## A CENTRAL LIMIT THEOREM FOR MARKOV PROCESSES THAT MOVE BY SMALL STEPS

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We consider a family  $X_n^\theta$  of discrete-time Markov processes indexed by a positive "step-size" parameter  $\theta$ . The conditional expectations of  $\Delta X_n^\theta$ ,  $(\Delta X_n^\theta)^2$ , and  $|\Delta X_n^\theta|^3$ , given  $X_n^\theta$ , are of the order of magnitude of  $\theta$ ,  $\theta^2$ , and  $\theta^3$ , respectively. Previous work has shown that there are functions f and g such that  $(X_n^\theta - f(n\theta))/\theta^{\frac{1}{2}}$  is asymptotically normally distributed, with mean 0 and variance g(t), as  $\theta \to 0$  and  $n\theta \to t < \infty$ . The present paper extends this result to  $t = \infty$ . The theory is illustrated by an application to the Wright-Fisher model for changes in gene frequency.

1. Introduction and overview. Let J be a bounded set of positive numbers with infimum 0. For every  $\theta \in J$ , let  $\{X_n^{\theta}\}_{n\geq 0}$  be a Markov process with stationary transition probabilities in a Borel subset  $I_{\theta}$  of the real line R. The parameter  $\theta$  is an index of the magnitude of  $\Delta X_n^{\theta} = X_{n+1}^{\theta} - X_n^{\theta}$ . We will be concerned with the asymptotic behavior of the distribution of  $X_n^{\theta}$  as  $n \to \infty$  and  $\theta \to 0$ .

The following assumptions, or their higher dimensional analogs, are in force throughout the paper:

$$E(\Delta X_n^{\theta} | X_n^{\theta} = x) = \theta w(x) + O(\theta^2)$$

(1.2) 
$$\operatorname{Var}\left(\Delta X_{n}^{\theta} \mid X_{n}^{\theta} = x\right) = \theta^{2} s(x) + o(\theta^{2})$$

(1.3) 
$$E(|\Delta X_n^{\theta}|^3 | X_n^{\theta} = x) = O(\theta^3),$$

uniformly over  $x \in I_{\theta}$ . Thus the error terms in (1.1) and (1.2) satisfy

$$\sup_{\theta \in J, x \in I_{\theta}} |O(\theta^2)|/\theta^2 < \infty$$

and

$$\sup\nolimits_{x\in I_{\theta}}|o\left(\theta^{2}\right)|/\theta^{2}\to0$$

as  $\theta \to 0$ . Let I be the closed convex hull of  $\bigcup_{\theta \in J} I_{\theta}$ . We assume that  $I_{\theta}$  approximates I as  $\theta \to 0$  in the sense that, for any  $x \in I$ ,

$$\inf_{y \in I_{\theta}} |y - x| \to 0$$

as  $\theta \to 0$ . The functions w and s are defined throughout I, s is Lipschitz, and w has a bounded Lipschitz derivative.

Under these assumptions the differential equations

$$f'(t) = w(f(t))$$

and

$$g'(t) = 2w'(f(t))g(t) + s(f(t))$$

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have unique solutions f(t) = f(t, x) and g(t) = g(t, x) with f(0) = x and g(0) = 0, where x is an arbitrary point of I. Suppose that  $x_{\theta} \in I_{\theta}$  and  $X_{\theta}^{\theta} = x_{\theta}$  a.s., and let

$$Z_n^{\theta} = (X_n^{\theta} - f(n\theta, x_{\theta}))/\theta^{\frac{1}{2}}$$
.

Let  $\mathcal{L}(Z)$  be the distribution of a random variable Z, and let  $\mathcal{N}(\mu, \sigma^2)$  be the normal distribution with mean  $\mu$  and variance  $\sigma^2$ . It has been established previously ([6] Theorem 8.1.1) that

$$\mathcal{L}(Z_n^{\theta}) \to \mathcal{N}(0, g(t, x))$$

as  $\theta \to 0$ ,  $x_{\theta} \to x$ , and  $n\theta \to t < \infty$ . Moreover, it can be shown that the distribution over C[0, T] of the random polygonal line  $Z^{\theta}$  with vertices  $Z^{\theta}(n\theta) = Z_n^{\theta}$  converges weakly to the distribution of the diffusion Z satisfying the stochastic differential equation

$$dZ(t) = w'(f(t))Z(t) dt + s(f(t))^{\frac{1}{2}} dB(t)$$

and the initial condition Z(0) = 0 a.s., where B is Brownian motion. Weak convergence theorems of this type have been established in similar contexts by Rosén [9] and Kurtz [4].

These results are the background for the present study. We shall consider the limit of  $\mathcal{L}(Z_n^{\theta})$  as  $\theta \to 0$  and  $n\theta \to \infty$  under certain additional assumptions. Our main project is to prove the following theorem, which was announced in ([8] Theorem 3.2(ii)).

Theorem 1. Suppose that I is bounded, w has a unique zero  $\lambda$ , and  $w'(\lambda) < 0$ . Then

$$(1.6) \mathcal{L}(Z_n^{\theta}) \to \mathcal{N}(0, g(\infty))$$

as  $\theta \to 0$  and  $n\theta \to \infty$ , where

$$g(\infty) = \lim_{t \to \infty} g(t, x) = s(\lambda)/2|w'(\lambda)|$$
.

The limiting process in Theorem 1 places no constraint on  $x_{\theta}$ . This implies that (1.6) holds uniformly over  $x_{\theta}$  in the following sense. Let d be a metric (or pseudometric) on probability distributions over R, such that  $d(\mathcal{L}_n, \mathcal{L}) \to 0$  whenever  $\mathcal{L}_n \to \mathcal{L}$  weakly. Then

$$\sup\nolimits_{x_{\theta}\in I_{\theta}}d(\mathcal{L}(Z_{n}^{\theta}),\,\mathcal{N}(0,\,g(\infty)))\to 0$$

as  $\theta \to 0$  and  $n\theta \to \infty$ . Proceeding further in this direction, we may combine (1.5) with Theorem 1 to obtain the rather striking conclusion that

(1.7) 
$$\sup_{n\geq 0, x_{\theta}\in I_{\theta}} D(n, \theta, x_{\theta}) \to 0$$

as  $\theta \rightarrow 0$ , where

$$D(n, \theta, x_{\theta}) = d(\mathcal{L}(Z_n^{\theta}), \mathcal{N}(0, g(n\theta, x_{\theta}))).$$

For if (1.7) were not true, there would be a c > 0 and sequences  $\theta_k$ ,  $n_k$ , and

 $x_k \in I_{\theta_k}$  such that  $\theta_k \to 0$  as  $k \to \infty$ , but

$$(1.8) D(n_k, \theta_k, x_k) \ge c$$

for all  $k \ge 1$ . We could, moreover, choose these sequences in such a way that  $n_k \theta_k \to t \le \infty$  and  $x_k \to x$ . It follows from (1.5) and (1.6) that  $\mathscr{L}(Z_n^{\theta}) \to \mathscr{N}(0, g(t, x))$  as  $k \to \infty$ , where  $\theta = \theta_k$ ,  $n = n_k$ , and  $g(\infty, x) = \lim_{t \to \infty} g(t, x)$ . Furthermore, it can be shown that g is continuous over  $[0, \infty] \times I$ , so  $\mathscr{N}(0, g(n_k \theta_k, x_k)) \to \mathscr{N}(0, g(t, x))$ . Therefore, by the triangle inequality,  $D(n_k, \theta_k, x_k) \to 0$  as  $k \to \infty$ , contradicting (1.8).

Another corollary of Theorem 1 is obtained by permitting  $\theta$  to approach 0 after  $n \to \infty$ . Suppose that  $\mathcal{L}(X_n^{\theta})$  converges weakly as  $n \to \infty$ , for every fixed  $\theta$ . Let  $\mathcal{L}(\theta)$  be the corresponding limit of  $\mathcal{L}(Z_n^{\theta})$ , or, equivalently, of  $\mathcal{L}(z_n^{\theta})$ , where

$$Z_n^{\theta} = (X_n^{\theta} - \lambda)/\theta^{\frac{1}{2}}$$

 $(f(t) \rightarrow \lambda \text{ as } t \rightarrow \infty)$ . It follows easily from (1.6) that

$$(1.9) \mathcal{L}(\theta) \to \mathcal{N}(0, g(\infty))$$

as  $\theta \to 0$ . Some special results of this type were established in [5]. It is also a consequence of Theorem 1 that (1.9) holds for an arbitrary family  $\mathcal{L}(\theta)$  of stationary distributions of  $z_n^{\theta}$ . This implies Part B of Theorem 10.1.1(i) of [6].

The proof of Theorem 1 is given in Sections 2 and 3. Section 4 presents an application to the Wright-Fisher model for the evolution of gene frequency under the influence of mutation, selection, and random drift. In that context,  $\theta = (2N)^{-\frac{1}{2}}$ , where N is the population size, and the function f is the classical deterministic approximation to gene frequency for very large populations (see [2] Section 2.3). The results (1.6) and (1.7), which relate to the distribution of the error of this approximation, appear to be new even for this much studied model.

Section 5 gives a multidimensional analog of Theorem 1, and an illustrative application to a mathematical learning model.

2. Conditional moments of  $\Delta Z_n^{\theta}$ . A basic component of the proof of Theorem 1 is Lemma 1.

Lemma 1. Under the hypotheses of Theorem 1,  $E((Z_n^{\theta})^2)$  is bounded over all  $\theta \in J$ ,  $x_{\theta} \in I_{\theta}$ , and  $n \geq 0$ .

This result follows immediately from Theorem 3.2(i) of [8]. The latter theorem assumes that J is an interval and  $I_{\theta} = I$  for all  $\theta$ , but these assumptions are not used in the proof. It emerges in the course of the proof that, for some  $K < \infty$  and  $\alpha > 0$ ,

$$|f(t, x) - \lambda| \le Ke^{-\alpha t}$$

for all  $x \in I$  and  $t \ge 0$ . Since w' and s satisfy Lipschitz conditions, we have

$$(2.1) |w'(f(t,x)) - w'(\lambda)| \le Ke^{-\alpha t}$$

and

$$|s(f(t, x)) - s(\lambda)| \le Ke^{-\alpha t}$$

for suitable new constants K.

Henceforth we suppress the  $\theta$  superscript on  $Z_n^{\theta}$ , and let  $\nu_n = f(n\theta, x_{\theta})$ . The purpose of this section is to establish Lemma 2.

LEMMA 2.

(2.3) 
$$E(\Delta Z_n | Z_n) = \theta w'(\nu_n) Z_n + o(\theta)$$

(2.4) 
$$E((\Delta Z_n)^2 | Z_n) = \theta s(\nu_n) + o(\theta)$$

$$E(|\Delta Z_n|^3 | Z_n) = o(\theta),$$

where the quantities  $o(\theta)$  satisfy  $E(|o(\theta)|)/\theta \to 0$  as  $\theta \to 0$ , uniformly over  $x_{\theta} \in I_{\theta}$  and  $n \ge 0$ .

Proof. Since w and w' are bounded,

$$f''(t) = w'(f(t))w(f(t))$$

is too. Thus

$$\Delta \nu_n = \theta w(\nu_n) + O(\theta^2)$$

uniformly over  $x_{\theta}$  and n. This expression and (1.1) imply that

(2.6) 
$$E(\Delta Z_n | Z_n) = \theta^{-\frac{1}{2}} (E(\Delta X_n | X_n) - \Delta \nu_n) \\ = \theta^{\frac{1}{2}} (w(X_n) - w(\nu_n)) + O(\theta^{\frac{3}{2}}).$$

Since w' is Lipschitz, this yields

$$E(\Delta Z_n | Z_n) = \theta w'(\nu_n) Z_n + \theta^{\frac{3}{2}} O(|Z_n|^2) + O(\theta^{\frac{3}{2}}),$$

which, in view of Lemma 1, is of the form (2.3).

Turning to the proof of (2.4), we begin by writing

(2.7) 
$$E((\Delta Z_n)^2 | Z_n) = \theta^{-1} \operatorname{Var} (\Delta X_n | X_n) + E(\Delta Z_n | Z_n)^2.$$

As a consequence of (2.6),

(2.8) 
$$E(\Delta Z_n | Z_n) = \theta O(|Z_n|) + O(\theta^{\frac{3}{2}}),$$

so that

$$E(\Delta Z_n \mid Z_n)^2 \leq K(\theta^2 \mid Z_n \mid^2 + \theta^3)$$

and

(2.9) 
$$E(\Delta Z_n | Z_n)^2 = o(\theta)$$

by Lemma 1. Next, (1.2) yields

(2.10) 
$$\theta^{-1} \operatorname{Var} (\Delta X_n | X_n) = \theta s(X_n) + o(\theta)$$
$$= \theta s(\nu_n) + \theta^{\frac{3}{2}} O(|Z_n|) + o(\theta)$$
$$= \theta s(\nu_n) + o(\theta)$$

by Lemma 1. Substituting (2.9) and (2.10) into (2.7), we obtain (2.4).

Finally,

$$E(|\Delta Z_n|^3 | Z_n) \le 4\theta^{-\frac{3}{2}} (E(|\Delta X_n|^3 | X_n) + |\nu_n|^3)$$
  
\$\leq K\theta^\frac{3}{2}\$

as a consequence of (1.3) and the boundedness of w. This implies (2.5).

3. A general central limit theorem. In view of (2.1), (2.2), Lemma 1, and Lemma 2, Theorem 1 is a corollary of Theorem 2.

THEOREM 2. Suppose that  $Z_n^{\theta}$ ,  $n \geq 0$ ,  $\theta \in J$ , is a family of real-valued stochastic processes such that

(3.1) 
$$E(\Delta Z_n^{\theta} | Z_n^{\theta}) = \theta a(n, \theta) Z_n^{\theta} + o(\theta)$$

(3.2) 
$$E((\Delta Z_n^{\theta})^2 | Z_n^{\theta}) = \theta b(n, \theta) + o(\theta)$$

$$E(|\Delta Z_n^{\theta}|^3 | Z_n^{\theta}) = o(\theta),$$

where

$$\sup_{n\geq 0} E(|o(\theta)|)/\theta \to 0$$

as 
$$\theta \to 0$$
,

(3.4) 
$$a(n, \theta) \rightarrow a \quad and \quad b(n, \theta) \rightarrow b$$

as  $\theta \to 0$  and  $n\theta \to \infty$ , and a < 0. Suppose also that

$$\sup_{n\geq 0, \theta\in J} E((Z_n^{\theta})^2) < \infty.$$

Then  $\mathcal{L}(Z_n^{\theta}) \to \mathcal{N}(0, \sigma^2)$  as  $\theta \to 0$  and  $n\theta \to \infty$ , where  $\sigma^2 = b/2|a|$ .

Proof. Let

$$h_n(\gamma) = h_n^{\theta}(\gamma) = E(\exp(i\gamma Z_n)).$$

Then

$$(3.6) h_{n+1}(\gamma) = E(\exp(i\gamma Z_n))E(\exp(i\gamma \Delta Z_n | Z_n)).$$

Expanding  $\exp(i\gamma \Delta Z_n)$  up to terms of third order in  $\gamma$  and using (3.1)—(3.3) we obtain

(3.7) 
$$\Delta h_n(\gamma) = \theta \gamma a(n, \theta) h_n'(\gamma) - \theta 2^{-1} \gamma^2 b(n, \theta) h_n(\gamma) + d_n(\gamma),$$

where

$$(3.8) |d_n(\gamma)| \le \theta \varepsilon_\theta |\gamma|$$

and  $\varepsilon_{\theta}$  is our generic notation for a quantity that depends only on  $\theta$  and approaches 0 as  $\theta$  approaches 0. This estimate is valid for all  $n \ge 0$  as long as  $\gamma$  is bounded,  $|\gamma| \le \Gamma$ . From (3.7) it follows that

(3.9) 
$$\Delta h_n(\gamma) = \theta \gamma a h_n'(\gamma) - \theta 2^{-1} \gamma^2 b h_n(\gamma) + d_n(\gamma) + e_n(\gamma),$$

where, in view of (3.4) and (3.5),

$$(3.10) |e_n(\gamma)| \le \theta c(n, \theta) |\gamma|,$$

and  $c(n, \theta)$  is a quantity that approaches 0 as  $\theta \to 0$  and  $n\theta \to \infty$ . The inequality (3.10) presupposes  $|\gamma| \leq \Gamma$ .

Let

$$v(\gamma) = \exp(2^{-1}\gamma^2\sigma^2)$$
,  
 $H_n(\gamma) = v(\gamma)h_n(\gamma)$ ,  
 $D_n(\gamma) = v(\gamma)d_n(\gamma)$ ,  
 $E_n(\gamma) = v(\gamma)e_n(\gamma)$ ,

and note that

$$v(\gamma)h_n'(\gamma) = H_n'(\gamma) - \sigma^2 \gamma H_n(\gamma) .$$

Thus multiplication of (3.9) by  $v(\gamma)$  yields

(3.11) 
$$\Delta H_n(\gamma) = \theta \gamma a H_n'(\gamma) + D_n(\gamma) + E_n(\gamma).$$

As a consequence of (3.8) and (3.10),

$$|D_n(\gamma)| \le \theta \varepsilon_\theta |\gamma|$$

and

$$(3.13) |E_n(\gamma)| \le \theta c(n, \theta)|\gamma|$$

for  $|\gamma| \leq \Gamma$ .

Let

$$\gamma_j = (1 + \theta a)^j \xi ,$$

where  $\xi$  is fixed for the remainder of the proof. Assuming  $\theta \leq 1/|a|$ ,

$$|\gamma_i| \le e^{a\theta j} |\xi| .$$

In particular,  $\gamma_i$  is bounded by  $|\xi| = \Gamma$  for all j and  $\theta$ .

For any  $0 \le m \le M$ , define  $\mathcal{H}_m = \mathcal{H}_m(M, \theta)$  by

$$\mathcal{H}_m = H_m(\gamma_{M-m})$$
.

Then  $\mathcal{H}_M = H_M(\xi)$ . Suppose that  $\mathcal{H}_M - \mathcal{H}_k \to 0$  as  $\theta \to 0$  and  $k\theta \to \infty$ , while  $\mathcal{H}_k \to 1$  as  $\theta(M-k) \to \infty$ . Then choosing k = [M/2] we see that  $H_M(\xi) \to 1$ ,

$$h_{\scriptscriptstyle M}(\xi) \to \exp\left(-2^{-1}\xi^2\sigma^2\right)$$
,

and  $\mathcal{L}(Z_M) \to \mathcal{N}(0, \sigma^2)$  as  $\theta \to 0$  and  $M\theta \to \infty$ , as the theorem asserts. Thus it remains only to show that  $\mathcal{H}_M = \mathcal{H}_k \to 0$  and  $\mathcal{H}_k \to 1$ .

It may assist the reader in understanding the proof that  $\mathcal{H}_M - \mathcal{H}_k \to 0$  to regard (3.11) as an approximation to the partial differential equation

$$\frac{\partial H(t,\gamma)}{\partial t} = \gamma a \frac{\partial H(t,\gamma)}{\partial \gamma}.$$

For any constant g,  $(t, ge^{-at})$  is a characteristic base curve of this equation ([1] page 63), so

$$\frac{d}{dt} H(t, ge^{-at}) = 0.$$

Since  $\gamma_{M-m}$  approximates  $\gamma_M e^{-am\theta}$ , we expect  $\mathcal{H}_m = H_m(\gamma_{M-m})$  to approximate  $H(m\theta, \gamma_M e^{-am\theta})$ . Thus  $\mathcal{H}_m$  should vary little with m.

Clearly

$$\Delta \mathcal{H}_{m-1} = A_m - B_m$$

for  $m \ge 1$ , where

$$A_m = H_m(\gamma_{M-m}) - H_{m-1}(\gamma_{M-m})$$

and

$$B_m = H_{m-1}(\gamma_{M+1-m}) - H_{m-1}(\gamma_{M-m})$$
.

Thus

$$(3.15) |\mathcal{H}_{M} - \mathcal{H}_{k}| \leq \sum_{m=k+1}^{M} |A_{m} - B_{m}|.$$

Now

(3.16) 
$$B_{m} = \Delta \gamma_{M-m} H'_{m-1}(\gamma_{M-m}) + F_{m-1} \\ = \theta a \gamma_{M-m} H'_{m-1}(\gamma_{M-m}) + F_{m-1},$$

where

(3.17) 
$$|F_{m-1}| \leq 2^{-1} |\Delta \gamma_{M-m}|^2 \max_{|\gamma| \leq |\xi|} |H''_{m-1}(\gamma)| \leq K \theta^2 e^{2a\theta(M-m)}$$

by virtue of (3.14) and (3.5). When the expression (3.16) for  $B_m$  is subtracted from (3.11) for  $A_m$ , the leading terms cancel, so that

$$A_m - B_m = D_{m-1}(\gamma_{M-m}) + E_{m-1}(\gamma_{M-m}) - F_{m-1}.$$

Applying the estimates (3.12), (3.13), (3.14), and (3.17) to (3.15), we obtain

$$\begin{aligned} |\mathscr{H}_{M} - \mathscr{H}_{k}| &\leq (\varepsilon_{\theta} + \sup_{n \geq k} c(n, \theta))\theta \sum_{m=k+1}^{M} e^{a\theta(M-m)} \\ &\leq (\varepsilon_{\theta} + \sup_{n \geq k} c(n, \theta))\theta/(1 - e^{a\theta}). \end{aligned}$$

Since  $\varepsilon_{\theta} \to 0$  as  $\theta \to 0$ , and  $c(n, \theta) \to 0$  as  $\theta \to 0$  and  $n\theta \to \infty$ , it follows that  $\mathscr{H}_M = \mathscr{H}_k \to 0$  as  $\theta \to 0$  and  $k\theta \to \infty$ .

Note, finally, that

$$|h_k(\gamma_{M-k}) - 1| \le |\gamma_{M-k}|E(|Z_k|)$$
  
$$\le K|\gamma_{M-k}|$$

by (3.5). Since  $\gamma_{M-k} \to 0$  as  $\theta(M-k) \to \infty$ , we have  $h_k(\gamma_{M-k}) \to 1$  and thus

$$\mathcal{H}_k = h_k(\gamma_{M-k})v(\gamma_{M-k}) \to 1$$

as  $\theta(M-k) \to \infty$ . This completes the proof.

4. The Wright-Fisher model. Suppose that there are two alleles,  $A_1$  and  $A_2$ , at a certain chromosomal locus in a diploid population of N individuals. Let i be the number and x = i/2N the proportion of  $A_1$  genes in the population at any time. According to the model (see [2] Section 4.8), values  $X_n$  of x in successive generations form a finite Markov chain with transition probabilities

$$p_{ij} = \binom{2N}{j} \pi_i^{j} (1 - \pi_i)^{2N-j}$$
,

where

$$\pi_i = (1 - u)\pi_i^* + v(1 - \pi_i^*)$$

and

$$\pi_i^* = \frac{(1+s_1)x^2 + (1+s_2)x(1-x)}{(1+s_1)x^2 + 2(1+s_2)x(1-x) + (1-x)^2}.$$

The constants  $s_1$ ,  $s_2$ , u, and v control selection pressure and mutation rate. The fitnesses of the genotypes  $A_1 A_1$  and  $A_1 A_2$ , relative to that of  $A_2 A_2$ , are  $1 + s_1$  and  $1 + s_2$ , respectively. The probability that an  $A_1$  gene mutates to  $A_2$  is u, while the probability that  $A_2$  mutates to  $A_1$  is v.

To apply Theorem 1 to this model, we assume that these parameters are proportional to  $\theta = (2N)^{-\frac{1}{2}}$ :  $s_i = \bar{s}_i \theta$ ,  $u = \bar{u}\theta$ , and  $v = \bar{v}\theta$ , where  $\bar{u}, \bar{v} \ge 0$ . The routine verification of the assumptions in the second paragraph of Section 1 is given in ([6] Section 18.1), where it is also shown that s(x) = x(1-x) and

(4.1) 
$$w(x) = \overline{v} - (\overline{u} + \overline{v})x + x(1 - x)(\overline{s}_2 + (\overline{s}_1 - 2\overline{s}_2)x)$$

on I = [0, 1]. Thus Theorem 1 applies whenever w has a unique root  $\lambda$  and  $w'(\lambda) < 0$  (i.e.,  $\lambda$  is stable).

The following conditions are sufficient but by no means necessary for this:  $\bar{u} > 0$ ,  $\bar{v} > 0$ , and  $\bar{s}_1 \le 2\bar{s}_2$ . (Proof. Since  $w(0) = \bar{v} > 0$  and  $w(1) = -\bar{u} < 0$ , w has at least one zero in (0, 1). If  $\bar{s}_1 = 2\bar{s}_2$ , w is quadratic or linear, and uniqueness and stability certainly obtain. If  $\bar{s}_1 < 2\bar{s}_2$ , the coefficient of  $x^3$  is positive, so w has a root above 1 and a root below 0. Thus w has only one root  $\lambda$  in (0, 1) and it must satisfy  $w'(\lambda) < 0$ .) The inequality  $\bar{s}_1 \le 2\bar{s}_2$  admits a number of genetically significant special cases:

- (i) no dominance,  $\bar{s}_1 = 2\bar{s}_2$ ;
- (ii) favored gene completely dominant,  $\bar{s}_1 = \bar{s}_2 > 0$  or  $\bar{s}_1 < \bar{s}_2 = 0$ ; and
- (iii) heterozygote advantage,  $\bar{s}_1 < \bar{s}_2 > 0$ .

Writing  $X_n^N$  and  $x^N$  for  $X_n^\theta$  and  $x_\theta$ , the conclusion of Theorem 1 can be expressed as follows:

$$(2N)^{\frac{1}{2}}[X_n^N - f(n/(2N)^{\frac{1}{2}}, x^N)] \sim \mathcal{N}(0, g(\infty))$$

as  $N \to \infty$  and  $n/N^{\frac{1}{2}} \to \infty$ . The occurrence of the fourth root on the left is noteworthy. (We observe that the related results in lines 13 and 22 on page 259 of [6] should have fourth roots instead of square roots.)

To see the relation of Theorem 1 to other diffusion approximations of the Wright-Fisher model, suppose that the mutation and selection parameters are proportional to a parameter  $\varepsilon > 0$ , i.e.,  $s_i = \bar{s}_i \varepsilon$ ,  $u = \bar{u}\varepsilon$ , and  $v = \bar{v}\varepsilon$ . Theorem 1 pertains directly to  $\varepsilon = (2N)^{-\frac{1}{2}}$ , but it turns out that this result is typical of those obtained when  $\varepsilon \to 0$  sufficiently slowly that  $N\varepsilon \to \infty$ . If the function w given in (4.1) satisfies the hypotheses of Theorem 1, then  $(\varepsilon N)^{\frac{1}{2}}X_n$ , suitably centered, is asymptotically normally distributed as  $\varepsilon \to 0$ ,  $N\varepsilon \to \infty$ , and  $n\varepsilon \to \infty$ . This generalization of Theorem 1 will be proved in a subsequent paper. For a clear heuristic analysis of the asymptotic behavior of  $X_n$  when  $\varepsilon \to 0$  and  $N\varepsilon \to \infty$ , see Section 9 of [3].

Suppose now that  $\varepsilon = (2N)^{-1}$ . In this case,  $\mathscr{L}(X_n^N) \to \mathscr{L}(t, x)$  as  $x^N \to x$ ,  $N \to \infty$ , and  $n\varepsilon \to t < \infty$ , where  $\mathscr{L}(t, x)$  is a nondegenerate distribution associated with a diffusion on I([6] page 260). The standard diffusion approximations of population genetics are of this type ([2] Section 5.1). This result, like the

analogous result (1.5), is valid whether or not the function w in (4.1) satisfies the hypotheses of Theorem 1. One would like to know what auxiliary conditions, if any, must be imposed to insure that  $\mathcal{L}(X_n^N)$  converges to  $\lim_{t\to\infty} \mathcal{P}(t,x)$  as  $x^N\to x$ ,  $N\to\infty$ , and  $n\varepsilon\to\infty$ .

5. Multidimensional case. Suppose that the assumptions of the first two paragraphs of Section 1 are in force, except that  $X_n^{\theta}$  is k dimensional, and the conditional variance in (1.2) is replaced by the conditional covariance matrix. Then (1.5) is valid, where the asymptotic covariance matrix g(t) = g(t, x) satisfies

$$g'(t) = w'(f(t))g(t) + g(t)w'(f(t))^* + s(f(t)),$$

and \* indicates transposition ([6] Theorem 8.1.1). Theorem 3 is the multidimensional analog of Theorem 1.

THEOREM 3. Suppose that the following additional conditions obtain: I is bounded, there is a point  $\lambda$  such that  $w(\lambda) = 0$ , and there is an inner product [x, y] on  $R^k$  such that

$$[x-\lambda, w(x)] < 0$$

for all  $x \in I$ ,  $x \neq \lambda$ , and

$$[z, w'(\lambda)z] < 0$$

for all  $z \in \mathbb{R}^k$ ,  $z \neq 0$ . Then

$$\mathcal{L}(Z_n^{\theta}) \to \mathcal{N}(0, g(\infty))$$

as  $\theta \to 0$  and  $n\theta \to \infty$ , where  $g(\infty)$  is the unique solution of the system

$$w'(\lambda)g(\infty) + g(\infty)w'(\lambda)^* + s(\lambda) = 0$$

of linear equations.

Obviously (5.1) implies that  $\lambda$  is the only zero of w. The most general inner product on  $R^k$  is [x, y] = (x, Py), where (x, y) is the Euclidean inner product and P is a positive definite matrix.

Theorem 3 can be established by a straightforward generalization of the proof of Theorem 1. This involves establishing the multidimensional generalizations of Lemmas 1 and 2 and Theorem 2. We omit details.

Theorem 3 is applicable to the Zeaman-House-Lovejoy learning model [7], which describes how a human or lower animal might learn to attend to a certain "relevant" dimension of a multidimensional stimulus. In this rather complex model,  $X_n$  is two dimensional and I is the closed unit square. There are six learning rate parameters,  $\varphi_1$ ,  $\varphi_2$ ,  $\varphi_3$ ,  $\varphi_4$ ,  $\theta_1$ ,  $\theta_2$ , and two payoff probability parameters,  $\pi_B$  and  $\pi_W$ . To apply Theorem 3, we assume that the learning rate parameters are all proportional to a single parameter  $\theta$ , i.e.,  $\varphi_i = \theta \bar{\varphi}_i$  and  $\theta_j = \theta \bar{\theta}_j$ , where  $\bar{\varphi}_i$  and  $\bar{\theta}_j$  are positive constants. It can be shown that the hypotheses of Theorem 3 are satisfied if and only if one of the following conditions holds: (i)  $\pi_B < 1$  and  $\pi_W < 1$ , or (ii)  $\max(\pi_B, \pi_W) = 1$ ,  $\min(\pi_B, \pi_W) < 1$ , and

 $\bar{\varphi}_1 > \bar{\varphi}_3(\pi_B + \pi_W)/2$ . In either case we can take  $[x, x'] = x_1 x_1' + c x_2 x_2'$  for c sufficiently large. Under condition (i),  $\lambda$  is in the interior of I, while under condition (ii), it is one of the corners.

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