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Finiteness of Equilibria Set for a Nonepistatic Selection under Multilocus Mendel Dominance

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Abstract—It is proven that the equilibria set under multilocus Mendel dominance in a population with any fixed-recombination coefficients is finite generically if the selection is nonepistatic in Karlin's sense. © 2003 Elsevier Science Ltd. All rights reserved.

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In the recent work [1], a general theorem on a finiteness of the equilibria set under phenotypical selection in a multilocus multiallele population with any fixed-recombination coefficients was established. As a consequence, the following result has been proved.

THEOREM 1. (See [1, Corollary 8.2].) *The equilibria set under multilocus Mendel dominance is finite generically.*

In this context, the *genericity* means that the finiteness takes place except for a proper algebraic subset E of the fitness space. The elements of this space are positive vectors of form $\lambda = (\lambda(g, h))$ where g and h run over the set Γ of the gamete genotypes, so that the pairs (g, h) determine the zygote genotypes. The value $\lambda(g, h)$ is called the *fitness coefficient* for the corresponding zygote. This value is symmetric, $\lambda(h, g) = \lambda(g, h)$, moreover, it is invariant with respect to the permutations of homological chromosomes, see [1, Section 6].

Every gamete g can be formally written as the formal product $g_1 \dots g_l$ where g_i is the allele carrying by g at the i^{th} locus, $1 \leq i \leq l$. We identify the set of loci under consideration with $L = \{1, \dots, l\}$. We will say that the multilocus fitness vector is *decomposable* if there is a function P such that

$$\lambda(g, h) = P\left(\lambda^{(1)}(g_1, h_1), \dots, \lambda^{(l)}(g_l, h_l)\right),$$

with some single-locus fitness vectors $\lambda^{(1)}, \dots, \lambda^{(l)}$. This concept includes two important partic-

ular cases: the *additive selection*,

$$\lambda(g, h) = \sum_{i=1}^l \lambda^{(i)}(g_i, h_i),$$

and the *multiplicative selection*

$$\lambda(g, h) = \prod_{i=1}^l \lambda^{(i)}(g_i, h_i),$$

both can be considered as the simplest manifestations of the *nonepistasis* (see [2] for a relevant discussion). Karlin [3] suggested a sort of “interpolation” between additive and multiplicative selection he named the *generalized nonepistatic selection*. This model (in a slightly modified notation) is

$$\lambda(g, h) = \sum_{U \subset L} c(U) \prod_{i \in U} \lambda^{(i)}(g_i, h_i), \tag{1}$$

where $c(U)$ are some nonnegative coefficients, at least one of them is positive. The additive/multiplicative cases are included in (1) with $c(U) = \delta_{U,L}$ or $c(U) = \delta_{|U|,1}$, respectively. (Here, δ is Kronecker’s symbol.)

Now let us clarify what is the Mendel multilocus dominance mentioned in Theorem 1. In a formal sense, this is the direct product of the Mendel single-locus structures, see [1, Section 4]. At the diallele locus with the alleles A and a such that A dominates a , the fitness coefficients are $\lambda(Aa) = \lambda(AA)$ and $\lambda(aa)$. In this situation, we consider $\lambda(AA)$ and $\lambda(aa)$ as independent variables, the coordinates of the two-dimensional fitness vector. A well-known elementary result in this case is that if $\lambda(AA) \neq \lambda(aa)$, then *the number of equilibria is either two or infinity*.

Now we consider two-locus diallele population and suppose that there is the Mendel dominance at each locus. Let the fitness coefficients at the first locus be $\lambda(AA) = \lambda(Aa) = \alpha$, $\lambda(aa) = \varepsilon$, and similarly, let $\lambda(BB) = \lambda(Bb) = \beta$, $\lambda(bb) = \omega$ at the second one. The typical zygotes are $aabb$, $aaBB$, $AAbb$, and $AABB$. Denote their fitness coefficients by $\lambda_1, \lambda_2, \lambda_3, \lambda_4$. All other zygotes have the same fitness coefficients, for example, $\lambda(AaBb) = \lambda(AABB)$, etc.

The selection rule (1) yields

$$\lambda_1 = c_0 + c_1\varepsilon + c_2\omega + c_{12}\varepsilon\omega, \tag{2}$$

$$\lambda_2 = c_0 + c_1\varepsilon + c_2\beta + c_{12}\varepsilon\beta, \tag{3}$$

$$\lambda_3 = c_0 + c_1\alpha + c_2\omega + c_{12}\alpha\omega, \tag{4}$$

$$\lambda_4 = c_0 + c_1\alpha + c_2\beta + c_{12}\alpha\beta, \tag{5}$$

where $c_0 = c(\emptyset)$, $c_1 = c(\{1\})$, $c_2 = c(\{2\})$, $c_{12} = c(L)$, $L = \{1, 2\}$.

By substitution from (2)–(5) into the equations of the exceptional set E (related to Theorem 1), we obtain a system of algebraic equations for $\alpha, \varepsilon, \beta, \omega$. These equations determine the *exceptional set* \tilde{E} in the space of parameters $\alpha, \varepsilon, \beta, \omega$. The equations of \tilde{E} are of form

$$\sum_{i_1, i_2, i_3, i_4} P_{i_1, i_2, i_3, i_4}^{(j)}(c_0, c_1, c_2, c_{12}) \alpha^{i_1} \varepsilon^{i_2} \beta^{i_3} \omega^{i_4} = 0, \tag{6}$$

where $P_{i_1, i_2, i_3, i_4}^{(j)}$ are some polynomials, $1 \leq j \leq N$, N is the total number of them. We show that at least one of polynomials $P_{i_1, i_2, i_3, i_4}^{(j)}$ is not identically zero.

Formulas (2)–(5) yield a linear transformation $(c_0, c_1, c_2, c_{12}) \mapsto (\lambda_1, \lambda_2, \lambda_3, \lambda_4)$. Its determinant is

$$\begin{vmatrix} 1 & \varepsilon & \omega & \varepsilon\omega \\ 1 & \varepsilon & \beta & \varepsilon\beta \\ 1 & \alpha & \omega & \alpha\omega \\ 1 & \alpha & \beta & \alpha\beta \end{vmatrix} = -(\alpha - \varepsilon)^2(\beta - \omega)^2.$$

This is different from zero if $\alpha \neq \varepsilon$ and $\beta \neq \omega$. Without loss of generality, one can assume that this condition is fulfilled. Take $(\lambda_1, \lambda_2, \lambda_3, \lambda_4) \notin E$ and solve the linear system (2)–(5) with respect to c_0, c_1, c_2, c_{12} . (The solution cannot be positive, no matter.) By definition of equations (6), at least one of them turns out to be broken. Thus, there exists a nonvanishing polynomial among $P_{i_1, i_2, i_3, i_4}^{(j)}$.

Now the system of all equations

$$P_{i_1, i_2, i_3, i_4}^{(j)}(c_0, c_1, c_2, c_{12}) = 0,$$

coming from (6) defines a proper algebraic subset $C \subset \mathbf{R}^4$.

THEOREM 2. *If $(c_0, c_1, c_2, c_{12}) \notin C$, then the set of equilibria under selection rule (2)–(5) is finite generically with respect to $\alpha, \varepsilon, \beta, \omega$.*

PROOF. Take any above-mentioned four-tuple (c_0, c_1, c_2, c_{12}) . Then the exceptional set $\tilde{E} \subset \mathbf{R}^4$ defined by (6) is a proper algebraic subset. For any four-tuple $(\alpha, \varepsilon, \beta, \omega) \notin \tilde{E}$, the corresponding four-tuple $(\lambda_1, \lambda_2, \lambda_3, \lambda_4) \notin E$. The latter means, by definition of E , that the equilibria set is finite. ■

The same method works for the generalized nonepistatic selection (1) under the Mendel dominance at each of l diallele loci takes place. The only assertion we need to prove is the following.

LEMMA. *Let for any $i = 1, \dots, l$ some distinct numbers $\lambda_0^{(i)}$ and $\lambda_1^{(i)}$ be given. Then the system of 2^l linear equations*

$$\lambda_{\nu_1, \dots, \nu_l} = \sum_{U \subset L} c(U) \prod_{i \in U} \lambda_{\nu_i}^{(i)} \quad (\nu_i \in \{0, 1\}, 1 \leq i \leq l) \tag{7}$$

is solvable with respect to the unknowns $\{c(U) : U \subset L\}$, whichever are 2^l numbers $\lambda_{\nu_1, \dots, \nu_l}$.

PROOF. Since we have as many equations as there are unknowns in (7), it is sufficient to show that the system with all $\lambda_{\nu_1, \dots, \nu_l} = 0$ has the only trivial solution. This is obvious for $l = 1$ and we actually know this for $l = 2$. Let us pass from $(l - 1)$ to l by induction.

Assume that

$$\sum_{U \subset L} c(U) \prod_{i \in U} \lambda_{\nu_i}^{(i)} = 0 \quad (\nu_i \in \{0, 1\}, 1 \leq i \leq l). \tag{8}$$

It is sufficient to prove that all $c(U) = 0$ for $U \neq \emptyset$. For definiteness, we will prove that $c(U) = 0$ if $1 \in U$. Consider two subsystems of (8) which correspond to $\nu_1 = 0$ and $\nu_1 = 1$, respectively:

$$\sum_{1 \in U \subset L} c(U) \lambda_0^{(1)} \prod_{i \in U \setminus \{1\}} \lambda_{\nu_i}^{(i)} + \sum_{1 \notin U \subset L} c(U) \prod_{i \in U} \lambda_{\nu_i}^{(i)} = 0 \tag{9}$$

and

$$\prod_{1 \in U \subset L} c(U) \lambda_1^{(1)} \prod_{i \in U \setminus \{1\}} \lambda_{\nu_i}^{(i)} + \sum_{1 \notin U \subset L} c(U) \prod_{i \in U} \lambda_{\nu_i}^{(i)} = 0. \tag{10}$$

Subtracting (10) from (9) and cancelling $\lambda_0^{(1)} - \lambda_1^{(1)} \neq 0$, we get

$$\sum_{1 \in U \subset L} c(U) \prod_{i \in U \setminus \{1\}} \lambda_{\nu_i}^{(i)} = 0.$$

Thus, the system is of form (7) with $L \setminus \{1\}$ instead of L . By the induction, assumption $c(U) = 0$ for $1 \in U$. ■

As a result, we obtain the following theorem.

THEOREM 3. *In the l -locus diallele population with the Mendel dominance at each locus, there exists a proper algebraic subset \mathcal{CR}^{2^l} such that if $(c(U) : U \subset L) \notin \mathcal{C}$, then under generalized nonepistatic selection rule (1), the set of equilibria is finite generically with respect to the single-locus fitness coefficients.*

In addition, *the number of equilibria does not exceed $3^{|\Gamma|-1}$* , see [1, Corollary 8.1].

Note that Theorem 3 says nothing for *a priori* given selective weights $c(U)$ in (1), in particular, for the additive or the multiplicative selection. However, in the additive case, the finiteness result follows from [4, Theorem 9.6.13 and Corollary 9.13]. The generic finiteness of the equilibria set under the multiplicative selection has been recently proved in [5].

REFERENCES

1. Y. Lyubich, V. Kirzhner and A. Ryndin, Mathematical theory of phenotypical selection, *Advanced in Appl. Math.* **26**, 330–352 (2001).
2. P.A.P. Moran, Unsolved problems in evolutionary theory, In *Proc. 5th Berkeley Symp. on Math. Stat. Prob.*, California, (1967).
3. S. Karlin, Principles of polymorphism and epistasis for multilocus systems, *Proc. Nat. Acad. Sci. USA* **76**, 541–545 (1979).
4. Y. Lyubich, *Mathematical Structures in Population Genetics, Biomathematics, Volume 22*, Springer, Berlin, (1992).
5. V. Kirzhner and Y. Lyubich, On the finiteness of multiplicative selection equilibria, (Preprint) (2001).