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The complexity of the Approximate Multiple Pattern Matching Problem for random strings

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15 – Abstract

We describe a multiple string pattern matching algorithm which is well-suited for approximate 16 search and dictionaries composed of words of different lengths. We prove that this algorithm has 17 optimal complexity rate up to a multiplicative constant, for arbitrary dictionaries. This extends to 18 arbitrary dictionaries the classical results of Yao [SIAM J. Comput. 8, 1979], and Chang and Marr 19

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1 The problem 27

1.1 Definition of the problem 28

Let Σ be an alphabet of s symbols, $\xi = \xi_0 \dots \xi_{n-1} \in \Sigma^n$ a word of n characters (the input 29 text string), $D = \{w_1, \ldots, w_\ell\}, w_i \in \Sigma^*$ a collection of words (the *dictionary*). We say that 30 $w = x_1 \dots x_m$ occurs in ξ with final position j if $w = \xi_{j-m+1}\xi_{j-m+2}\cdots\xi_j$. We say that w 31 occurs in ξ with final position j, with no more than k errors, if the letters x_1, \ldots, x_m can 32 be aligned to the letters $\xi_{j-m'}, \ldots, \xi_j$ with no more than k errors of insertion, deletion or 33 substitution type, i.e., it has *Levenshtein distance* at most k to the string $\xi_{i-m'} \dots \xi_i$ (see an 34 example in Figure 1). Let $r_m(D)$ be the number of distinct words of length m in D. We call 35 $\mathbf{r}(D) = \{r_m(D)\}_{m>1}$ the content of D, a notion of crucial importance in this paper. 36

The approximate multiple string pattern matching problem (AMPMP), for the datum 37 (D,ξ,k) , is the problem of identifying all the pairs (a,j) such that $w_a \in D$ occurs in ξ with 38 final position j, and no more than k errors (cf. Figure 1). This is a two-fold generalisation of 39 the classical string pattern matching problem (PMP), for which the exact search is considered, 40 and the dictionary consists of a single word. 41

A precise historical account of this problem, and a number of theoretical facts, are 42 presented in Navarro's review [8]. The first seminal works have concerned the PMP. Results 43



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Figure 1 Typical output of an approximate multiple string pattern matching problem, on an English text (alphabet of 26 characters plus the space symbol .). In this case k = 2 and r(D) = (0, 0, 0, 0, 1, 0, 2, 1, 0, ...). The symbols D, S and ⁱ stand for deletion, substitution and insertion errors, while X corresponds to an insertion or a substitution.

included the design of efficient algorithms (notably Knuth-Morris-Pratt and Boyer-Moore), 44 and have led to the far-reaching definition of the Aho–Corasick automata [1, 3, 7, 11]. In 45 particular, Yao [11] is the first paper that provide rigorous bounds for the complexity of 46 PMP in random texts. To make a long story short, it is argued that an interesting notion of 47 complexity is the asymptotic average fraction of text that needs to be accessed (in particular, 48 at least at this stage, it is not the time complexity of the algorithm), and is of order $\ln(m)/m$ 49 for a word of length m. The first works on approximate search, yet again for a single word 50 (APMP), are the description of the appropriate data structure, in [10, 4], and, more relevant 51 to our aims here, the derivation of rigorous complexity bounds in Chang and Marr [5]. Yet 52 again in simplified terms, if we allow for k errors, the complexity result of Yao is deformed 53 into order $[\ln(m) + k]/m$. More recent works have concerned the case of dictionaries composed 54 of several words, all of the same length [9],¹ however, also at the light of unfortunate flaws in 55 previous literature, the rigorous derivation of the average complexity for the MPMP has been 56 missing even in the case of words of the same length, up to our recent paper [2], where it is 57 established that the Yao scaling $\ln(m)/m$ is (roughly) modified into $\max_m \ln(mr_m)/m$ (a 58 more precise expression is given later on). By combining the formula of Chang and Marr for 59 APMP, and our formula for MPMP, it is thus natural to expect that the AMPMP may have 60 a complexity of the order $\max_m [\ln(mr_m) + k]/m$. This paper has the aim of establishing a 61 result in this fashion. 62

Of course, the present work uses results, ideas and techniques already presented in [2], 63 for the PMPM. A main difference is that in [2] we show that, for any dictionary, a slight 64 modification of an algorithm by Fredriksson and Grabowski [6] is optimal within a constant, 65 while this is not true anymore for approximate search with Levenshtein distance (we expect 66 that it remains optimal for approximate search in which only substitution errors are allowed, 67 although we do not investigate this idea here). As a result, we have to modify this algorithm 68 more substantially, by combining it with the algorithmic strategy presented in Chang and 69 Marr [5], and including one more parameter (to be tuned for optimality). This generalised 70 algorithm is presented in Section 2.2. 71

Also, a large part of our work in [2] is devoted to the determination of a relatively tight lower bound, while the determination of the upper bound consists of a simple complexity analysis of the Fredriksson–Grabowski algorithm. Here, instead, we will make considerable efforts in order to determine an upper bound for the complexity of our algorithm, which is

¹ This is the reason why, before our paper [2], which deals with dictionaries having words of different length, the forementioned notion of "content" of a dictionary did not appear in the literature.

	exact	approximate
single word	$C_{\text{Yao}} \frac{\ln m}{m}$ (Yao)	$C_{\rm CM} \frac{\ln m + k}{m}$ (Chang and Marr)
dictionary	$C_1^{\text{ex}} \frac{1}{m_{\min}} + C_2^{\text{ex}} \max_m \frac{\ln(smr_m)}{m}$	$\frac{C_1k + C_1'}{m_{\min}} + C_2 \max_m \frac{\ln(smr_m)}{m}$

Table 1 Summary of average complexities for exact and approximate search, for a single word or on arbitrary dictionaries. The results are derived from Yao [11], Chang and Marr [5], our previous paper [2], and the present paper, respectively.

the content of Section 2.4, while we will content ourselves of a rather crude lower bound, derived with small effort in Section 1.3 by combining the results of [5] and [2].

78 1.2 Complexity of pattern matching problems

In our previous paper [2] we have established a lower bound for the (exact search) multiple pattern matching problem, in terms of the size s of the alphabet, and the content $\mathbf{r} = \{r_m\}$ of the dictionary, involving the length m_{\min} of the shortest word in the dictionary, and a function $\phi(\mathbf{r})$ with the specially simple structure $\phi(\mathbf{r}) = \max_m f(m, r_m)$. More precisely, calling $\Phi_{\text{aver}}(\mathbf{r})$ (resp. $\Phi_{\max}(\mathbf{r})$) the average over random texts, of the average (res. maximum) over dictionaries D of content \mathbf{r} , of the asymptotic fraction of text characters that need to be accessed, we have

Theorem 1 (Bassino, Rakotoarimalala and Sportiello, [2]). Let $s \ge 2$ and $m_{\min} \ge 2$, and define $\kappa_s = 5\sqrt{s}$. For all contents \mathbf{r} , the complexity of the MPMP on an alphabet of size ssatisfies the bounds

⁸⁹
$$\frac{1}{\kappa_s} \left(\phi(\boldsymbol{r}) + \frac{1}{2s \, m_{\min}} \right) \leqslant \Phi_{\text{aver}}(\boldsymbol{r}) \leqslant \Phi_{\max}(\boldsymbol{r}) \leqslant 2 \left(\phi(\boldsymbol{r}) + \frac{1}{2s \, m_{\min}} \right) ,$$
 (1)

90 where

$$_{91} \qquad \phi(\boldsymbol{r}) := \max_{m} \frac{1}{m} \ln(s \, m \, r_m) \,. \tag{2}$$

Note a relative factor ln s between the statement of the result above, and its original
formulation in [2], due to a slightly different definition of complexity.

As we have anticipated, such a result is in agreement with the result of Yao [11], for 94 dictionaries composed of a single word, which is simply of the form $\ln(m)/m$. Combining 95 this formula with the complexity result for APMP, derived in Chang and Marr [5], it 96 is natural to expect that the AMPMP has a complexity whose functional dependence 97 on k and r is as in Table 1. Indeed, the bottom-right corner of the table is consistent 98 both with the entry above it, and the entry at its left. Furthermore, it is easily seen 99 that, up to redefining the constants, several other natural guesses would have this same 100 functional form in disguise. Let us give some examples of this mechanism. Write $X \ge$ 101 $a_{L/U}Y + b_{L/U}Z$ as a shortcut for $a_LY + b_LZ \leq X \leq a_UY + b_UZ$. Now, suppose that we 102 establish that $\Phi(\mathbf{r}, k) \ge a_{L/U} (k+1)/m_{\min} + b_{L/U} \max_m (\ln(mr_m) + k)/m$. Then we also 103 have $\Phi(\mathbf{r},k) \ge a'_{L/U}(k+1)/m_{\min} + b_{L/U} \max_m \ln(mr_m)/m$, with $a'_U = a_U + b_U$ (and all 104 other constants unchanged). On the other side, if we have $\Phi(\mathbf{r}, k) \ge a_{L/U}(k+1)/m_{\min} + c_{L/U}(k+1)/m_{\min}$ 105 $b_{L/U} \max_{m} \ln(mr_m)/m$, with $a_L > b_L$, then we also have $\Phi(\mathbf{r}, k) \geq a_{L/U} (k+1)/m_{\min} + b_L$ 106 $b'_{L/U} \max_{m} (\ln(mr_m) + k)/m$, with $b'_{L} = a_{L} - b_{L}$. 107

¹⁰⁸ The precise result that we obtain in this paper is the following:

Theorem 2. For the AMPMP, with k errors and a dictionary D of content $\{r_m\}$, the complexity rate $\Phi(D)$ is bounded in terms of the quantity

11
$$\tilde{\Phi}(D) := \frac{C_1(k+1)}{m_{\min}} + C_2 \max_m \frac{\ln(smr_m)}{m}$$
 (3)

112 as

1

¹¹³
$$\frac{1}{C_1 + \kappa_s C_2} \widetilde{\Phi}(D) \leqslant \Phi(D) \leqslant \widetilde{\Phi}(D),$$
 (4)

114 with $a = \ln(2s^2/(2s+1))$, $a' = \ln(4s^2-1)$, $\kappa_s = 5\sqrt{s}$ (as in Theorem 1) and

¹¹⁵₁₁₆
$$C_1 = \frac{a+2a'}{a};$$
 $C_2 = \frac{2(a+2a')}{aa'} = \frac{2}{a'}C_1.$ (5)

117 **1.3** The lower bound

¹¹⁸ Now, let us derive a lower bound of the functional form as in Table 1 for the AMPMP, by ¹¹⁹ combining our results in [2] for the MPMP and the results in [5] for the APMP. Let us first ¹²⁰ observe a simple fact. Suppose that we have two bounds $A^{\text{LB}}(\mathbf{r}, k) \leq \Phi(\mathbf{r}, k) \leq A^{\text{UB}}(\mathbf{r}, k)$ ¹²¹ and $B^{\text{LB}}(\mathbf{r}, k) \leq \Phi(\mathbf{r}, k) \leq B^{\text{UB}}(\mathbf{r}, k)$ (with $A^{\text{LB}}(\mathbf{r}, k)$ and $B^{\text{LB}}(\mathbf{r}, k)$ positive). Then, for all ¹²² functions $p(\mathbf{r}, k)$, valued in [0, 1], we have

$$p(\boldsymbol{r},k)A^{\mathrm{LB}}(\boldsymbol{r},k) + (1-p(\boldsymbol{r},k))B^{\mathrm{LB}}(\boldsymbol{r},k) \leqslant \Phi(\boldsymbol{r},k) \leqslant A^{\mathrm{UB}}(\boldsymbol{r},k) + B^{\mathrm{UB}}(\boldsymbol{r},k).$$

We want to exploit this fact by using as bounds $A^{\text{LB}/\text{UB}}(\boldsymbol{r},k)$ our previous result for the 124 exact search, and as lower bound $B^{\text{LB}}(\boldsymbol{r},k)$ the simple quantity $(k+1)/m_{\text{min}}$. Then, later 125 on, in Section 2, we will work on the determination of a bound $B^{\text{UB}}(\boldsymbol{r},k)$ which has the 126 appropriate form for our strategy above to apply. Let us discuss why $\Phi(\mathbf{r}, k) \ge (k+1)/m_{\min}$. 127 We will prove that this quantity is a bound to the minimal density of a *certificate*, over a 128 single word of length $m = m_{\min}$, and text ξ . A certificate, as described in [11], is a subset 129 $I \subseteq \{1, \ldots, n\}$ such that, for the given text, the characters $\{\xi_i\}_{i \in I}$ imply that no occurrences 130 of words of the dictionary may be possible, besides the ones which are fully contained in I. 131 Some reflection shows that: (1) for the interesting case m > k, the smallest density |I|/n of 132 a certificate is realised on a *negative certificate*, that is, on a text ξ with no occurrences of 133 the word w; (2) the smallest density is realised, for example, by the text $\xi = bbb \cdots b$, and 134 the word $w = aaa \cdots a$; (3) in such a certificate, we must have read at least k + 1 characters 135 in every interval of size m, otherwise the alignment of w to this portion of text, in which we 136 perform all the substitutions on the disclosed characters, would still be a viable candidate. 137 Note in particular that deletion and insertion errors do not lead to higher lower bounds 138 (although, for large m, they lead to bounds which are only slightly smaller). 139

As a result, recalling the expression for the lower bound in Theorem 1, by choosing $p(\mathbf{r}, k)$ to satisfy $\frac{p}{1-p} = \frac{\kappa_s C_2}{C_1}$ we have

$$_{^{142}} \Phi(\boldsymbol{r},k) \ge (1-p)\frac{k+1}{m_{\min}} + \frac{p}{\kappa_s}\phi(\boldsymbol{r}) = \frac{p}{\kappa_s C_2} \left(C_1 \frac{k+1}{m_{\min}} + C_2 \phi(\boldsymbol{r}) \right) = \frac{C_1 \frac{k+1}{m_{\min}} + C_2 \phi(\boldsymbol{r})}{C_1 + \kappa_s C_2} \,.$$

This proves the lower bound part of Theorem 2. Note that we could confine all the dependence from $\{r_m\}$ to the function ϕ (in particular, the choice $\frac{p}{1-p} = \frac{\kappa_s C_2}{C_1}$ only depends on the size of the alphabet s).

¹⁴⁶ **2** The (q, L) search algorithm, and the upper bound

¹⁴⁷ 2.1 Definition of alignment

We define a partial alignment α of the word $w = x_1 \dots x_m$ to the portion of text $\xi_{i_1} \dots \xi_{i_2}$, with k errors, and boundary parameters $(\varepsilon, \varepsilon') \in \mathbb{N}$, as the datum $\alpha = (w; i_1, i_2; \varepsilon, \varepsilon'; u)$, where u is a string in $\{C, S_a, D, I_a\}^*$, (these letters stand for correct, substitution, deletion and insertion, respectively, and the index a runs from 1 to s). Two integer parameters (for example i_2 and ε') are not independent, as they are deduced (say) from i_1, ε and the length of u. Indeed, say that the string u has m_C symbols C, m_D symbols D, m_S symbols of type S_a (for all a's altogether) and m_I symbols of type I_a , then

155	$k = m_S + m_D + m_I$	(number of errors)
156	$\varepsilon + \varepsilon' = m - (m_C + m_S + m_D)$	(portion of the word on the sides)
157 158	$i_2 - i_1 + 1 = m_C + m_S + m_I$	(length of the aligned portion of text)

¹⁵⁹ The alignment has the following pattern (with a dash – denoting a skipped character, in the text or in the word):

161

For example, if w = counteroffers, in our reference text of Figure 1 we have the alignment $\alpha = (w; i_1, i_2; \varepsilon, \varepsilon'; u) = (w; 14, 24; 3, 1; u)$ with k = 4 and $u = CCCCI_{\Box}CCS_{\Box}S_{o}I_{u}C$, as indeed

$$i_{\underline{1}} = 14 \qquad i_{\underline{2}} = 24$$

$$\xi = \cdots \text{ the } . \text{ winter } . \text{ of } . \text{ our } . \text{ discontent}$$

$$u = CCCCI_{\Box}CCS_{\Box}S_{\circ}I_{u}C$$

$$w = \underbrace{\operatorname{counter } r \text{ of } r \text{ er } r \text{ s}}_{\varepsilon = 3}$$

This example shows an important feature of this notion: several strings u may correspond 166 to equivalent alignments among the same word and the same portion of text, and with the 167 same offset ε . For example, the three last errors of $u = \cdots S_{\Box}S_{\circ}I_{u}C$ can be replaced as in 168 $u' = \cdots S_{\Box} I_0 S_u C$ or as in $u'' = \cdots I_{\Box} S_0 S_u C$. As the underlying idea in producing an upper 169 bound from an explicit algorithm is to analyse the algorithm while using the union bound on 170 the possible alignments, it will be useful to recognise classes of equivalent alignments, and, 171 in the bound, 'count' just the classes, instead of the elements (we are more precise on this 172 notion in Section 2.3). 173

We define a *full alignment* to be likewise a partial alignment, but with $\varepsilon = \varepsilon' = 0$. That is, the goal of any algorithm for the AMPMP is to output the list of (say) positions i_2 of the full alignments among the given text and dictionary. Note that we can always complete a partial alignment with k errors and boundary parameters ($\varepsilon, \varepsilon'$) to a full alignment with no more than $k + \varepsilon + \varepsilon'$ errors, and no less than k errors, by including substitution or insertion errors at the two sides.

We define a *c-block partial alignment* as the generalisation of the notion of partial alignment to the case in which the portion of text consists of c non-adjacent blocks. In this

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Figure 2 Typical outcome for the search of the pattern deformed pattern in our reference text. In this example L = 3 and q = 12, the number of full blocks is $c(\alpha) = 2$, and can be aligned to the disclosed portion of the text (denoted by underline) with k = 3 errors: one deletion on the first block, one insertion in the second block, and one deletion somewhere in between the two blocks. On the bottom line, another alignment of the same word, in which, instead of inserting the letter **r** in the second block, we have substituted **n** by **r**, still with k = 3. These two alignments are sufficiently different to contribute separately to our estimate of the complexity, within our version of the union bound (because the values of ε are different).

case, besides the natural alignment parameters ε , ε' , and $i_{1,a}$, $i_{2,a}$, and u_a , for the blocks $a = 1, \ldots, c$, we have c - 1 parameters $\delta_a \in \mathbb{Z}$, associated to the offset between the alignment of the word to the blocks with index a and a + 1. As a result, in order to extend a c-block partial alignment to a full alignment, we need to perform at least $-\delta_a$ further insertion errors, $+\delta_a$ further deletion errors, depending on the sign of δ_a , for each of the c - 1 intervals between the portions of text. That is, any c-block partial alignment α with k errors can be completed to a full alignment with no less than $k + \sum_a |\delta_a|$ errors.

Note that in the following we will *not* need to count all of the possible ways in which these deletions or insertions can be performed, as it may seem natural in a naïve perspective on the use of the union bound. This fact will allow us to efficiently bound the number of possible multi-block partial alignments arising in our algorithm analysis (instead of counting directly the possible full alignments, which would result in a too large bound).

¹⁹⁴ 2.2 The algorithm

¹⁹⁵ Here we introduce an algorithm for AMPMP, concentrating on the pertinent notion of ¹⁹⁶ complexity, which is the ratio between the number of accesses to the text and the length of ¹⁹⁷ the text, and neglecting all implementation issues, and analysis of time complexity.

The algorithm is determined by two integers q and L, such that $k+1 \leq L < q \leq m_{\min} - k$. 198 The emerging inequality $2k + 1 < m_{\min}$ is not a limitation, as when this inequality is not 199 satisfied we have to read a fraction $\Theta(1)$ of the text, and in this regime there is no point in 200 showing that some algorithm can reach a complexity which is optimal up to a multiplicative 201 constant. When L = 1, the algorithm coincides with the one described by Fredriksson and 202 Grabowski [6], and already analysed in detail in [2] for the MPMP. When we have a single 203 word of length m, and q has the maximal possible value q = m - k, the algorithm coincides 204 with the one used by Chang and Marr [5] for their proof of complexity of the APMP. As 205 we will see in Section 2.5, choosing the optimal values of q and L for a given dictionary D206 (when the words are of different length) is not a trivial task. 207

Call the interval $\xi_{bq}\xi_{bq+1}\cdots\xi_{bq+L-1}$ of the text ξ the *b*-th block of text. The text is thus 208 decomposed in a list of blocks of length L, and of intervals between the blocks, of length 209 q-L. To every possible full alignment α of the word w to the text, are associated two 210 integers: $c(\alpha)$ is the number of blocks which are fully contained in the alignment, and $b(\alpha)$ is 211 the index of the rightmost of these blocks. Furthermore, we define c(w) as the minimum of 212 $c(\alpha)$ among the possible alignments involving w (indeed, it is either $c(\alpha) = c(w)$ for all α , or 213 $c(\alpha) \in \{c(w), c(w) + 1\}$ for all α , and, of course, at fixed q and L, c(w) only depends on the 214 length |w| of the word). 215

Our algorithm accesses the text in three steps, namely, for every block index $b = 0, 1, \ldots, \lceil n/q \rceil - 1$:

We read all the characters ξ_i of the text, for $bq \leq i < bq + L$, that is we read the *b*-th block;

We consider the possible c-block partial alignments α (with $c = c(\alpha)$) such that $b(\alpha) = b$, and associated to the intervals of text read so far. If any of these alignments is not excluded or determined positively, we read also the characters ξ_i for i = bq - 1, bq - 2, ...,one by one, in this order, up to when all partial alignments are either excluded, or reach $\varepsilon = 0$. For a given instance of the problem, call $\mathcal{E}_L(b)$ (left-excess at block b) the set of positions of further characters that we need to access by this second step (with indices shifted so that the block starts at 1), and $e_L(b) = |\mathcal{E}_L(b)|$.

If at the previous step we still have partial alignments which are not excluded, we read also the characters at positions i = bq + L, bq + L + 1, ..., in this order, up to when all partial alignments are either excluded, or completed to a full alignment. Similarly to above, introduce $\mathcal{E}_R(b)$ and $e_R(b) = |\mathcal{E}_R(b)|$ (right-excess at block b).

An example with $c(\alpha) = 2$ is in Figure 2. Note that, at all steps, the pattern of the accessed part of the text consists of some blocks of length L and spacing q, plus one rightmost block with length $L' \ge L$ and spacing $q' \le q$. A typical situation within the second step is as follows (here c = 5, L = 3, q = 8, L' = 12 and q' = 7):

²³⁶ Call $\mathcal{E}(b) = \mathcal{E}_L(b) \cup \mathcal{E}_R(b)$, and $e(b) = e_L(b) + e_R(b)$. Call Ψ_h^{exact} the average over random ²³⁷ texts of the indicator function for the event that $e(b) \ge h$. Clearly, the average complexity ²³⁸ rate of our algorithm is bounded by the expression

$$_{^{239}} \qquad \Phi_{\text{alg}}(D) \leqslant \frac{L + \mathbb{E}(e(b))}{q} = \frac{L + \sum_{h \ge 1} \Psi_h^{\text{exact}}}{q} \,,$$

235

where the average is taken over random texts, at fixed dictionary. Note that, because of our choice of range for q and L, $c(\alpha) \ge 1$ for all α , and $c(|w|) \ge 1$ for all w.

Let α be a full alignment associated to the block *b*. Call $\mathcal{E}[\alpha]$ the set of extra positions of the text (besides the blocks) that we need to access in order to determine the alignment α . Then clearly $\mathcal{E}(b) = \bigcup_{\alpha} \mathcal{E}[\alpha]$.

245 **2.3** Proof strategy for the upper bound

Our proof strategy is to prove that there exists a choice of parameters L and q, with the properties that $q = \Theta(m_{\min})$, $L/q = \Theta(\phi(\mathbf{r}(D)))$, and $\mathbb{E}(e(b)) = \Theta(1)$. This last condition is equivalent to the requirement that Ψ_h^{exact} is a summable series, and we will see that indeed the first can be bounded by a geometric series, and the second is rather small. Up to calculating the pertinent multiplicative constants, such a pattern would imply the functional form of the complexity anticipated in Section 1.2.

The idea is that the exact calculation of $\mathbb{E}(e(b))$ or of Ψ_h^{exact} , even at q and L fixed (which is easier than optimising w.r.t. these parameters), is rather difficult, but we can produce a simpler upper bound by:

For alignments α with $c(\alpha) > 1$, neglect the information coming from the e(b') extra characters that we have accessed at blocks b' < b. This allows to separate the analysis on the different blocks of text.

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²⁵⁸ Naïvely, for different (full) alignments α , we could perform a *union bound*, that is, ²⁵⁹ $e(b) = |\mathcal{E}(b)| = |\bigcup_{\alpha} \mathcal{E}[\alpha]| \leq \sum_{\alpha} |\mathcal{E}[\alpha]|$, which thus separates the analysis over the ²⁶⁰ different alignments. We will make an improved version of this bound, namely we use this

bound, not with full alignments, but rather with "classes of equivalent partial alignments". As we anticipated, the crucial point is that we count partial alignments instead of full alignments. A further slight improvement of the bound comes from considering these 'classes of equivalent partial alignments', instead of just the partial alignments. These two facts are motivated by the same argument, that we now elucidate.

Consider the two following notions: (1) Each set $A_h(w)$ of partial alignments is partitioned into classes I. (2) There is a subset $\bar{A}_h(w) \subseteq A_h(w)$ of alignments, that we shall call *basic alignments*. Now, suppose that the two following properties hold: (i) $I \cap \bar{A}_h(w) \neq \emptyset$ for all classes I of $A_h(w)$. (ii) For each $\alpha \in I$, there exists a $\bar{\alpha} \in I \cap \bar{A}_h(w)$, such that $\mathcal{E}(\alpha) \subseteq \mathcal{E}(\bar{\alpha})$. In this case it is easily established that the bound above can be improved into $e(b) = |\mathcal{E}(b)| = |\bigcup_{\alpha} \mathcal{E}[\alpha]| \leq \sum_{\bar{\alpha}} |\mathcal{E}[\bar{\alpha}]|$, where the sum runs only on basic partial alignments. Thus, calling $\Psi_h := \sum_{w \in D} \sum_{\alpha \in \bar{A}_h(w)} \mathbb{P}(|\mathcal{E}[\bar{\alpha}]| \geq h)$, we have $\Psi_h \geq \Psi_h^{\text{exact}}$.

We propose the following definition of basic alignment. Let α be in $A_h(w)$. In the string u, suppose that we write C_a instead of C, whenever the well-aligned character is a, and D_a when the deleted character is a (this is clearly just a bijective decoration of u). For $\alpha \in \overline{A}_h(w)$, we require that there are no occurrences of $C_a I_a$ as factors of u (as these are equivalent to $I_a C_a$), of $C_a D_a$ (as these are equivalent to $D_a C_a$) and of $I_a D_b$ or $D_b I_a$ (as these are equivalent to C_a or S_a , depending if a = b or not). If α can be obtained from α' by a sequence of these rewriting rules, then α and α' are in the same class I.

It is easy to see that this definition of basic alignment and classes has the defining properties above.

²⁸² **2.4** Evaluation of an upper bound at q and L fixed

Let us call $p_{c,h,\varepsilon'}(w)$ the probability that, for a given word w and parameter ε' , there exists an alignment $\alpha \in A_h(w)$, to a text consisting of c-1 blocks of length L and one block of length L+h, which is visited by the algorithm (that is, it makes at most k errors), that is, in particular,

$$\Psi_h \leqslant \sum_{\varepsilon'=0}^{q-1} p_{c,h,\varepsilon'}(w) \,. \tag{6}$$

289 We have the important fact

▶ Proposition 3.

2

$$p_{c,h,\varepsilon'}(w) \leqslant \beta s^{-(cL+h)} B_{cL+h+c-1,k} \tag{7}$$

²⁹¹ for all ε' , where $\beta = \frac{(2s-1)L+k}{(2s-1)L-k}$ and $B_{L,k} = (2s-1)^k {\binom{L+k}{k}}$.

The proof of this proposition is slightly complicated, and is presented in Appendix A. Note however that for the special case c = 1, and with exactly k errors (instead of at most k errors), the bound $s^{-(L+h)}(2s)^k {L+k \choose k}$ can be established trivially. Also note that the bound does not depend on ε' , and, in particular, it only depends on $h = |\mathcal{E}_L| + |\mathcal{E}_R|$ for the alignments α at given w and ε' , and not separately on the two summands.

We are now ready to evaluate the expressions for the upper bound on the quantity Ψ_h in (6), in light of (7). Call $R_c = \sum_{m: c(m)=c} r_m = \sum_{m=qc+L-1}^{q(c+1)+L-2} r_m$, and $p_{c,h}$ as q times the

²⁹⁹ RHS of (7) (that is, an upper bound to $\sum_{\varepsilon'=0}^{q-1} p_{c,h,\varepsilon'}(w)$). We have the bound

$$\sum_{h} \Psi_{h} \leqslant \sum_{c} R_{c} \sum_{h} p_{c,h} = \sum_{c} R_{c} \sum_{h} \beta q \, s^{-(cL+h)} B_{cL+h+c-1,k} \,.$$
(8)

302 Recalling that

$$\sum_{h\geq 0} s^{-h} \binom{a+k+h}{k} \leqslant \frac{1}{1-\frac{1}{s}\frac{a+k+1}{a+1}} \binom{a+k}{k}$$

(and that $q < m_{\min}$), substituting in (8) gives

$$\Phi_{\text{alg}}(D) \leqslant \frac{1}{q} \left(L + \beta q \sum_{c} R_{c} \frac{1}{1 - \frac{1}{s} \frac{cL + k + c}{cL + c}} s^{-cL} {cL + c - 1 + k \choose k} (2s - 1)^{k} \right) \\
\leqslant \frac{1}{q} \left(L + \frac{\beta m_{\min}(2s - 1)^{k}}{1 - \frac{1}{s} \frac{L + k}{L}} \sum_{c} R_{c} s^{-cL} {c(L + 1) + k \choose k} \right).$$
(9)

306 We want to prove that

$$_{307} \qquad \Phi_{\text{alg}}(D) \leqslant \frac{C_1 k + C_1'}{m_{\min}} + C_2 \max_m \frac{\ln(smr_m)}{m},$$
(10)

with suitable constants C_1 , C'_1 and C_2 (it will turn out at the end that we can set $C'_1 = C_1$ and C_1 , C_2 to be as in Theorem 2, but at this point it is convenient to let them be three separate variables). This would prove the upper bound part of Theorem 2.

Note that, if $k/m_{\min} \ge 1/C_1$, the upper bound expression (10) is larger than the trivial bound $\Phi_{\text{alg}}(D) \le 1$, and there is nothing to prove. So we can assume that $k/m_{\min} < 1/C_1$.

313 **2.5** Optimisation of q and L

We now have to analyse the expression (9), in order to understand which values of q and Lmake the bound smaller. The sum over c is the most complicated term. We simplify it by using the fact that, for all $\xi \in \mathbb{R}^+$, $\ln {a+k \choose k} \leq k \ln(1+\xi) + a \ln(1+\xi^{-1})$, which gives

317
$$T := m_{\min}(2s-1)^k \sum_c R_c \, s^{-cL} \binom{c(L+1)+k}{k}$$

305

$$\leq \sum_{c} \frac{1}{c^2} \exp\left[-c\left(LA - \frac{1}{c}\left(\ln(R_c m_{\min}) + k\ln((1+\xi)(2s-1))\right) - \frac{\ln c^2}{c} - \ln(1+\xi^{-1})\right)\right]$$
$$= \sum_{c} \frac{1}{c^2} \exp\left[-c(LA - \phi'(c) - \ln(1+\xi^{-1}))\right],$$
(11)

321 where $A = \ln(s\xi/(1+\xi))$, $A' = \ln((1+\xi)(2s-1))$ and

$$_{322} \qquad \phi'(c) = \frac{\ln(c^2 R_c m_{\min}) + kA'}{c} \,. \tag{12}$$

Ultimately, we want to choose L such that T is bounded by a constant, as its summands over c are bounded by a convergent series. With this goal, let c^* be the value maximising the expression $\phi'(c)$, and ϕ^* the value of the maximum. The sum above is then bounded by

³²⁶
$$\sum_{c} \frac{1}{c^2} \exp[-c(LA - \phi^* - \ln(1 + \xi^{-1}))].$$

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For any value of ξ such that A > 0 (that is, for $\xi > (s-1)^{-1}$), there exists a positive smallest

value of
$$L$$
 such that the exponent in the expression above is negative. So we set

329
$$L^* = \left\lceil \frac{\phi^* + \ln(1 + \xi^{-1})}{A} \right\rceil$$
,

(as the choice of ξ is free, we can tune it at the end so that the ratio is an integer), and recognise that the RHS of equation (11), specialised to $L = L^*$, is bounded by $\sum_c \frac{1}{c^2} = \pi^2/6$. Note that

334 so that

335
$$\frac{L^*}{k} \ge \frac{A'}{A} = \frac{\ln((1+\xi)(2s-1))}{\ln(s\xi/(1+\xi))},$$

which implies that we can set $\beta = \frac{2s-1+A/A'}{2s-1-A/A'}$, and

$$_{337} \qquad \frac{1}{1 - \frac{1}{s}\frac{L+k}{L}} \leqslant \frac{1}{1 - \frac{1}{s}(1 + A/A')} = \frac{1}{1 - \frac{1}{s}\frac{\ln(s\xi(2s-1))}{\ln((1+\xi)(2s-1))}} \,.$$

Now, let us choose $q = \lfloor \frac{m_{\min}-k}{2} \rfloor$, which coincides with the choice of the analogous parameter in Chang and Marr [5]. This is the largest possible value such that $c(w) \ge 1$ for all $w \in D$. With this choice,

$$_{341} \qquad \frac{1}{q} \leqslant \frac{2}{m_{\min}} \, \frac{C_1}{C_1 - 1} \, .$$

³⁴² Collecting the various factors calculated above, we get that the expression (9) is bounded by

$$\Phi_{\text{alg}}(D) \leqslant \frac{2}{m_{\min}} \frac{C_1}{C_1 - 1} \left(L^* + \frac{\beta \frac{\pi^2}{6}}{1 - \frac{1}{s} (1 + A/A')} \right).$$

We are left with two tasks: choosing suitable values for ξ and C_1 (both of order 1), and recognising that the expression for L^* (and for ϕ^*) can be related to the quantity $\phi(\mathbf{r})$ in (2). Let us start from the latter. Note that, as for any $m \ge m_{\min}$

$$_{^{347}} \qquad \frac{m-k}{q} - 2 \leqslant c(m) \leqslant \frac{m}{q}$$

we can write² $m \leq m_{\min}c(m) \leq s^2m$, which gives

349
$$\max_{c} \frac{1}{c} \ln(c^2 m_{\min} R_c) \leq \max_{m} \frac{m_{\min}}{m} \ln(s^2 m^2 r_m) \leq 2m_{\min} \phi(\mathbf{r}).$$

As, of course $\max_c(X(c) + Y(c)) \leq \max_c X(c) + \max_c Y(c)$, we have in particular that

$$\phi^* \leq 2m_{\min}\phi(\mathbf{r}) + kA' \qquad L^* \leq \frac{2m_{\min}\phi(\mathbf{r}) + kA' + \ln(1+\xi^{-1})}{A} ,$$

² Because $s \ge 2$, and we anticipate that, under our choice, $C_1 \ge 5$, thus

$$m \leq 2(m-k-q) \leq m_{\min}\left(\frac{m-k}{q}-2\right) \leq m_{\min}c(m) \leq m_{\min}\frac{m}{q} \leq 2\left(\frac{C_1}{C_1-1}\right)m \leq s^2m$$
.



Figure 3 Plot of the constant $C_1(s)$, $C'_1(s)$ and $C_2(s)$, as given by the expressions in (13) (respectively, in blue, green and red). The asymptotic values are 5, $5\pi^2/12$ and 0 respectively.

353 which thus implies

$$\begin{split} \Phi_{\mathrm{alg}}(D) &\leqslant \frac{2}{m_{\min}} \frac{C_1}{C_1 - 1} \left(\frac{2m_{\min}}{A} \phi(\mathbf{r}) + k \frac{A'}{A} + \frac{\beta \frac{\pi^2}{6}}{1 - \frac{1}{s}(1 + A/A')} + \frac{\ln(1 + \xi^{-1})}{A} \right) \\ &= \frac{2C_1}{C_1 - 1} \left[\frac{A'}{A} \frac{k}{m_{\min}} + \left(\frac{\beta \frac{\pi^2}{6}}{1 - \frac{1}{s}(1 + \frac{A}{A'})} + \frac{\ln(1 + \xi^{-1})}{A} \right) \frac{1}{m_{\min}} + \frac{2}{A} \phi(\mathbf{r}) \right] \,. \end{split}$$

354

Let us choose $C_1 = 2A'/A + 1$. The expression above simplifies into

$$\Phi_{\rm alg}(D) \leqslant \frac{C_1 k}{m_{\rm min}} + \frac{2A' + A}{AA'} \left[\left(\frac{A\beta \frac{\pi^2}{6}}{1 - \frac{1}{s}(1 + \frac{A}{A'})} + \ln(1 + \xi^{-1}) \right) \frac{1}{m_{\rm min}} + 2\phi(\boldsymbol{r}) \right],$$

³⁵⁷ in particular, this justifies the notation C_1 , which in the introduction was chosen to denote ³⁵⁸ the coefficient in front of the $\frac{k}{m_{\min}}$ summand. Now we shall choose the optimal value of ξ . ³⁵⁹ The dependence on ξ is mild, provided that we are in the appropriate range $\xi > 1/(s-1)$. ³⁶⁰ The choice of ξ , in turns, determines the ratio between the lower and upper bound, which ³⁶¹ has the functional form $C'_1 + \kappa_s C_2$ (with notations as in the theorem). A choice which is ³⁶² a good trade-off among the three summands in this expression, and for which the analytic ³⁶³ expression is relatively simple, is to take $\xi = 2s$. Under this choice we have

$$\begin{array}{ll} _{364} & \quad C_1 = 1 + 2 \frac{\ln(4s^2 - 1)}{\ln(2s^2/(2s + 1))} \,, \qquad C_2 = \frac{4}{\ln(2s^2/(2s + 1))} + \frac{2}{\ln(4s^2 - 1)} \,, \\ _{365} & \quad C_1' = \frac{C_2}{2} \left[\ln \frac{2s + 1}{2s} + \frac{\beta \pi^2}{6} \frac{s \ln(2s^2/(2s + 1)) \ln(4s^2 - 1)}{(s - 1) \ln(4s^2 - 1) - \ln(2s^2/(2s + 1)))} \right] \end{array}$$

or, in a more compact way, calling $a = A|_{\xi=2s} = \ln(2s^2/(2s+1))$ and $a' = A'|_{\xi=2s} = \ln(4s^2-1)$, and substituting back the value of β ,

₃₆₉
$$C_1 = \frac{a+2a'}{a},$$
 $C_2 = \frac{a+2a'}{a}\frac{2}{a'},$ (13a)

$$C_{1}^{370} = \frac{a+2a'}{a} \left(\frac{\ln s - a}{a'} + \frac{\pi^{2}}{6} \frac{(2s-1)a' + a}{(2s-1)a' - a} \frac{as}{(s-1)a' - a} \right).$$
(13b)

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The behaviour in s of these constants is depicted in Figure 3.

It can be verified that, with our choice of ξ , $C'_1 < C_1$ for all $s \ge 2.^3$ we can replace C'_1 by C_1 in the functional form (10) for the bound on $\Phi_{\text{alg}}(D)$, and thus obtain the statement of

³⁷⁵ Theorem 2. This concludes our proof.

³⁷⁶ — References

377	1	Alfred V. Aho and Margaret J. Corasick. Efficient string matching: an aid to bibliographic
378		search. Communications of the ACM, 18(6):333–340, 1975.

 Frédérique Bassino, Tsinjo Rakotoarimalala, and Andrea Sportiello. The complexity of the Multiple Pattern Matching Problem for random strings. In 2018 Proceedings of the Fifteenth Workshop on Analytic Algorithmics and Combinatorics (ANALCO), pages 40–53. SIAM, 2018.

Robert S Boyer and J Strother Moore. A fast string searching algorithm. Communications of the ACM, 20(10):762-772, 1977.

William I. Chang and Eugene L. Lawler. Sublinear approximate string matching and biological
 applications. Algorithmica, 12(4-5):327–344, 1994.

William I. Chang and Thomas G. Marr. Approximate string matching and local similarity.
 In Combinatorial Pattern Matching - CPM 1994, volume 807 of Lecture Notes in Computer
 Science, pages 259–273. Springer, 1994.

Kimmo Fredriksson and Szymon Grabowski. Average-optimal string matching. Journal of Discrete Algorithms, 7(4):579–594, 2009.

7 Donald E. Knuth, James H. Morris, and Vaughan R. Pratt. Fast pattern matching in strings.
 SIAM Journal on Computing, 6(2):323–350, 1977.

³⁹³ 8 Gonzalo Navarro. A guided tour to approximate string matching. ACM Computing Surveys
 (CSUR), 33(1):31-88, 2001.

³⁹⁵ 9 Gonzalo Navarro and Kimmo Fredriksson. Average complexity of exact and approximate
 ³⁹⁶ multiple string matching. *Theoretical Computer Science*, 321(2-3):283-290, 2004.

Peter H. Sellers. The theory and computation of evolutionary distances: pattern recognition.
 Journal of Algorithms, 1(4):359–373, 1980.

Andrew Chi-Chih Yao. The complexity of pattern matching for a random string. SIAM
 Journal on Computing, 8(3):368-387, 1979.

³ One way to see this is by proving that both $C_1(s)$ and $C'_1(s)$ decrease monotonically as functions on the real interval $[2, +\infty[$, that $\lim_{s\to\infty} C_1(s) = 5$, that $C_1(s) > C'_1(s)$ for $s \in \{2, 3, \ldots, 7\}$, and that $C'_1(8) < 5$.

A Proof of Proposition 3

401

In this section we evaluate an upper bound to $p_{c,h,\varepsilon'}$, which is the probability that, for a given word w with c(|w|) = c, the disclosed text composed of c-1 intervals of size L and one interval of size L + h corresponds to at least one basic alignment α by making no more than k errors. The statement of the result, equation (14) below, is given in Proposition 3. Let us introduce the recurring quantity

407
$$B_{L,k} := (2s-1)^k \binom{L+k}{k}$$

First, let us analyse the case in which we have a single block, and exactly k errors. For w a word of length m, it is clear that the result depends only on the $m - \varepsilon'$ left-most characters of the word, not on the ε' right-most ones, so we can assume without loss of generality that $\varepsilon' = 0$. Call $H_{L,k}(m)$ the number of different words of length L obtained by transforming the suffixes of w and making exactly k errors. We have

⁴¹³ \triangleright Proposition A.1. For all $L \ge k \ge 1$, $H_{L,k} \le B_{L,k}$.

⁴¹⁴ **Proof.** Note that the analogous statement with 2s - 1 replaced by 2s in $B_{L,k}$ is trivial, as we ⁴¹⁵ have exactly 2s types of errors (one deletion, s insertions and s - 1 substitutions), and the ⁴¹⁶ counting of their possible positions in the string u is a function of the length of the string, ⁴¹⁷ bounded from above by the worst case, associated to all insertion errors.

We can gain the factor 2s - 1 instead of 2s by restricting to basic alignments, but this 418 requires a finer analysis involving generating functions. Let us call f(u, y, z) the generating 419 function such that $[u^a y^L z^k] f(u, y, z)$ is the number of basic alignments of length L obtained 420 by transforming a word of length a and making exactly k errors. Calculating f(u, y, z) exactly 421 is a difficult task, and the result would depend on w as a word, not only on m = |w|, but we 422 will calculate a simpler upper bound $f_{\rm UB}(u, y, z)$, which in particular only depends on m. In 423 this context, a generating-function upper bound is an upper bound for partial sums, that 424 is $g \succeq f$ if $\sum_{h=0}^{k} [u^a y^L z^h](g(u, y, z) - f(u, y, z)) \ge 0$ for all L and a. Let us construct f_{UB} by starting from $f_0(u, y, z) := \frac{uy}{1-uy}$, which is the generating function f specialised to z = 0, 425 426 and let us introduce the various types of errors one at the time. 427

The first operation corresponds to allow for *insertion* errors. The restriction to basic 428 alignments, however, brings to a subtlety. For example, starting with a word w = abcd, in 429 order to get the alignment *aaabcd* we can proceed in several ways: *aaabcd* or *aaabcd* or by 430 *aaabcd* (bold letters correspond to insertions). Under the notion of basic alignment we avoid 431 to overcount these manifestly equivalent alignments, as of these expressions we would only 432 keep the latter, aaabcd, that is, at the left of a letter a we can only insert letters different 433 from a. On the other hand, at the right end of the word one can insert strings consisting of 434 any character of the alphabet. 435

 f_{ii} Calling f_i the generating function in which insertion errors are allowed, we thus get

$$_{437} \qquad f_i(u,y,z) = \frac{1}{1 - syz} f(u,y,z) \big|_{uy \to uy\left(\frac{1}{1 - (s-1)z}\right)} = \frac{uy}{(1 - syz)(1 - uy - (s-1)yz)}$$

We now introduce deletion errors, which, consistently, we allow only on the characters of the initial string (not on the ones which have just been insterted). Thus, any given original character can be either left as is, or deleted. This gives the generating function $f_{i,d}$, with

441
$$f_{i,d}(u,y,z) = f_i(u,y,z)|_{uy \to uy+uz} = \frac{uz+uy}{(1-syz)(1-uy-uz-(s-1)yz)}$$

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Finally, for substitution errors, again we can either substitute any initial character with one of the s-1 other characters of the alphabet, or leave it unchanged, which brings to $f_{i,d,s}$, with

445
$$f_{i,d,s} = f_{i,d}(u,y,z)|_{uy \to uy+(s-1)uyz} = \frac{u(syz - yz + y + z)}{(1 - syz)(1 - uz - (s-1)(u+1)yz)}$$

⁴⁴⁶ Note that, by this procedure, we have already produced an upper bound, as $f_{i,d,s} \succeq f$ (in ⁴⁴⁷ the sense defined above). Note also that it is *not* $f_{i,d,s} = f$, because, for example, we have ⁴⁴⁸ overcounted the equivalent cases in which in a word $w = \cdots aa \cdots$ we have deleted the first ⁴⁴⁹ or the second character.

If the word w is shorter than L + k, we may miss some alignments because they do not fit in the text interval. As we are evaluating an upper bound, we can restrict to the case in which w is long enough for this not to happen, and thus sum over all suffixes by just setting u = 1, and conclude that $H_{L,k} \leq [y^L z^k] f'(1, y, z)$. Thus, in order to conclude, we must show that $[y^L z^k] f'(1, y, z) \leq B_{L,k}$. Let us call

455
$$F_{L,k} = [y^L z^k] \frac{1}{(syz - 1)(2syz - 2yz + y + z - 1)}$$

We can rewrite the inequality above as $H_{L,k} \leq F_{L-1,k} + F_{L,k-1} + (s-1)F_{L-1,k-1}$, and thus, if we can prove that $F_{L,k} \leq B_{L,k}$, for all pairs of integers $L \geq k$, we could conclude in light of the fact that

⁴⁵⁹
$$H_{L,k} \leq B_{L-1,k} + B_{L,k-1} + (s-1)B_{L-1,k-1} = (2s-1)^k \binom{L+k}{k} - R_{L,k},$$

where $R_{L,k} = (2s-1)^{k-1} \left(2(s-1) \frac{k-1}{L} {L+k-2 \choose k-1} \right)$ is indeed easily checked to be non-negative for all $L \ge k \ge 1$.

462 So, to finish the proof, let us show that $F_{L,k} \leq B_{L,k}$. First,

$$F_{L,k} = [y^L z^k] \left(\frac{1}{1 - syz} + \frac{2syz - 2yz + y + z}{1 - 2syz + 2yz - y - z} \right)$$
$$= \delta_{L,k} s^k + F_{L-1,k} + F_{L,k-1} + 2(s-1)F_{L-1,k-1}$$

463

464 Since $L \ge k \ge 1$, we have $R_{L,k} \ge \delta_{L,k} s^k$ for $s \ge 2$, and $B_{L,k} \ge F_{L,k} \ge H_{L,k}$.

To conclude, we just check the boundary conditions in the recursion above for $F_{L,k}$ and $B_{L,k}$, which again are in agreement with the inequality. Indeed we have, for $(L,k) \in$ $\{(0,0), (0,1), (1,0)\}, F_{0,0} = B_{0,0} = 1, B_{0,1} = 2s - 1 \ge 1 = F_{0,1}$ and $B_{1,0} = 4s - 2 \ge 3s =$ $F_{1,0}$.

Now we want to deal with the more general case, in which we have more than one block, and we sum over the number of errors up to k. We will prove a more general statement, in which we have c blocks of lengths L_1, \ldots, L_c , separated by gaps of lengths q_1, \ldots, q_{c-1} , which in particular is so general to allow us to treat in one stroke the case in which we add characters at the left or at the right of the b-th algorithm block.

Similarly to the argument above, in order to produce an upper bound we can set without loss of generality that $\varepsilon' = 0$, all the q_i 's are larger than k and that m is larger than $\sum L_i + \sum q_i + k$, as any variant of this would give no more alignments. So, we will call $p_{L_1,...,L_c;k}$ the corresponding quantity, in which the dependence from the q_i 's and m has been dropped.



Figure 4 Example of multi-interval alignment analysed for the estimate of $p_{L_1,...,L_c;k}$.

For multi-block partial alignments, we have parameters $\delta_1, \ldots, \delta_{c-1}$ for the offset among the different consecutive blocks of the partial alignment, and, if we have an offset δ_i in the alignment of two blocks, we have to perform at least $|\delta_i|$ deletions or insertions errors when completing the partial alignment to a full one (cf. figure 4).

483 Calling $\bar{L} = \sum_{i=1}^{c} L_i$, this leads to the following sum

$$p_{L_1,...,L_c;k} \leqslant s^{-\bar{L}} \sum_{t=0}^k \sum_{\Delta=0}^t \sum_{\substack{k_1,k_2,...,k_c \in \mathbb{N} \\ k_1+k_2+...+k_c=t-\Delta}} \sum_{\substack{\delta_1,\delta_2,...,\delta_{c-1} \in \mathbb{Z} \\ \delta_1+\delta_2+...+\delta_{c-1}=\Delta}} B_{L_1,k_1} B_{L_2,k_2} \dots B_{L_c,k_c} \,.$$

From the Vandermonde convolution formula, $\sum_{i=0}^{k} {l_{1}+i \choose i} {l_{2}+k-i \choose k-i} = {l_{1}+l_{2}+k+1 \choose k}$, which implies $\sum_{h} B_{L_{1},h} B_{L_{2},k-h} = B_{L_{1}+L_{2}+1,k}$, we can simplify the expression above into

487
$$p_{L_1,...,L_c;k} \leqslant s^{-\bar{L}} \sum_{t=0}^k \sum_{\Delta=0}^t \sum_{\substack{\delta_1,\delta_2,...,\delta_{c-1} \in \mathbb{Z} \\ \delta_1+\delta_2+...+\delta_{c-1}=\Delta}} B_{\bar{L}+c-1,t-\Delta}.$$

488 The sum over the δ_i 's gives

$$\sum_{\substack{\delta_1,\delta_2,\ldots,\delta_{c-1}\in\mathbb{Z}\\\delta_1+\delta_2+\ldots+\delta_{c-1}=\Delta}} 1 = [z^{\Delta}] \left(\frac{1+z}{1-z}\right)^{c-1}$$

⁴⁹⁰ that is, by recognising that $B_{L,k-h} \leq B_{L,k} \left(\frac{k}{(2s-1)L}\right)^h$, we get

491
$$p_{L_1,...,L_c;k} \leqslant s^{-\bar{L}} B_{\bar{L}+c-1,k} \left(\frac{(1+z)^{c-1}}{(1-z)^c} \right) \Big|_{z=\frac{k}{(2s-1)(\bar{L}+c-1)}}.$$

This is all we shall say at this level of generality. Now note that, in our patterns, $\bar{L}+c-1 \ge cL$ (and $k \le L$), so that, in this range of parameters,

$${}^{494} \qquad p_{L_1,\dots,L_c;k} \leqslant s^{-\bar{L}} B_{\bar{L}+c-1,k} \left(\frac{(1+z)^{c-1}}{(1-z)^c} \right) \Big|_{z=\frac{1}{(2s-1)c}\frac{k}{L}} \leqslant \frac{(2s-1)+\frac{k}{L}}{(2s-1)-\frac{k}{L}} s^{-\bar{L}} B_{\bar{L}+c-1,k} .$$
(14)