisons	Attempts	Comparisons	Attempts	Comparisons
4	100	133	148	6899	7719	10197	10857	38402	39254	38459	39310
5	100	271	297	12930	14404	18352	19484	83041	85608	83229	85795
6	100	364	402	21315	23957	28737	30713	154185	158658	154638	159112
7	100	402	447	22237	24731	29554	31372	180131	185979	180783	186631
8	100	536	592	21495	23841	28051	29727	191651	197606	192232	198187
9	100	776	859	24919	28257	31525	33891	240397	248454	241177	249235
10	100	1579	1756	31603	35360	40451	43195	333436	345747	334643	346956
11	100	619	669	32797	36438	41812	44450	367043	377922	368994	379873
12	100	1667	1872	30928	35067	38553	41374	367780	384834	369501	386555

Table 9: The number of attempts and comparisons performed to search for a set of patterns that do not exist in the text

	TSW		BR		BM		KMP		BF	
Pattern										
length	Attempts	Comparisons								
4	132894	142538	135240	151799	193892	206747	768769	777901	768769	777941
5	113904	121958	115988	129713	156915	166626	768768	777900	768768	777940
6	96782	98008	98075	98201	135651	135728	768767	777899	768767	777939
7	85970	86986	87081	87141	115235	115270	768766	777898	768766	777938
8	78722	82124	81500	86587	100541	103498	768765	777897	768765	777937
9	72136	76506	75625	82879	91793	96158	768764	777896	768764	777936
10	65890	69354	68904	74522	85619	89079	768763	777895	768763	777935
11	60094	61964	62235	64683	80370	82038	768762	777894	768762	777934
12	56774	60319	60911	67164	74262	78281	768761	777893	768761	777933

Book1. The results of taking 100 words are similar to that of taking a single word with different lengths. As shown in Table 8, TSW algorithm best performance is when we search for words selected from the end of Book1. In case of a complete mismatch, as in Table 9, the average number of comparisons and attempts of the TSW algorithm is the minimum; this is because the shift value in most cases reaches m+2.

CONCLUSION

In this research, we presented a fast pattern matching algorithm The Two Sliding Windows algorithm TSW which makes use of two sliding windows. It employs the main idea of BR by maximizing the shift value and using two sliding windows rather than using one sliding window to scan all text characters as in BR algorithm. The TSW uses two arrays; each array is a one dimensional array of size (m-1). The arrays are used to store the calculated shift values for the two sliding windows, while the original BR algorithm uses a two-dimensional array.

We evaluated TSW performance by using a text string and various set of patterns. Also in the algorithm, during the pre-processing phase we reduced the memory required by using one-dimensional arrays for the pattern characters only. The concept of searching the text from both sides simultaneously gives TSW algorithm a preference over other algorithms in the number of comparisons and attempts especially if the pattern searched for occurs at the end of the text. In future research, we intend to implement the TSW algorithm on real parallel processors to minimize the number of comparisons and attempts. Also we intend to implement the idea of the two sliding windows on other algorithms such as KMP and BM.

REFERENCES

- Wang, Y. and H. Kobayashi, 2006. High performance pattern matching algorithm for network security. IJCSNS, 6: 83- 87. URL:http://paper.ijcsns.org/07_book/200610/ 200610A3.pdf
- Navarro, G. and M. Raffinot, 2002. Flexible Pattern Matching in Strings-Practical On-line Search Algorithms for Texts and Biological Sequences. First Edition. Cambridge University Press, New York. ISBN: 0-521-81 307-7
- Crochemore, M. and W. Rytter, 2002. Jewels of Stringology. First Edition. WorldScientific, Singapore. ISBN: 9789810247829
- Smyth, W.F., 2003. Computing Patterns in Strings. First Edition. Pearson Addison Wesley. United States. ISBN: 978-0-201-39839-7
- Charras, C. and T. Lecroq, 2004. Handbook of Exact String Matching Algorithms. First Edition. King's College London Publications. ISBN: 0954300645
- Knuth, D.E., J.H. Morris and V.R. Pratt, 1977. Fast pattern matching in strings. SIAM J. Comput., 6: 323-350.
- 7. Horspool, R.N., 1980. Practical fast searching in strings. Software Practice Experience, 10: 501-506.
- Berry, T. and S. Ravindran, 1999. A fast string matching algorithm and experimental results. In: Proceedings of the Prague Stringology Club Workshop '99, Liverpool John Moores University, pp: 16-28.
- Crochemore, M. and D. Perrin, 1991. Two-way string-matching. ACM, 38: 651-675. DOI: http://doi.acm.org/10.1145/116825.116845
- Thathoo, R. *et al.*, 2006. TVSBS: A fast exact pattern matching algorithm for biological sequences. Current Sci., 91: 47-53. URL: http://profile.iiita.ac.in/avirmani 02/TVSBS.pdf

- Boyer, R.S. and J.S. Moore, 1977. A fast string searching algorithm. Commun. ACM., 20: 762-772. DOI:10.1145/359842.359859
- Michael, T.G. and Roberto Tamassia, 2002. Algorithm Design, Foundations, Analysis and Internet Examples. First Edition. John Wiley and Sons, Inc, USA. ISBN: 0-471-38365-1
- He, L., F. Binxing and J. Sui, 2005. The wide window string matching algorithm. Theor. Compu. Sci., 332: 391-404. DOI: 10.1016/j.tcs.2004.12.002
- Hume, A. and D. Sunday, 1991. Fast string searching. Software Practice Experience, 21: 1221-1248. DOI: 10.1002/spe.4380211105
- Lecroq, T., 1995. Experimental results on string matching algorithms. Software-practice and Experience, 25: 727-765. DOI: 10.1002/spe.4380250703
- Davies G., and Bowsher S., 1996. Algorithms for pattern matching, Software-Practice and Experience,16:575-601. DOI:10.1002/spe.4380160608
- Tsai, T.H., 2003. Average case analysis of the boyer-moore algorithm. Source. Random Struct. Algorithms, 28: 481-498. DOI: 10.1002/rsa.v28:4
- Frantisek F., Christopher G. Jennings and W. F. Smyth, 2007. A simple fast hybrid patternmatching algorithm, J. Discrete Algorithms 5: 682– 695. DOI: 10.1016/j.jda.2006.11.004
- T. Bell, J. Cleary, and I. Witten, Text Compression. First Edition. Englewood Cliffs, N.J.: Prentice Hall, 1990. ISBN: 0-13-911991-4