

CRITICAL PHENOMENA IN SEQUENCE MATCHING

BY RICHARD ARRATIA¹ AND MICHAEL S. WATERMAN²

University of Southern California

Dedicated to the memory of Mark Kac.

We give a generalization of the result of Erdős and Rényi on the length R_n of the longest head run in the first n tosses of a coin. Consider two independent sequences, $X_1 X_2 \cdots X_m$ and $Y_1 Y_2 \cdots Y_n$. Suppose that X_1, X_2, \dots are i.i.d. μ , and Y_1, Y_2, \dots are i.i.d. ν , where μ and ν are possibly different distributions on a common finite alphabet S . Let $p \equiv P(X_1 = Y_1) \in (0, 1)$. The length of the longest matching consecutive subsequence is $M_{m,n} \equiv \max\{k: X_{i+r} = Y_{j+r} \text{ for } r = 1 \text{ to } k, \text{ for some } 0 \leq i \leq m - k, 0 \leq j \leq n - k\}$. For m and $n \rightarrow \infty$ with $\log(m)/\log(mn) \rightarrow \lambda \in (0, 1)$, our result is that there is a constant $K \equiv K(\mu, \nu, \lambda) \in (0, 1]$ such that $P(\lim M_{m,n}/\log_{1/p}(mn) = K) = 1$. The proof uses large deviation methods. The constant K is determined from a variational formula involving the Kullback-Liebler distance or relative entropy. A simple necessary and sufficient condition for $K = 1$ is given. For the case $m = n$ ($\lambda = 1/2$) and $\mu = \nu$, $K = 1$. The set of (μ, ν, λ) for which $K = 1$ has nonempty interior. The boundary of this set is the location of a phase transition. The results generalize to more than two sequences and to Markov chains. A strong law of large numbers is given for the proportion of letters within the longest matching word; the limiting proportion exhibits critical behavior, similar to that of K .

1. Introduction. This paper gives a generalization of the result of Erdős and Rényi on the length of the longest run of heads in the first n tosses of a coin. Our motivation is the comparison of DNA sequences, which are sometimes modeled as sequences of i.i.d. letters, or as letters of a Markov chain, with different distributions used for different sequences; see Smith, Waterman, and Sadler (1983).

Consider two sequences of length n , $X_1 X_2 \cdots X_n$ and $Y_1 Y_2 \cdots Y_n$. The length of the longest consecutive match, without shifts, is

$$(1) \quad R_n \equiv \max\{m: X_{i+k} = Y_{i+k} \text{ for } k = 1 \text{ to } m, \text{ for some } 0 \leq i \leq n - m\}.$$

The length of the longest consecutive match, allowing shifts, is

$$(2) \quad M_n \equiv \max\{m: X_{i+k} = Y_{j+k} \text{ for } k = 1 \text{ to } m, \text{ for some } 0 \leq i, j \leq n - m\}.$$

Suppose that the two sequences X_1, X_2, \dots and Y_1, Y_2, \dots are independent, with all letters chosen from a common finite alphabet S . Assume that X_1, X_2, \dots

Received March 1984; revised December 1984.

¹Supported by NSF Grant MCS-8301960.

²Supported by a grant from the System Development Foundation.

AMS 1980 subject classifications. Primary 60J10; secondary 68G10, 94A17.

Key words and phrases. Entropy, Kullback-Liebler distance, large deviations, sequence matching.

are i.i.d. (μ), and Y_1, Y_2, \dots are i.i.d. (ν), where μ and ν are probability distributions on S . Let $p \equiv P(X_1 = Y_1) = \sum_{a \in S} (\mu_a \nu_a)$, and assume that $p \in (0, 1)$.

To compute the length R_n of the longest match without shifts, the two sequences of letters may first be reduced to a single sequence of "heads" and "tails," with a "head" reported for the i th toss when $X_i = Y_i$. Thus R_n is the length of the longest head run in the first n tosses of a p -biased coin, described by the Erdős-Rényi law [Rényi (1970)]:

$$(3) \quad P\left(\lim_{n \rightarrow \infty} R_n / \log_{1/p}(n) = 1\right) = 1.$$

For the length M_n of the longest match with shifts, in the case $\mu = \nu$, it is shown in Arratia and Waterman (1985) that $P(\lim_{n \rightarrow \infty} M_n / \log_{1/p}(n) = 2) = 1$, so that

$$(4) \quad P\left(\lim_{n \rightarrow \infty} M_n / R_n = 2\right) = 1.$$

Loosely speaking, allowing shifts between two independent sequences *with the same distribution* doubles the length of the longest match. To see that M_n might grow like $2 \log_{1/p}(n)$, note that a match of length $m = \lceil 2 \log_{1/p}(n) \rceil$ starting from X_i and Y_j occurs with probability $p^m \approx n^{-2}$, which balances against $\approx n^2$ choices for (i, j) . However, if μ and ν are not "close," in a sense to be made precise later, then allowing shifts will not double the length of the longest match, i.e., (4) does not hold.

For a class of examples in which we can explicitly determine when allowing shifts doubles the length of the longest match, let X_1, X_2, \dots be a sequence of fair coin tosses, and let Y_1, Y_2, \dots be an independent sequence of biased coin tosses, with $\theta = P(Y_1 = \text{heads}) \in [0, 1]$. For all θ , $p = \frac{1}{2}$, so by (3), R_n grows like $\log_2(n)$. In the case $\theta = \frac{1}{2}$, the two sequences have the same distribution, so that M_n satisfies (4). In the case $\theta = 1$, the Y sequence is pure heads, so that $M_n \equiv R_n$ is the length of the longest head run in $X_1 X_2 \dots X_n$, i.e., allowing shifts has no effect on the length of the longest match, and (4) does not hold. What happens for intermediate cases, when one sequence represents a fair coin and the other sequence represents a biased but nondegenerate coin? Part of the answer, given by Theorem 1, is that (4) holds iff $\theta \in [x, 1 - x]$, where $x = 0.11002786 \dots$ is the smaller solution of $(x)\log(x) + (1 - x)\log(1 - x) = -(\log 2)/2$.

Theorem 1 states that if X_1, X_2, \dots is i.i.d. (μ) and Y_1, Y_2, \dots is i.i.d. (ν), with all letters independent and $p = P(X_1 = Y_1) \in (0, 1)$, then there exists a constant $C = C(\mu, \nu) \in [1, 2]$ such that

$$(5) \quad P\left(\lim_{n \rightarrow \infty} M_n / R_n = C\right) = 1.$$

[In the notation used in the summary and in Section 6, $C(\mu, \nu) \equiv 2K(\mu, \nu, 1/2)$.]

Let $\alpha \equiv \alpha(\mu, \nu)$ be the distribution on S corresponding to matching a single pair of letters:

$$(6) \quad \alpha_a \equiv (\mu_a \nu_a) / p = P(X_1 = Y_1 = a | X_1 = Y_1).$$

A necessary and sufficient condition for $C = 2$ is that

$$(7) \quad \begin{aligned} \sum (\mu_a \nu_a / p) \log(\mu_a) &\leq (\log p) / 2 \\ \text{and } \sum (\mu_a \nu_a / p) \log(\nu_a) &\leq (\log p) / 2, \end{aligned}$$

or equivalently, after a little manipulation,

$$(7') \quad H(\alpha, \nu) \leq (1/2) \log(1/p) \quad \text{and} \quad H(\alpha, \mu) \leq (1/2) \log(1/p).$$

Here $H(\cdot, \cdot)$ is the relative entropy or Kullback-Leibler distance: $H(\alpha, \nu) \equiv \sum \alpha_a \log(\alpha_a / \nu_a) \geq 0$, with $H(\alpha, \nu) = 0$ iff $\alpha = \nu$.

Let $H(\alpha) \equiv -\sum \alpha_a \log(\alpha_a) \geq 0$ be the entropy of α . Note that $H(\alpha, \mu) + H(\alpha, \nu) = -H(\alpha) + \log(1/p)$, so that if $H(\alpha, \mu) = H(\alpha, \nu)$ (in particular if $\mu = \nu$) then $H(\alpha, \mu) = H(\alpha, \nu) = [-H(\alpha) + \log(1/p)]/2 \leq (1/2) \log(1/p)$ so by (7'), $C(\mu, \nu) = 2$. Furthermore, if $\nu = \mu$ and μ is nontrivial, then α is nontrivial, so $H(\alpha) > 0$, and $H(\alpha, \mu) = H(\alpha, \nu) < (1/2) \log(1/p)$. It follows from (7') that for a fixed nontrivial distribution μ , $C(\mu, \nu) = 2$ for all distributions ν in some neighborhood of μ .

2. Further discussion. For any distributions μ and ν , it is very easy to get an upper bound on $M_n - 2 \log_{1/p}(n)$, as follows. For $m = 1, 2, \dots$, define the event

$$(8) \quad A_{ij} \equiv \{X_{i+1} \cdots X_{i+m} = Y_{j+1} \cdots Y_{j+m}\}$$

that some "witness" to the event $\{M \geq m\}$ appears at positions i in the X sequence and j in the Y sequence. Note that $P(A_{ij}) = p^m$ for each choice of i and j . Thus if $m = 2 \log_{1/p} n + b$ is an integer, so that $p^m = n^{-2} p^b$, we have $P(M \geq m) = P(\bigcup_{0 \leq i, j \leq n-m} A_{ij}) < n^2 p^m = p^b$. Write $\lfloor x \rfloor$ for the greatest integer $\leq x$, and write $\lceil x \rceil$ for the least integer $\geq x$. Using $m = \lceil (2 + \epsilon) \log_{1/p} n \rceil$ yields $P(M_n / (\log_{1/p} n) > 2 + \epsilon) < n^{-\epsilon}$, and an argument using the Borel-Cantelli lemma along a skeleton of times $n_k \equiv \lceil p^{-k} \rceil$ implies that $1 = P(\limsup(M_n / (\log_{1/p} n)) \leq 2)$.

The idea behind the proof of Theorem 1 is contained in the following calculation, which shows directly how condition (7) arises. Let $m = \lceil 2 \log_{1/p}(n) \rceil$, so that $n^2 p^m \in (p, 1]$. For each "word" $w \in S^m$ let E_w be the event that w appears within both $X_1 X_2 \cdots X_n$ and $Y_1 Y_2 \cdots Y_n$:

$$(9) \quad \begin{aligned} E_w &\equiv E_{w,n} \\ &\equiv \{w = X_{i+1} \cdots X_{i+m} = Y_{j+1} \cdots Y_{j+m} \text{ for some } 0 \leq i, j \leq n - m\}. \end{aligned}$$

From the independence of the two sequences,

$$(10) \quad \begin{aligned} &P(E_w) \\ &= P\left(\bigcup_{0 \leq i \leq n-m} \{w = X_{i+1} \cdots X_{i+m}\}\right) P\left(\bigcup_{0 \leq j \leq n-m} \{w = Y_{j+1} \cdots Y_{j+m}\}\right) \\ &< [(n \mu^m(w)) \wedge 1] [(n \nu^m(w)) \wedge 1]. \end{aligned}$$

Now with unions and sums taken over $w \in S^m$,

$$\begin{aligned}
 (11) \quad P(M \geq m) &= P\left(\bigcup_w E_w\right) \leq \sum_w P(E_w) \\
 &< \sum_w [(n\mu^m(w)) \wedge 1][(n\nu^m(w)) \wedge 1] \\
 (12) \quad &\leq \sum_w n\mu^m(w)n\nu^m(w) \\
 &= n^2 p^m \in (p, 1],
 \end{aligned}$$

using $p = \sum_a (\mu_a \nu_a)$ to get the final equality. By the weak law of large numbers, most of the contribution to the sum at (12) comes from words w in which the proportions of letters are approximately those of the distribution α at (6). The condition (7) is that for words w with proportions α , both $n\mu^m(w)$ and $n\nu^m(w)$ are not larger than $np^{m/2} \in (p^{1/2}, 1]$, so that the truncations " $\wedge 1$ " in the line preceding (12) have a negligible effect on the sum.

In the general case, $C = C(\mu, \nu) \in [1, 2]$ is defined by the requirement that with $m = \lceil C \log_{1/p} n \rceil$, the sum in (11) is ≈ 1 , in the sense that $0 = \lim_{n \rightarrow \infty} ((1/m) \log[\sum_{w \in S^m} P(E_w)])$. To show that $M_n/R_n \rightarrow C$ in probability, only minor modification of the above calculation is needed. The upper bound on M , $\forall \epsilon > 0 P(M_n/\log_{1/p} n > C + \epsilon) \rightarrow 0$, is easily proved; it suffices to use $m = \lceil (C + \epsilon) \log_{1/p} n \rceil$, and show that $P(M \geq m) = P(\bigcup_w E_w) \leq \sum P(E_w) \rightarrow 0$ as $n \rightarrow \infty$. To get the lower bound, $\forall \epsilon > 0 P(M_n/\log_{1/p} n > C - \epsilon) \rightarrow 1$, is more difficult; a bound on correlations is needed. For each word $w \in S^m$ consider the event G_w that w appears at a multiple of m within both $X_1 X_2 \cdots X_n$ and $Y_1 Y_2 \cdots Y_n$:

$$(13) \quad G_w \equiv G_{w,n} \equiv \bigcup_{0 \leq i, j \leq (n/m)-1} \{w = X_{im+1} \cdots X_{(i+1)m} = Y_{jm+1} \cdots Y_{(j+1)m}\},$$

so that $\bigcup_w G_w \subset \{M \geq m\}$. For $m = \lceil (C - \epsilon) \log_{1/p} n \rceil$, calculation shows that $\sum_w P(G_w) \rightarrow \infty$. For $w \neq v \in S^m$, the events G_w and G_v are negatively correlated (Lemma 1), so that $\sum P(G_w) \rightarrow \infty$ implies $P(\bigcup_w G_w) \rightarrow 1$.

3. Distinguishing matches by the proportions of letters involved. Let $\text{Pr}(S) = \{\gamma \in R^d: \gamma_a \geq 0, \sum \gamma_a = 1\}$ be the set of probability measures on our finite alphabet $S \equiv \{1, 2, \dots, d\}$, and for $m = 1, 2, \dots$, for any word $w \in S^m$, let $L(w) \in \text{Pr}(S)$ be the vector whose a th component is the proportion of letter a among the letters of w :

$$\text{for } a \in S, \quad L(w)_a \equiv (1/m) \sum_{1 \leq i \leq m} 1(w_i = a).$$

For $U \subset \text{Pr}(S)$ define the length $M_{n,U}$ of the longest match between $X_1 \cdots X_n$ and $Y_1 \cdots Y_n$ with proportions in U :

$$(14) \quad M_{n,U} \equiv \max\{m: X_{i+1} \cdots X_{i+m} = Y_{j+1} \cdots Y_{j+m} = w \text{ for some } w \text{ with } L(w) \in U, \text{ for some } 0 \leq i, j \leq n - m\}.$$

Given μ and ν with $p \equiv \sum \mu_a \nu_a \in (0, 1)$, for $\gamma, \beta \in \text{Pr}(S)$ and $c > 0$ define (with the convention that $\log 0 = -\infty$, but $0 \log 0 = 0$)

$$\begin{aligned}
 b(\gamma, \beta, c) &\equiv (1/c)\log(1/p) + \sum \gamma_a \log \beta_a, \\
 (15) \quad f(\gamma, c) &\equiv H(\gamma) + 0 \wedge b(\gamma, \mu, c) + 0 \wedge b(\gamma, \nu, c), \\
 g(\gamma) &\equiv \inf\{c: f(\gamma, c) < 0\}.
 \end{aligned}$$

Informally, $f(\gamma, c)$ represents $1/m$ times the log of the contribution to the sum in the line before (12), from words w having $L(w)$ near γ , if $m = \lceil c \log_{1/p} n \rceil$. Note that $f(\gamma, \cdot)$ is nonincreasing, and if $H(\gamma) > 0$ and $f(\gamma, c) = 0$, then $f(\gamma, \cdot)$ is strictly decreasing in a neighborhood of c . Thus for nontrivial $\gamma \in \text{Pr}(S)$, $g(\gamma)$ is the unique value for c for which $f(\gamma, c) = 0$. If $\gamma = \delta_a$ is the point mass on the letter $a \in S$ and $q \equiv \min(\mu_a, \nu_a) > 0$, then $g(\gamma) = (\log p)/(\log q) \in (0, 2)$.

The expressions for f and g in (15) allow a remarkable degree of simplification. Let $f_0(\gamma, c) \equiv H(\gamma)$; $f_1(\gamma, c) \equiv H(\gamma) + b(\gamma, \mu, c)$; $f_2(\gamma, c) \equiv H(\gamma) + b(\gamma, \nu, c)$; and $f_3(\gamma, c) \equiv H(\gamma) + b(\gamma, \mu, c) + b(\gamma, \nu, c)$; so that $f \equiv \min(f_0, f_1, f_2, f_3)$. For $i = 1, 2, 3$, define $g_i(\gamma)$ by the requirement that $f_i(\gamma, g_i(\gamma)) = 0$, so that $g(\gamma) = \min_{1 \leq i \leq 3} g_i(\gamma)$. Now

$$f_1(\gamma, c) \equiv -\sum \gamma_a \log \gamma_a - (\log p)/c + \sum \gamma_a \log \mu_a = -H(\gamma, \mu) - (\log p)/c,$$

so that $g_1(\gamma) = \log(1/p)/H(\gamma, \mu)$ and $g_2(\gamma) = \log(1/p)/H(\gamma, \nu)$. Also

$$\begin{aligned}
 f_3(\gamma, c) &\equiv -\sum \gamma_a \log \gamma_a - 2(\log p)/c + \sum \gamma_a \log \mu_a \nu_a \\
 &= \sum \gamma_a \log(\mu_a \nu_a / (p \gamma_a)) - (2 - c)(\log p)/c \\
 &= -H(\gamma, \alpha) - (2 - c)(\log p)/c,
 \end{aligned}$$

so that $g_3(\gamma) = (2 \log(1/p))/(\log(1/p) + H(\gamma, \alpha))$. Thus

$$(16) \quad g(\gamma) = \min \left\{ \frac{\log(1/p)}{H(\gamma, \mu)}, \frac{\log(1/p)}{H(\gamma, \nu)}, \frac{2 \log(1/p)}{\log(1/p) + H(\gamma, \alpha)} \right\}.$$

THEOREM 1. *If X_1, X_2, \dots are i.i.d. (μ) and Y_1, Y_2, \dots are i.i.d. (ν), with all letters independent and $p = P(X_1 = Y_1) \in (0, 1)$, then for any open $U \subset \text{Pr}(S)$, $M_{n,U}/(\log_{1/p} n)$ converges a.s. to $\sup_{\gamma \in U} g(\gamma)$. In particular, $1 = P(\lim_{n \rightarrow \infty} M_n / \log_{1/p} n = C(\mu, \nu))$ where*

$$C(\mu, \nu) = \sup_{\gamma \in \text{Pr}(S)} \min \left\{ \frac{\log(1/p)}{H(\gamma, \mu)}, \frac{\log(1/p)}{H(\gamma, \nu)}, \frac{2 \log(1/p)}{\log(1/p) + H(\gamma, \alpha)} \right\}$$

and $C(\mu, \nu) = 2$ if and only if both $H(\alpha, \nu), H(\alpha, \mu) \leq (1/2)\log(1/p)$.

PROOF. Fix an open nonempty set $U \subset \text{Pr}(S)$ and let $c = \sup_{\gamma \in U} g(\gamma)$.

First we prove the lower bound, that $P(M_{n,U} > (c - \epsilon)\log_{1/p} n \text{ eventually}) = 1$. If $c = 0$ (which occurs iff there is some letter $a \in S$ with $\alpha_a = 0$ and $\gamma_a > 0 \forall \gamma \in U$), then the lower bound is automatic. Assume that $c > 0$. Let $\epsilon > 0$ be given; we may assume that $\epsilon < c$. Fix a particular nontrivial $\beta \in U$ for which $g(\beta) > c - \epsilon$. From the strict monotonicity of $f(\beta, \cdot)$ in a neighborhood of $g(\beta)$,

it follows that $f(\beta, c - \epsilon) > 0$. Let $\delta = f(\beta, c - \epsilon)/5$. Fix an open set V with $\beta \in V \subset U$ for which the final two terms in expression (15) for $f(\cdot, c - \epsilon)$ vary by at most δ from their values at β , so that $\forall \gamma \in V, 0 \wedge b(\gamma, \mu, c - \epsilon) \geq 0 \wedge b(\beta, \mu, c - \epsilon) - \delta$, and similarly with ν in place of μ .

The number of words w of length m with proportions $L(w)$ in V is at least $\exp(m(H(\beta) - \delta))$, if m is sufficiently large, by Lemma 2. Let

$$T \equiv T(V, n, m) \equiv \sum_{w \in S^m: L(w) \in V} 1(G_{w,n}),$$

so that with $m = \lceil (c - \epsilon)\log_{1/p} n \rceil$,

$$\{T \neq 0\} = \bigcup_{w \in S^m: L(w) \in V} G_{w,n} \subset \{M_{n,V} > (c - \epsilon)\log_{1/p} n\}.$$

Using Lemma 3, for sufficiently large n we have

$$\begin{aligned} (1/m)\log(ET) &\geq H(\beta) - \delta \\ &\quad + 0 \wedge b(\beta, \mu, c - \epsilon) - \delta + 0 \wedge b(\beta, \nu, c - \epsilon) - \delta - \delta \\ &= f(\beta, c - \epsilon) - 4\delta = \delta > 0, \end{aligned}$$

so that $ET > \exp(m\delta)$ for large n . Using Chebyshev's inequality and then Lemma 1 to get $\text{var}(T) < ET$,

$$\begin{aligned} P(M_{n,V} > (c - \epsilon)\log_{1/p} n) &\geq P(T \neq 0) \\ &> 1 - \text{var}(T)/\{E(T)\}^2 \\ &> 1 - 1/E(T) \\ &> 1 - \exp(-m\delta). \end{aligned}$$

A Borel-Cantelli argument along the skeleton of times $n_k \equiv \lceil p^{-k} \rceil$ implies that $1 = P(M_{n,V} > (c - \epsilon)\log_{1/p} n \text{ eventually})$. Hence $1 = P(M_{n,U} > (c - \epsilon)\log_{1/p} n \text{ eventually})$.

Now we prove the upper bound. For each $\gamma \in U, c \geq g(\gamma)$ implies $f(\gamma, c + \epsilon/2) < 0$. Hence at least one of the two terms $b(\gamma, \mu, c + \epsilon/2), b(\gamma, \nu, c + \epsilon/2)$ is < 0 , and not controlled by the truncation with 0. With $\delta = (1/5)\log(1/p) [(c + \epsilon/2)^{-1} - (c + \epsilon)^{-1}]$, it follows that for all $\gamma \in U, f(\gamma, c + \epsilon) \leq -5\delta < 0$.

Each of the three terms in expression (15) for f is continuous, and $\text{Pr}(S)$ is compact, so that we can pick a finite collection $\{\gamma_i, V_i\}$ such that $U \subset \bigcup_i V_i$, and for each $i, \gamma_i \in V_i \subset U$, and for all $\gamma \in V_i, H(\gamma) < H(\gamma_i) + \delta, 0 \wedge b(\gamma, \mu, c + \epsilon) < 0 \wedge b(\gamma_i, \mu, c + \epsilon) + \delta$, and $0 \wedge b(\gamma, \nu, c + \epsilon) < 0 \wedge b(\gamma_i, \nu, c + \epsilon) + \delta$.

The number of words $w \in S^m$ with proportions $L(w) \in V_i$ is less than $\exp[m(H(\gamma_i) + 2\delta)]$, for sufficiently large m , by Lemma 2. Let

$$T_i \equiv T(V_i, n, m) \equiv \sum_{w \in S^m: L(w) \in V(i)} 1(E_{w,n}),$$

so that with $m = \lceil (c + \epsilon)\log_{1/p} n \rceil, \{M_{n,V(i)} \geq (c + \epsilon)\log_{1/p} n\} \subset \{T_i \neq 0\}$. Using

the upper bound on $P(E_{w,n})$ from Lemma 3, for large n we have

$$\begin{aligned} (1/m)\log(ET_i) &\leq H(\gamma_i) + 2\delta + b(\gamma_i, \mu, c + \epsilon) + \delta + b(\gamma_i, \nu, c + \epsilon) + \delta \\ &= f(\gamma_i, c + \epsilon) + 4\delta \leq -\delta < 0, \end{aligned}$$

so that $ET_i < \exp(-m\delta)$ for large n .

A Borel-Cantelli argument with $n_k \equiv \lfloor p^{-k} \rfloor$ implies that for each i , $0 = P(M_{n, \nu(i)} > (c + \epsilon)\log_{1/p} n$ infinitely often). Hence $1 = P(M_{n,U} < (c + \epsilon)\log_{1/p} n$ eventually). \square

LEMMA 1. *Let $X_1, X_2, \dots, Y_1, Y_2, \dots$ be independent S -valued variables, let integers m and n be fixed, and for any two distinct $w, v \in S^m$, consider the events G_w and G_v defined in (13). The events G_w and G_v are negatively correlated.*

PROOF. Writing $k \equiv \lfloor n/m \rfloor$, we have

$$\begin{aligned} P((G_w)^c \cap (G_v)^c) &= (1 - \mu^m(w) - \mu^m(v))^k (1 - \nu^m(w) - \nu^m(v))^k \\ &\leq (1 - \mu^m(w))^k (1 - \mu^m(v))^k (1 - \nu^m(w))^k (1 - \nu^m(v))^k \\ &= P((G_w)^c)P((G_v)^c). \end{aligned} \quad \square$$

LEMMA 2. *Let $S = \{1, 2, \dots, d\}$ and let $U \subset \text{Pr}(S)$ be an open subset of the set of probability measures on S . The number of words of length m with proportions in U grows like $\exp(m \sup_{\gamma \in U} H(\gamma))$, i.e.,*

$$\lim_{m \rightarrow \infty} (1/m)\log(|\{w \in S^m: L(w) \in U\}|) = \sup_{\gamma \in U} H(\gamma).$$

PROOF. This result is contained in the theory of large deviations of sums of independent R^d -valued random vectors, as in Bahadur (1971). We present a simple proof, in order to prepare the way for Lemma 4 and to keep this paper self-contained. Now $|\{w \in S^m: (L(w) \in U)\}| = \sum m! / (m_1! \cdots m_d!)$, where the sum is taken over integers m_1, \dots, m_d for which $\sum m_i = m$ and $\gamma = (m_1/m, \dots, m_d/m) \in U$. From $n \log n - n + 1 < \log(n!) < (n + 1)\log(n + 1) - n$ it follows that

$$\begin{aligned} H(\gamma) - m^{-1} \sum (1 + \log(m_i + 1)) &< m^{-1} \log(m! / [m_1! \cdots m_d!]) \\ &< H(\gamma) + m^{-1} \log m, \end{aligned}$$

where $\gamma = (m_1/m, \dots, m_d/m) \in \text{Pr}(S)$. The lower bound on $(1/m)\log(|\{w \in S^m: L(w) \in U\}|)$ is demonstrated by taking a single choice of (m_1, \dots, m_d) with proportions $\gamma = (m_1/m, \dots, m_d/m)$ whose entropy $H(\gamma)$ approximates $\sup_{\gamma \in U} H(\gamma)$. For the upper bound, note that the number of terms in the sum is $\leq m^d$, and $(1/m)\log(m^d) \rightarrow 0$ as $m \rightarrow \infty$. \square

LEMMA 3. *Suppose X_1, X_2, \dots are i.i.d. (μ) and Y_1, Y_2, \dots are i.i.d. (ν) , with all letters independent. Let $c > 0$ and $p \in (0, 1)$ be given and let $m \equiv$*

$m(n, c) \equiv \lceil c \log_{1/p} n \rceil$. Let $w \in S^m$ have proportions $L(w)$ such that $L(w_a) = 0$ whenever $\mu_a \nu_a = 0$. Then the function f defined in (15) and the events $E_{w,n}$ and $G_{w,n}$ defined in (9) and (13) satisfy

$$f(L(w), c) - H(L(w)) - \varepsilon < (1/m) \log P(G_{w,n}) < (1/m) \log P(E_{w,n}) < f(L(w), c) - H(L(w)),$$

where $\varepsilon = (2/m)[\log(4m) + \log(1/p)/c] \rightarrow 0$ as $n \rightarrow \infty$.

PROOF. To see the upper bound note that

$$(1/m)(\log n) \leq 1/(c \log_{1/p} n)(\log n) = \log(1/p)/c$$

and

$$(1/m) \log(\mu^m(w)) = (1/m) \sum_{1 \leq i \leq m} \log(\mu_{w(i)}) = \sum_{a \in S} L(w)_a \log(\mu_a),$$

so that

$$(1/m) \log[1 \wedge (n\mu^m(w))] \leq 0 \wedge b(L(w), \mu, c).$$

Thus

$$P(E_w) = P\left(\bigcup_{0 \leq i \leq n-m} \{w = X_{i+1} \cdots X_{i+m}\}\right) P\left(\bigcup_{0 \leq j \leq n-m} \{w = Y_{j+1} \cdots Y_{j+m}\}\right) < [(n\mu^m(w)) \wedge 1][(n\nu^m(w)) \wedge 1],$$

and hence $(1/m) \log P(E_{w,n}) < 0 \wedge b(L(w), \mu, c) + 0 \wedge b(L(w), \nu, c) = f(L(w), c) - H(L(w))$.

For the lower bound, by independence

$$P(G_w) = P\left(\bigcup_{0 \leq i \leq n/m-1} \{w = X_{mi+1} \cdots X_{mi+m}\}\right) \cdot P\left(\bigcup_{0 \leq j \leq n/m-1} \{w = Y_{mj+1} \cdots Y_{mj+m}\}\right) = [1 - (1 - \mu^m(w))^{\lfloor n/m \rfloor}][1 - (1 - \nu^m(w))^{\lfloor n/m \rfloor}].$$

For all $z \in [0, 1]$ and $n = 0, 1, 2, \dots$, $1 - (1 - z)^n \geq (1/2)(nz \wedge 1)$, so that

$$P(G_w) \geq (1/4)(\lfloor n/m \rfloor \mu^m(w) \wedge 1)(\lfloor n/m \rfloor \nu^m(w) \wedge 1) \geq 1/(4m^2)(n\mu^m(w) \wedge 1)(n\nu^m(w) \wedge 1).$$

Since $m - 1 \leq c \log_{1/p} n$, $(1/m) \log(n) \geq \log(1/p)/c - \log(1/p)/(mc)$. Thus for all n and w ,

$$(1/m) \log P(G_w) \geq f(L(w), c) - H(L(w)) - (2/m)[\log(2m) + \log(1/p)/c].$$

□

4. A strong law of large numbers. Informally, Theorem 1 says that for the longest consecutive match between $X_1 X_2 \cdots X_n$ and $Y_1 Y_2 \cdots Y_n$ with proportions near a given distribution γ , the length relative to $\log_{1/p} n$ tends almost

surely to $g(\gamma)$. Now the function $g: \text{Pr}(S) \rightarrow [0, 2]$ is continuous, and we will prove that g achieves its maximum $C(\mu, \nu)$ at a unique distribution β . It then follows easily from Theorem 1 that for any neighborhood U of β , the longest match with proportions in U will be longer than the longest match with proportions not in U , almost surely as $n \rightarrow \infty$. Thus the proportions of *all* matching words of maximal length tend almost surely to β , as $n \rightarrow \infty$.

Depending on μ and ν , the distribution α of letters in a simple match may or may not be the distribution β which maximizes g . For the coin tossing example discussed in Section 1, $\mu = (0.5, 0.5)$ and $\nu = \alpha = (1 - \theta, \theta)$, any case having $0 < H(\nu) < (1/2)\log 2$ gives an example with $\beta \neq \alpha$.

THEOREM 2. *In the setup of Theorem 1, there is a unique $\beta \in \text{Pr}(S)$ such that*

$$g(\beta) = C(\mu, \nu) = \sup_{\gamma \in \text{Pr}(S)} g(\gamma).$$

If $C(\mu, \nu) = 2$ (in particular, if $H(\alpha, \mu) = H(\alpha, \nu)$), then $\beta = \alpha$. As $n \rightarrow \infty$, the proportions of letters, in all words of maximal length common to both $X_1 X_2 \cdots X_n$ and $Y_1 Y_2 \cdots Y_n$, tend almost surely to β :

$$1 = P\left(0 = \limsup_{n \rightarrow \infty} \left\{ |\beta - L(w)| : w = X_{i+1} \cdots X_{i+m} = Y_{j+1} \cdots Y_{j+m} \right. \right. \\ \left. \left. \text{for some } 0 \leq i, j \leq n - m, \text{ with } m = M_n \right\} \right).$$

PROOF. To see that g achieves its maximum at a unique distribution β , consider the expression for g in (16): $g = \min_{1 \leq i \leq 3} g_i$. Since g_1 and g_2 have no local maxima in the interior of $\text{Pr}(S)$, g achieves its maximum either at α , where g_3 has its unique maximum, or else on one of the surfaces $g_i = g_j$. A maximum for g on the surface $g_1 = g_2$ is easily ruled out, since $g_1(\gamma) = g_2(\gamma) = c > 0$ implies $f_1(\gamma, c) = f_2(\gamma, c) = 0$ and thus $f_3(\gamma, c) = -H(\gamma) < 0$, so that $g_3(\gamma) < c$.

If $g_1(\gamma) = g_3(\gamma) = c > 0$ then $0 = f_1(\gamma, c) = f_3(\gamma, c)$ so that $f_2(\gamma, c) = H(\gamma) > 0$ and hence $g_2(\gamma) > c$ so that $g(\gamma) = c = g_3(\gamma)$. On the surface $\{\gamma: g_1(\gamma) = g_3(\gamma)\} \equiv \{\gamma: \log(1/p) + H(\gamma, \alpha) = 2H(\gamma, \mu)\}$, g_3 is maximized by minimizing $H(\gamma, \alpha)$. It follows from the strict convexity of $H(\cdot, \alpha)$ and of $H(\cdot, \mu)$ that there is a unique γ_μ which achieves this. Similarly, there is a unique γ_ν which maximizes $g(\gamma)$ given the constraint $g_2 = g_3$. It remains to show that $g(\gamma_\mu) \neq g(\gamma_\nu)$. If $g(\gamma_\mu) = g(\gamma_\nu)$, then $\gamma_\mu \neq \alpha$ so $2 > g_3(\gamma_\mu) = g_1(\gamma_\mu)$ and hence $H(\alpha, \mu) \geq H(\gamma_\mu, \mu) > (1/2)\log(1/p)$. The same argument yields $H(\alpha, \nu) > (1/2)\log(1/p)$, which is impossible, since $H(\alpha, \mu) + H(\alpha, \nu) = -H(\alpha) + \log(1/p) \leq \log(1/p)$. We have shown that there exist a distribution β such that $g(\beta) > g(\gamma)$ for all $\gamma \neq \beta$.

Since $g(\gamma) \leq (2 \log(1/p)) / (\log(1/p) + H(\gamma, \alpha))$ by (16), and $H(\gamma, \alpha) \geq 0$ with equality iff $\gamma = \alpha$, it follows that if $C(\mu, \nu) = 2$, then for $\gamma \neq \alpha$, $g(\gamma) < 2 = g(\alpha)$.

Given $\varepsilon > 0$, let $U = \{\gamma \in \text{Pr}(S): |\gamma - \beta| > \varepsilon\}$. Let $\delta = (1/2)(g(\beta) - \sup_{\gamma \in U} g(\gamma))$; $\delta > 0$ since $\text{Pr}(S)$ is compact and g is continuous. By Theorem 1, there is a random N which is almost surely finite, such that for all

$n > N$, $M_n/\log_{1/p} n > g(\beta) - \delta$ and $M_{n,U}/\log_{1/p} n < \sup_{\gamma \in U} g(\gamma) + \delta = g(\beta) - \delta$. Thus $n > N$ implies that $|\beta - L(w)| \leq \varepsilon$, for all w with $w = X_{i+1} \cdots X_{i+m} = Y_{j+1} \cdots Y_{j+m}$ for some $0 \leq i, j \leq n - m$, with $m = M_n$. \square

5. Matching between two different Markov processes. In this section we generalize Theorem 1 to the situation in which $X_1 X_2 \cdots X_n$ and $Y_1 Y_2 \cdots Y_n$ are independent sequences of letters governed by two different Markov transition mechanisms on the finite alphabet $S = \{1, 2, \dots, d\}$.

It is necessary to keep track of the proportions of *pairs* of letters that appear in adjacent positions. Note that for any word $w \in S^m$ and letter $i \in S$, the number of adjacent pairs in w that begin with i is equal to the number of pairs that end in i , provided that the word is wrapped around a circle so that the pair (last letter, first letter) is counted as one of the m pairs. Thus we define:

$$\text{for } w \in S^m, \quad \tilde{L}(w)_{ij} = (1/m) \sum_{1 \leq k \leq m} 1(w_k w_{k+1} = ij); \quad i, j \in S,$$

with the understanding that w_{m+1} is evaluated as w_1 . Let

$$B \equiv \left\{ q \in \text{Pr}(S^2) : \forall i, j \in S, q_{ij} > 0 \text{ and } \sum_{k \in S} q_{ik} = \sum_{k \in S} q_{ki} \right\},$$

be the set of strictly positive *balanced* proportions of pairs, so that for any word w , $\tilde{L}(w) \in \bar{B}$. For $q, r \in \text{Pr}(S^2)$ define

$$\tilde{H}(q) \equiv - \sum_{i, j \in S} q_{ij} \log \left(q_{ij} / \left(\sum_{k \in S} q_{ik} \right) \right)$$

and

$$\tilde{H}(q, r) \equiv \sum_{i, j} q_{ij} \log \left(\left(q_{ij} / \left(\sum_k q_{ik} \right) \right) / \left(r_{ij} / \left(\sum_k r_{ik} \right) \right) \right).$$

Note that if π and σ are the marginals of q and r , respectively, so that $\pi_i = \sum_k q_{ik}$ and $\sigma_i = \sum_k r_{ik}$, then $[q_{ij}/\pi_i]$ is the stochastic matrix governing a Markov process, and if $q \in B$ then (π_i) is the invariant measure: $\sum_i \pi_i (q_{ij}/\pi_i) = \sum_i q_{ij} = \pi_j$. Also $\tilde{H}(q, r) \geq 0$, with equality iff $q = r$, since $\tilde{H}(q, r) = \sum_i \pi_i [H(q_{i\cdot}/\pi_i, r_{i\cdot}/\sigma_i)]$. Note that $\tilde{H}(q, r) \leq H(q, r)$, with equality iff $\pi = \sigma$, since $\tilde{H}(q, r) = H(q, r) - H(\pi, \sigma)$.

LEMMA 4. Let $S = \{1, 2, \dots, d\}$ and let $U \subset \text{Pr}(S^2)$ be open. The number of words of length m with "proportions of pairs" in U grows like $\exp(m \sup_{q \in U \cap B} \tilde{H}(q))$, i.e.,

$$\lim_{m \rightarrow \infty} (1/m) \log(|\{w \in S^m : \tilde{L}(w) \in U\}|) = \sup_{q \in U \cap B} \tilde{H}(q).$$

PROOF. We give an elementary proof, but note that this result could also be proved by applying the large deviation theory in Donsker and Varadhan (1975) to the *two-step* Markov chain with state space S^2 and transition probabilities $p_{(i,j),(k,l)} = (1/d) \delta_{jk}$.

Let integers $m_{ij} > 1, i, j \in S$, be given, with the property that for each $i \in S$, $\sum_j m_{ij} = \sum_j m_{ji}$. Let $m = \sum_{ij} m_{ij}$ and let $m_i = \sum_j m_{ij}$, for each $i \in S$. Let $q_{ij} = m_{ij}/m$, for $i, j \in S$, so that $q \in B$. Elementary analysis of multinomial coefficients, as in Lemma 2, will complete the proof, once it is shown that

$$\prod_{i \in S} ((m_i - 1)! / (m_{i1}! \cdots (m_{i,i+1} - 1)! \cdots m_{id}!)) \leq |\{w \in S^m: \tilde{L}(w) = q\}| \leq |S| \prod_{i \in S} (m_i! / (m_{i1}! \cdots m_{id}!)),$$

with $d + 1$ identified as 1 in the lower bound. [The question of counting $\{w \in S^m: \tilde{L}(w) = q\}$ exactly is addressed in Billingsley (1961), Baum and Eagon (1966), and Zaman (1984).] A given word $w \in S^m$ with $\tilde{L}(w) = q$ determines, for each $i \in S$, a partition of the set $\{1, 2, \dots, m_i\}$ into subsets S_{i1}, \dots, S_{id} , with $|S_{ij}| = m_{ij}$ under the condition that $k \in S_{ij}$ if the k th appearance of letter i in the word is immediately followed by letter j . The word can be reconstructed from its starting letter w_1 and these partitions; this proves the upper bound.

The lower bound is the number of words satisfying the additional conditions that the last appearance of letter 1 is followed by letter 2, the last appearance of 2 is followed by a 3, ..., with the word ending in letter d . Let $n_{ij} = m_{ij} - \delta_{i,i+1}$, with the index $d + 1$ replaced by 1, so that $n_i = \sum_j n_{ij} = \sum_j n_{ji}$; i.e., $[n_{ij}]$ also satisfies the balance equations. Partition the set $\{1, 2, \dots, n_i\}$ into subsets S_{i1}, \dots, S_{id} , with $|S_{ij}| = n_{ij}$. These partitions determine a word w with $\tilde{L}(w) = q$, via the recipe: for $k \leq n_i$, the k th appearance of letter i is followed by letter j , iff $k \in S_{ij}$. The word begins with letter 1. When letter i appears for the $(1 + n_i)$ th time, all n_i pairs ending in i have been used up, and we put down a letter $i + 1$ and then continue to follow the partitions. This happens first with letter 1, then letter 2, ..., then letter d , at which point the word is completed. \square

THEOREM 3. *Let $X_1 X_2 \cdots$ and $Y_1 Y_2 \cdots$ be independent Markov chains on $S = \{1, 2, \dots, d\}$. Let $P = [p_{ij}]$ and $Q = [q_{ij}]$ be the transition matrices governing X and Y , respectively, with $p_{ij} > 0$ and $q_{ij} > 0$ for all $i, j \in S$. Let π and σ be the equilibrium distributions for X and Y , and define μ and $\nu \in B \subset \text{Pr}(S^2)$ by*

$$\mu_{ij} = \pi_i p_{ij}, \quad \nu_{ij} = \sigma_i q_{ij}, \quad i, j \in S.$$

Consider the substochastic matrix $R = [r_{ij}] \equiv [p_{ij} q_{ij}]$, and let $p, (r_i)$, and (l_i) be its principal eigenvalue and corresponding left and right positive eigenvectors, normalized so that $\sum l_i r_i = 1$. Since $[r_{ij} r_j / (p r_i)]$ is a stochastic matrix which governs a Markov process with equilibrium $(l_i r_i)$, we define $\alpha \in B$ by

$$\alpha_{ij} = l_i r_i r_j / p, \quad i, j \in S.$$

Define $\tilde{g}: \text{Pr}(S^2) \rightarrow [0, 2]$, using (16) with H replaced by \tilde{H} . Then for any open $U \subset \text{Pr}(S^2)$, $M_{n,U} / (\log_{1/p} n)$ converges a.s. to $\sup_{\gamma \in U \cap B} \tilde{g}(\gamma)$. In particular, $1 = P(\lim_{n \rightarrow \infty} M_n / \log_{1/p} n = C(P, Q))$, where

$$C(P, Q) = \sup_{\gamma \in \text{Pr}(S^2) \cap B} \min\{\log(1/p) / \tilde{H}(\gamma, \mu), \log(1/p) / \tilde{H}(\gamma, \nu), (2 \log(1/p) / (\log(1/p) + \tilde{H}(\gamma, \alpha)))\},$$

and $C(P, Q) = 2$ if and only if both $\tilde{H}(\alpha, \nu), \tilde{H}(\alpha, \mu) \leq (1/2)\log(1/p)$. Furthermore, there is a unique $\beta \in B$ such that $\tilde{g}(\beta) = C(P, Q)$ and

$$1 = P\left(0 = \limsup_{n \rightarrow \infty} \{|\beta - \tilde{L}(w)|: w = X_{i+1} \cdots X_{i+m} = Y_{j+1} \cdots Y_{j+m}\right.$$

for some $0 \leq i, j \leq n - m$, with $m = M_n\}$.

If $C(P, Q) = 2$, then $\beta = \alpha$.

PROOF. The proof follows those of Theorems 1 and 2, with minor changes such as the substitution of Lemma 4 in place of Lemma 2. In place of the events G_w involving nonoverlapping blocks of m letters, we apply Doeblin's method: Fix a letter $a \in S$ and consider blocks involving m successive returns to letter a . Details of this method in the context of matching with shifts are given in Arratia and Waterman (1985). The remaining modifications are routine. \square

6. Sequences with different lengths; more than two sequences. Comparison of DNA sequences often involves two sequences with very different lengths, such as 200 and 4000. Consider the length $M(m, n)$ of the longest consecutive matching between two sequences of lengths m and n , say $X_1 \cdots X_m$ and $Y_1 \cdots Y_n$. Even in the case where all $m + n$ letters are i.i.d., the limit of the ratio $(\log m)/(\log n)$ can have a critical role in determining first, whether or not $M(m, n)$ grows asymptotically like $\log_{1/p}(mn)$, and second, the composition of the best matching word.

Proceeding as in Section 3, we analyze $M(m, n)$ according to the proportions $L(w)$ of letters within the matching word w . Thus, for $U \subset \text{Pr}(S)$ let

$$M_U(m, n) \equiv \max\{t: X_{i+1} \cdots X_{i+t} = Y_{j+1} \cdots Y_{j+t} = w$$

for some w with $L(w) \in U$, for some $0 \leq i \leq m - t, 0 \leq j \leq n - t\}$,

so that when $U = \text{Pr}(S)$, $M_U(m, n) \equiv M(m, n)$.

THEOREM 4. Assume that X_1, X_2, \dots are i.i.d. (μ) and Y_1, Y_2, \dots are i.i.d. (ν) , with all letters independent and $p = P(X_1 = Y_1) \in (0, 1)$. Define $\alpha \in \text{Pr}(S)$ by $\alpha_a = \mu_a \nu_a / p$. Assume that m and $n \rightarrow \infty$, with $(\log m)/(\log mn) \rightarrow \lambda \in (0, 1)$. For $\lambda \in (0, 1)$ and $\gamma \in \text{Pr}(S)$ define

$$(17) \quad G(\gamma, \lambda) \equiv \min\{\lambda \log(1/p)/H(\gamma, \mu), (1 - \lambda) \log(1/p)/H(\gamma, \nu), \\ \log(1/p)/(\log(1/p) + H(\gamma, \alpha))\}.$$

Then for any open $U \subset \text{Pr}(S)$, $M_U(m, n)/(\log_{1/p}(mn))$ converges a.s. to $\sup_{\gamma \in U} G(\gamma, \lambda)$. In particular, with $K(\mu, \nu, \lambda) \equiv \sup_{\gamma \in \text{Pr}(S)} G(\gamma, \lambda) \in (0, 1]$, we have

$$1 = P\left(\lim_{n \rightarrow \infty} M(m, n)/\log_{1/p}(mn) = K(\mu, \nu, \lambda)\right), \text{ and}$$

$$(18) \quad K(\mu, \nu, \lambda) = 1 \quad \text{iff both } H(\alpha, \mu) \leq \lambda \log(1/p)$$

$$\text{and } H(\alpha, \nu) \leq (1 - \lambda) \log(1/p).$$

PROOF. The proof is very similar to the proof of Theorem 1. In place of f and g as defined at (15), we now use

$$F(\gamma, c, \lambda) \equiv H(\gamma) + 0 \wedge b(\gamma, \mu, c/\lambda) + 0 \wedge b(\gamma, \nu, c/(1 - \lambda)),$$

with the idea that $F(\gamma, c, \lambda)$ represents $1/t$ times the log of the contribution to $\sum_w [(m\mu^t(w)) \wedge 1][(n\nu^t(w)) \wedge 1]$, from words $w \in S^t$ having $L(w)$ near γ , when $t = \lceil c \log_{1/p}(mn) \rceil$. Elementary manipulation shows that $G(\gamma, \lambda) = \inf\{c: F(\gamma, c, \lambda) < 0\}$. The correspondence with the notation of Theorem 1 is that $F(\gamma, c, \frac{1}{2}) = f(\gamma, 2c)$, $2G(\gamma, \frac{1}{2}) = g(\gamma)$, and $2K(\mu, \nu, \frac{1}{2}) = C(\mu, \nu)$. \square

In the special case $\mu = \nu$, Theorem 4 says that if $(\log m)/(\log(mn)) \rightarrow \lambda \in (0, 1)$, then $M(m, n)$ is asymptotic to $\log_{1/p}(mn)$ iff $\lambda \in [\lambda_{cr}, 1 - \lambda_{cr}]$, where $\lambda_{cr} \equiv H(\alpha, \mu)/\log(1/p) \in [0, \frac{1}{2})$. Note that in this case, with $\mu = \nu$, the following are equivalent: $\lambda_{cr} = 0$; $H(\alpha, \mu) = 0$; $\alpha = \mu$; μ is the uniform distribution on S .

If $\beta \equiv \beta(\mu, \nu, \lambda)$ is the unique distribution on S for which $G(\beta, \lambda) = \sup_{\gamma \in \text{Pr}(S)} G(\gamma, \lambda)$, then as in Theorem 2, there is a strong law of large numbers for the composition of the best matching word: If m and $n \rightarrow \infty$ with $(\log m)/(\log(mn)) \rightarrow \lambda \in (0, 1)$, then with probability one, the proportions $L(w)$ of letters within any longest matching word w common to $X_1 \cdots X_m$ and $Y_1 \cdots Y_n$ tends to β . There are examples in which β varies nontrivially with λ , even with $\mu = \nu$, such as any biased coin tossing example, with $\mu = \nu = (1 - \theta, \theta)$, and $\theta \neq \frac{1}{2}$.

Theorem 1 can also be generalized to the case of $r \geq 2$ independent sequences, allowing r different distributions and r different lengths. As in Theorem 3, all of this can also be done for r independent Markov chains, allowing r different transition matrices. In either the i.i.d. or the Markov case, the expressions corresponding to F and G in the statement of Theorem 4 become quite complicated— F becomes the sum of $H(\gamma)$ plus r terms, each involving relative entropy and truncation, and the formula corresponding to (16) and (17) expresses G as a minimum of $2^r - 1$ smooth terms. The one result which remains reasonably simple is the necessary and sufficient condition for the length of the longest match to be asymptotic to $\log_{1/p}$ of the number of positions in which such a match might occur. This result is given, for the i.i.d. case, in Theorem 5.

THEOREM 5. *Suppose that for $j = 1$ to r , the letters X_1^j, X_2^j, \dots are i.i.d. (μ_j), where μ_1, \dots, μ_r are probability distributions on a finite alphabet S . Let $p \equiv \sum_{a \in S} \mu_1(a) \cdots \mu_r(a)$, and assume $p \in (0, 1)$. Define $\alpha \in \text{Pr}(S)$ by $\alpha(a) = \mu_1(a) \cdots \mu_r(a)/p$. Define the length $M \equiv M(n_1, \dots, n_r)$ of the longest word*

appearing, for $j = 1$ to r , within the first n_j letters of the j th sequence:

$$M \equiv \max \left\{ m: \phi \neq \bigcap_{j=1 \text{ to } r} \left\{ w \in S^m: X_{i+1}^j \cdots X_{i+m}^j = w, \right. \right. \\ \left. \left. \text{for some } 0 \leq i \leq n_j - m \right\} \right\}.$$

Suppose that $n_1, \dots, n_r \rightarrow \infty$ with $(\log n_j)/(\log(n_1 \cdots n_r)) \rightarrow \lambda_j > 0$, for $j = 1$

to r . Then there is a constant $K \equiv K(\mu_1, \dots, \mu_r; \lambda_1, \dots, \lambda_r) \in (0, 1]$ such that

$$1 = P(M/\log_{1/p}(n_1 \cdots n_r) \rightarrow K)$$

and

$$K = 1 \text{ iff } H(\alpha, \mu_j) \leq \lambda_j \log(1/p) \text{ for } j = 1 \text{ to } r.$$

PROOF. The argument is essentially the same as that for Theorems 1 and 4. \square

Acknowledgment. Guidance and motivation for this paper came from the work of Donsker and Varadhan (1975 and 1983).

REFERENCES

- ARRATIA, R. and WATERMAN, M. S. (1985). An Erdős-Rényi law with shifts. *Adv. in Math.* **55** 13-23.
- BAHADUR, R. R. (1971). *Some Limit Theorems in Statistics*. SIAM, Philadelphia.
- BAUM, L. E. and EAGON, J. A. (1966). The number of circular patterns compatible with a pseudo-symmetric connected graph. *Canad. J. Math.* **18** 237-239.
- BILLINGSLEY, P. (1961). Statistical methods in Markov chains. *Ann. Math. Statist.* **31** 12-40.
- DONSKER, M. D. and VARADHAN, S. R. S. (1975). Asymptotic evaluation for certain Markov process expectations for large time, I. *Comm. Pure Appl. Math.* **28** 1-47.
- DONSKER, M. D. and VARADHAN, S. R. S. (1983). Asymptotic evaluation for certain Markov process expectations for large time, IV. *Comm. Pure Appl. Math.* **36** 183-212.
- RÉNYI, A. (1970). *Foundations of Probability*. Holden-Day, San Francisco.
- SMITH, T. F., WATERMAN, M. S. and SADLER, J. R. (1983). Statistical characterization of nucleic acid sequence functional domains. *Nucleic Acids Res.* **11** 2205-2220.
- ZAMAN, A. (1984). Urn models for Markov exchangeability. *Ann. Probab.* **12** 223-229.

DEPARTMENT OF MATHEMATICS
UNIVERSITY OF SOUTHERN CALIFORNIA
LOS ANGELES, CALIFORNIA 90089-1113