

Selection-mutation balance models with epistatic selection

Article

Accepted Version

Kondratiev, Y. G., Kuna, T. and Ohlerich, N. (2008) Selection-mutation balance models with epistatic selection. *Condensed Matter Physics*, 11 (2). pp. 283-291. ISSN 1607-324X
Available at <https://centaur.reading.ac.uk/1397/>

It is advisable to refer to the publisher's version if you intend to cite from the work. See [Guidance on citing](#).

Published version at: <http://www.icmp.lviv.ua/journal/zbirnyk.54/index.html>

Publisher: Institute for Condensed Matter Physics

All outputs in CentAUR are protected by Intellectual Property Rights law, including copyright law. Copyright and IPR is retained by the creators or other copyright holders. Terms and conditions for use of this material are defined in the [End User Agreement](#).

www.reading.ac.uk/centaur

CentAUR

Central Archive at the University of Reading

Reading's research outputs online

Selection-mutation balance models with epistatic selection

2 Yu.G.Kondratiev, T.Kuna, N.Ohlerich

Universität Bielefeld, Postfach 10 01 31, D–33501 Bielefeld, Germany; BiBoS, Univ. Bielefeld, Germany

Received January 31, 2208

3 We present an application of birth-and-death processes on configuration spaces to a generalized mutation-
4 selection balance model. The model describes the aging of population as a process of accumulation of muta-
5 tions in a genotype. A rigorous treatment demands that mutations correspond to points in abstract spaces.
6 Our model describes an infinite-population, infinite-sites model in continuum. The dynamical equation which
7 describes the system, is of Kimura-Maruyama type. The problem can be posed in terms of evolution of states
8 (differential equation) or, equivalently, represented in terms of Feynman-Kac formula. The questions of interest
9 are the existence of a solution, its asymptotic behavior, and properties of the limiting state. In the non-epistatic
10 case the problem was posed and solved in [Steinsaltz D., Evans S.N., Wachter K.W., Adv. Appl. Math., 2005,
11 35(1)]. In our model we consider a topological space X as the space of positions of mutations and the influ-
12 ence of an epistatic potential.

13 **Key words:**

14 **PACS:** 02.50.Ga

15 1. The model

16 First recall some genetical concepts and notions, see e.g. [1]. A gene represents a (contiguous)
17 region of DNA coding. It may have different forms, called alleles. Thus an allele is one of the
18 variant forms of a gene that occupies a given locus (position) on a chromosome, i.e. alleles are
19 DNA sequences that code a gene. An individual's genotype for a certain gene is the collection
20 of alleles it consists of. A change of genetic material is called a mutation, and the affected allele
21 is called mutant allele. We call the “null genotype” the one which has wild-type alleles at every
22 locus and carries none of mutant alleles. So a wild-type allele is an allele which is considered to
23 be “normal” for the organism in question, as opposed to a mutant allele which appears due to
24 mutation. In this chapter we will use the word “genotype” in a sense somewhat different from the
25 mentioned above: a genotype represents a set of mutant alleles that an individual may carry. So in
26 contrast to the usual definition we are interested only in the set of mutant alleles, rather than in
27 the complete information about all alleles.

28 In this section we describe a model introduced by [8], which describes the aging of a population.
29 Let X be a Polish space, interpreted as the space of loci (i.e. positions of possible mutations). Denote
30 the Borel σ -algebra on X by $\mathcal{B}(X)$, and fix a Borel σ -finite measure σ on $(X, \mathcal{B}(X))$ – interpreted
31 as mutation rate. For simplicity, we assume that at each locus at most one mutation may occur.
32 A locally finite configuration of points in X (defined as usual) is interpreted as a genotype. Then
33 $\gamma = \emptyset$ plays the role of the null genotype (wild-type genotype). The set of all genotypes γ is thus
34 the configuration space $\Gamma := \Gamma(X)$. We assume that genotypes are influenced by a selection cost
35 Φ , which is a continuous function $\Phi : \Gamma \rightarrow \mathbb{R}$, e.g. $\Phi(\emptyset) = 0$, $\Phi(\gamma) > 0$, for $\gamma \neq \emptyset$.

The emergence of mutant alleles is described by a stochastic process, the state of the population
of genotypes at each fixed moment of time t is described by a probability measure μ_t on Γ . The
time development of the population is modelled by a Kimura-Maruyama type equation

$$\frac{d}{dt} \mu_t(F) = \mu_t \left(\int_X (F(\cdot \cup x) - F(\cdot)) d\sigma(x) \right) - \mu_t(F \cdot \Phi) + \mu_t(F) \mu_t(\Phi), \quad (1)$$

where $\mu_t(F) := \int_{\Gamma} F d\mu_t$, $F : \Gamma \rightarrow \mathbb{R}$ is a bounded cylindric function. The questions of interest for us are: the existence of solution μ_t , convergence of $\mu_t \rightarrow \mu$ for $t \rightarrow +\infty$ and properties of the obtained limiting state μ . The useful choice of time parameterization is to start the process in the remote past, namely at time $t = -T < 0$, in the state μ_{-T} . Then we arrive at $t = 0$ in the state $\mu_{0,T}$. The limiting state for long time is then given by

$$\lim_{T \rightarrow +\infty} \mu_{0,T} = \mu_0.$$

Next, we consider another representation of the model, which gives us an explicit solution of equation (1) with the help of the Feynman-Kac formula. Denote by $E := \Gamma(X)$. Remind that E is a Polish space. Let L be a Markov generator defined by

$$LF(\gamma) = \int_X (F(\gamma \cup x) - F(\gamma)) d\sigma(x)$$

for bounded cylindric functions $F : E \rightarrow \mathbb{R}$. The continuous function $\Phi : E \rightarrow \mathbb{R}$ will play the role of potential in Feynman-Kac formula. Rewriting (1) in terms of these notations we obtain

$$\frac{d}{dt} \mu_t^T(F) = \mu_t^T(LF) - \mu_t^T(F \cdot \Phi) + \mu_t^T(F) \mu_t^T(\Phi). \quad (2)$$

Denote by μ_t^T , $-T \leq t \leq 0$ the measure-valued dynamical system which is the solution of (2) for each bounded cylindric function $F : E \rightarrow \mathbb{R}$, started in $\mu_{-T}^T = \mu$.

The solution μ_t^T of (2) can be explicitly written as

$$\mu_t^T = \frac{1}{Z_t(\Phi)} e^{(t+T)(L-\Phi)^*} \mu,$$

where Z_t is the normalizing constant. Via Feynmann-Kac formula we can represent μ_t^T as

$$\mu_t^T(f) = \frac{\mathbb{E} \left[f(\xi_t^T) e^{-\int_{-T}^t \Phi(\xi_\tau^T) d\tau} \right]}{\mathbb{E} \left[1 \cdot e^{-\int_{-T}^t \Phi(\xi_\tau^T) d\tau} \right]},$$

where ξ_τ^T denotes the Markov process corresponding to the generator L , started in $\mu_{-T}^T = \mu$. Performing the limit $T \rightarrow +\infty$ gives us heuristically

$$\mu_0(f) = \int_{\Omega(\mathbb{R} \rightarrow E)} f(\xi(0)) d\nu^\Phi(\xi(\cdot)), \quad (3)$$

where

$$d\nu^\Phi(\xi(\cdot)) = \frac{1}{Z} e^{-\int_{-\infty}^0 \Phi(\xi(\tau)) d\tau} d\nu^0(\xi(\cdot)), \quad (4)$$

Z is the normalizing constant.

The aim of the following sections is to give proper sense to ν^Φ , defining the measure first in a bounded volume and for finite time and then going to the limit. By means of ν^Φ we derive the large time asymptotic for μ_0^T .

In the non-epistatic case the problem was posed and solved in [8]. The articles [3], [4] were motivated by this work, and treat the case of a more general potential – the epistatic one. In both articles the space of the possible positions of mutations is \mathbb{R}^d . The generalization to a topological space X seems important because of the nonlinear structure of the DNA. In our model we consider a topological space X as the space of positions of mutations and the influence of an epistatic potential.

2. Pure birth process

We define the pure birth Markov process on $\Gamma(X)$, starting with an empty configuration at time $t = -T$, via the generator

$$L_B F(\gamma) = \int_X (F(\gamma \cup y) - F(\gamma)) \sigma(dy) \quad (5)$$

for bounded cylinder functions $F(\gamma)$. In our interpretation this means that there were no mutant alleles at the beginning, in other words we start with the null genotype. As the time passes, the mutations gradually appear in some points $x_i \in X$ at times t_i , $-T < t_i \leq 0$, and then they stay forever.

Notational convention: for readability reasons we prefer to consider the following positive times. Nevertheless, we would like to consider 0 as the final time. Therefore, we reflect the time w.r.t. to the origin. So we consider our pure birth process on the space of marked configurations $\hat{\Gamma}(X, \mathbb{R}_+)$, which is defined by

$$\hat{\Gamma}(X, \mathbb{R}_+) = \{\hat{\gamma} = (\gamma, s(\gamma)) \mid \gamma \in \Gamma(X), s(\gamma) = \{s_x \mid x \in \gamma\}, s_x \in \mathbb{R}_+\}.$$

For more details about marked configuration spaces see cf. [1,6,7]. The spaces $\hat{\Gamma}(\Lambda, \mathbb{R}_+)$ and $\hat{\Gamma}(\Lambda, [0, T])$ are defined analogously. Denote the marked Poisson measure on $\hat{\Gamma}(X, [0, T])$ by ν_T^0 , and its restriction to $\hat{\Gamma}(\Lambda, [0, T])$ by $\nu_{\Lambda, T}^0$. It is well known that the marked Poisson measure ν_T^0 can be characterized by its Laplace transform

$$\int_{\hat{\Gamma}(X, [0, T])} e^{\langle f, \hat{\gamma} \rangle} d\nu_T^0(\hat{\gamma}) = \exp \left\{ \int_X \int_0^T (e^{f(x, t)} - 1) dt d\sigma(x) \right\}, \quad f \in C_0(X \times [0, T]). \quad (6)$$

The Markov birth process $\xi_\tau(\hat{\gamma}), 0 \leq \tau \leq T$ (time is going backwards, i.e. the process starts at T and ends at 0) on $(\hat{\Gamma}(X, [0, T]), \nu_T^0)$, corresponding to the generator (5) can be realized by

$$\xi_\tau : \hat{\Gamma}(X, [0, T]) \rightarrow \Gamma(X), \quad \xi_\tau(\hat{\gamma}) = \{x \in \gamma \mid \tau \leq s_x(\gamma)\}. \quad (7)$$

Further, we assume the effect of a selection cost function $\Phi : \Gamma \rightarrow \mathbb{R}_+$, which consists of two parts:

$$\Phi(\gamma) = \Phi_{ne}(\gamma) + \Phi_e(\gamma).$$

$\Phi_{ne}(\gamma)$ is the nonepistatic part, which describes the life costs of a mutation, and it is given by

$$\Phi_{ne}(\gamma) := \langle h, \gamma \rangle = \sum_{x \in \gamma} h(x), \quad h(x) \geq c > 0.$$

$\Phi_e(\gamma)$ is the epistatic part, which describes the coexistence costs of mutations, and it is defined by

$$\Phi_e(\gamma) := \sum_{\{x, y\} \subset \gamma} \phi(x; y),$$

conditions on ϕ are specified later.

As the configuration γ may contain, in general, infinite number of points, the above cost functions are well-defined only in a bounded region $\Lambda \subset X$.

For convenience we introduce the corresponding path space measure in two steps: first we consider only the effect of the nonepistatic part of the cost function and then take into consideration the effect of the epistatic part.

2.1. The effect of nonepistatic part of the potential

First we construct the path space measure ν^h on the space $\hat{\Gamma}(X, \mathbb{R}_+)$, obtained under the effect of Φ_{ne} . The restriction of ν^h to $\hat{\Gamma}(\Lambda, [0, T])$ is denoted by $\nu_{\Lambda, T}^h$, and defined for bounded $\Lambda \subset X$ as

$$d\nu_{\Lambda, T}^h(\hat{\gamma}_\Lambda) = \frac{1}{Z_{\Lambda, T}} \exp \left\{ - \int_0^T \Phi_{ne}^{T, \Lambda}(\xi_\tau(\hat{\gamma}_\Lambda)) d\tau \right\} d\nu_{\Lambda, T}^0(\hat{\gamma}_\Lambda), \quad (8)$$

where $Z_{\Lambda, T}$ is the normalizing constant

$$Z_{\Lambda, T} = \int_{\hat{\Gamma}(\Lambda, [0, T])} \exp \left\{ - \int_0^T \Phi_{ne}^{T, \Lambda}(\xi_\tau(\hat{\gamma}_\Lambda)) d\tau \right\} d\nu_{\Lambda, T}^0(\hat{\gamma}_\Lambda). \quad (9)$$

Then we obtain the measure ν^h as a limit of measures $\nu_{\Lambda, T}^h$, which are defined in a bounded volume Λ and for finite time T . $\nu_{\Lambda, T}^h$ is the so-called Gibbs perturbation of marked Poisson measure ν_T^0 plus.

First we will show that $\nu_{\Lambda, T}^h$ still remains a Poisson measure. For this purpose we calculate its intensity measure by computing the Laplace transform of $\nu_{\Lambda, T}^h$.

Lemma 2.1. *Let $F(\hat{\gamma}) = e^{\langle f, \hat{\gamma} \rangle}$, $f \in C_0(X \times [-T, 0])$ where*

$$\langle f, \hat{\gamma} \rangle := \sum_{(x, t_x) \in \hat{\gamma}} f(x, t_x) = \int_0^T \int_X f(x, s) \hat{\gamma}(dx, ds), \quad \hat{\gamma} \in \hat{\Gamma}(X, \mathbb{R}_+).$$

Then we have

$$\begin{aligned} & \int_{\hat{\Gamma}(\Lambda, [0, T])} F(\hat{\gamma}_\Lambda) \exp \left\{ - \int_0^T \Phi_{ne}^{T, \Lambda}(\xi_\tau(\hat{\gamma}_\Lambda)) d\tau \right\} d\nu_{\Lambda, T}^0(\hat{\gamma}_\Lambda) \\ &= \exp \left\{ \int_\Lambda \int_0^T (\exp \{ f(x, s) - sh(x) \} - 1) ds d\sigma(x) \right\}. \end{aligned}$$

Then the normalizing constant $Z_{\Lambda, T}$ is

$$Z_{\Lambda, T} = \exp \left\{ \int_\Lambda \int_0^T (\exp \{-sh(x)\} - 1) ds d\sigma(x) \right\}. \quad (10)$$

Calculating the integral of $F = e^{\langle f, \hat{\gamma} \rangle}$ w.r.t the measure $\nu_{\Lambda, T}^h$ we obtain

$$\begin{aligned} \int F(\hat{\gamma}_\Lambda) d\nu_{\Lambda, T}^h(\hat{\gamma}_\Lambda) &= \frac{\exp \left\{ \int_\Lambda \int_0^T (\exp \{ f(x, s) - sh(x) \} - 1) ds d\sigma(x) \right\}}{\exp \left\{ \int_\Lambda \int_0^T (\exp \{-sh(x)\} - 1) ds d\sigma(x) \right\}} \\ &= \exp \left\{ \int_\Lambda \int_0^T (e^{f(x, s)} - 1) e^{-sh(x)} ds d\sigma(x) \right\}. \end{aligned}$$

Thus $\nu_{\Lambda, T}^h$ is a marked Poisson measure on $\hat{\Gamma}(\Lambda, [0, T])$ with intensity measure $e^{-sh(x)} d\sigma(x) ds$.

Recall that we say that there exists a weak limit $\lim_{\Lambda \uparrow X} \rho_\Lambda = \rho$ if

$$\int F(\hat{\gamma}) d\rho_\Lambda(\hat{\gamma}) \xrightarrow{\Lambda \nearrow X} \int F(\hat{\gamma}) d\rho(\hat{\gamma}).$$

for all bounded cylinder functions $F \in \mathcal{FL}^0(\hat{\Gamma}(X, [0, T]))$. The set of cylinder functions $\mathcal{FL}^0(\hat{\Gamma}(X, [0, T]))$ is defined as the set of all measurable F such that there exists a $\Lambda \in \mathcal{B}_c(X)$ with

$$F(\hat{\gamma}) = F(\hat{\gamma} \upharpoonright_{\Lambda \times [0, T]}).$$

- 1 We are interested in the weak limit of $\nu_{\Lambda,T}^h$ for $\Lambda \uparrow X$, $T \rightarrow +\infty$. In the case considered here the
 2 limit does not depend on the order in which the limits are taken. First we can take, for example,
 3 $\Lambda \uparrow X$, then $T \rightarrow +\infty$. As a result we get the following statement:

Theorem 2.2. 1) *There exists a weak limit*

$$\lim_{\Lambda \uparrow X} \nu_{\Lambda,T}^h = \nu_T^h,$$

- 4 where ν_T^h is a marked Poisson measure on $\hat{\Gamma}(X, [0, T])$ with intensity measure $e^{-sh(x)}\sigma(dx)ds$.
 2) *There exists a weak limit*

$$\lim_{T \rightarrow +\infty} \nu_T^h = \nu^h,$$

- 5 where ν^h is a marked Poisson measure on $\hat{\Gamma}(X, \mathbb{R}_+)$ with the same intensity measure $e^{-sh(x)}\sigma(dx)ds$.
 6 Ultimately, the measure ν^h can also be described as a marked point field $\hat{\gamma} = (\gamma, s_\gamma)$, where γ is
 7 distributed according to $\pi_{\sigma/h(x)}$ - Poisson measure on $\Gamma(X)$ - with marks $s_x \in \mathbb{R}_+$ distributed
 8 independently with probability $p(ds) = h(x)e^{-h(x)s}ds$ on \mathbb{R}_+ .

The main object of our interest is the final distribution of mutations μ^h , i.e. it is the distribution of end points of the bars. Recall that we have chosen the time range so that the final time is 0. We obtain μ^h , similar to the construction above, as the limit of final distributions $\mu_{\Lambda,T}^0$ given in the bounded volume and for finite time. The measure $\mu_{\Lambda,T}^0$ on $\Gamma(X)$ is defined for $F(\eta) = e^{\langle f, \eta \rangle}$, $\eta \in \Gamma(X)$ by

$$\begin{aligned} \int_{\Gamma(X)} F(\gamma_\Lambda) d\mu_{\Lambda,T}^0(\gamma_\Lambda) &:= \int_{\hat{\Gamma}(\Lambda, [0, T])} F(\xi_0(\hat{\gamma}_\Lambda)) d\nu_{\Lambda,T}^h(\hat{\gamma}_\Lambda) \\ &= \frac{\int F(\xi_0(\hat{\gamma}_\Lambda)) \exp\left\{-\int_0^T \Phi_{ne}^{T,\Lambda}(\xi_t(\hat{\gamma}_\Lambda)) dt\right\} d\nu_{\Lambda,T}^0(\hat{\gamma}_\Lambda)}{\int \exp\left\{-\int_0^T \Phi_{ne}^{T,\Lambda}(\xi_t(\hat{\gamma}_\Lambda)) dt\right\} d\nu_{\Lambda,T}^0(\hat{\gamma}_\Lambda)}. \end{aligned} \quad (11)$$

By definition of $\mu_{\Lambda,T}^h$ the integral w.r.t. $\mu_{\Lambda,T}^h$ is given by

$$\int_{\hat{\Gamma}(\Lambda, [0, T])} F(\hat{\gamma}) d\mu_{\Lambda,T}^h(\hat{\gamma}) = \frac{\int_{\hat{\Gamma}(\Lambda, [0, T])} e^{T(L_\Lambda - \Phi_{ne}^{T,\Lambda})} F(\hat{\gamma}) d\mu_{\Lambda,T}^0(\hat{\gamma})}{\int_{\hat{\Gamma}(\Lambda, [0, T])} e^{T(L_\Lambda - \Phi_{ne}^{T,\Lambda})} 1 d\mu_{\Lambda,T}^0(\hat{\gamma})}, \quad (12)$$

- 9 so $\mu_{\Lambda,T}^0$ is the solution of (1) in the bounded volume Λ for finite time T , where $\Phi(\gamma) := \langle h, \gamma \rangle$.
 Note that for $f \in C_0(X)$, $\hat{\gamma} \in \hat{\Gamma}(X, \mathbb{R}_+)$ we have

$$\langle f, \xi_0(\hat{\gamma}) \rangle = \int_X \int_0^T f(x) \hat{\gamma}(dx, ds) = \langle F, \hat{\gamma} \rangle,$$

- 10 where $F(x, s) = f(x) \mathbb{1}_{[0, T]}(s)$. Therefore, the following lemma is the corollary of Lemma 2.1.

Lemma 2.3. *Let $F(\eta) = e^{\langle f, \eta \rangle}$, where $\eta \in \Gamma(X)$, $f \in C_0(X)$. Then*

$$\begin{aligned} &\int F(\xi_0(\hat{\gamma}_\Lambda)) \exp\left\{-\int_0^T \Phi_{ne}^{T,\Lambda}(\xi_t(\hat{\gamma}_\Lambda)) dt\right\} d\nu_{\Lambda,T}^0(\hat{\gamma}_\Lambda) \\ &= \exp\left\{\int_\Lambda \int_0^T (\exp\{f(x) - sh(x)\} - 1) ds d\sigma(x)\right\}. \end{aligned}$$

Now we calculate the integral in (11)

$$\begin{aligned} \int F(\gamma_\Lambda) d\mu_{\Lambda,T}^0(\gamma_\Lambda) &= \frac{\exp \left\{ \int_\Lambda \int_0^T (\exp \{f(x) - sh(x)\} - 1) ds d\sigma(x) \right\}}{\exp \left\{ \int_\Lambda \int_0^T (\exp \{-sh(x)\} - 1) ds d\sigma(x) \right\}} \\ &= \exp \left\{ \int_\Lambda (e^{f(x)} - 1) \frac{(1 - \exp \{-Th(x)\})}{h(x)} d\sigma(x) \right\}. \end{aligned}$$

Again, as before, we are interested in the weak limit of $\mu_{\Lambda,T}^0$ for $\Lambda \uparrow X$, $T \rightarrow +\infty$. The limit does not depend on the order as well. We can first take for example $\Lambda \uparrow X$, then $T \rightarrow +\infty$. As a result, we get the following statement:

Theorem 2.4. (cf. [8])

1) There exists a weak limit

$$\lim_{\Lambda \uparrow X} \mu_{\Lambda,T}^0 = \mu_T^h,$$

where μ_T^h is a Poisson measure on $\Gamma(X)$ with intensity

$$\frac{(1 - \exp \{-Th(x)\})}{h(x)} d\sigma(x).$$

The weak limit means that for all bounded cylinder functions $F \in \mathcal{FL}^0(\Gamma(X))$

$$\int F(\hat{\gamma}) d\mu_{\Lambda,T}^0(\hat{\gamma}) \xrightarrow{\Lambda \nearrow X} \int F(\hat{\gamma}) d\mu_T^h(\hat{\gamma}).$$

2) According to Lebesgues dominated convergence theorem, there exists a weak limit

$$\lim_{T \rightarrow +\infty} \mu_T^h = \mu^h,$$

where μ^h is a Poisson measure on $\Gamma(X)$ with intensity measure $\frac{1}{h(x)} \sigma$.

2.2. The effect of the epistatic part of the potential

Now we include the effect of the epistatic part of the potential $\Phi_e(\gamma)$. We consider the Gibbs perturbation of measure ν^h from Theorem 2.2 through Φ_e , i.e.

$$d\nu^{\beta,\phi}(\hat{\gamma}) = \frac{1}{Z_\beta} \exp \left\{ -\beta \int_0^{+\infty} \Phi_e(\xi_\tau(\hat{\gamma})) d\tau \right\} d\nu^h(\hat{\gamma}), \quad \beta > 0.$$

Again the construction is well-defined only for a bounded region $\Lambda \subset X$ and we first consider the restriction of measures to the space $\hat{\Gamma}(\Lambda, \mathbb{R}_+)$:

$$d\nu_\Lambda^{\beta,\phi}(\hat{\gamma}_\Lambda) = \frac{1}{Z_{\beta,\Lambda}} \exp \left\{ -\beta \int_0^{+\infty} \Phi_e^\Lambda(\xi_\tau(\hat{\gamma}_\Lambda)) d\tau \right\} d\nu_\Lambda^h(\hat{\gamma}_\Lambda). \quad (13)$$

We define the measure $\nu^{\beta,\phi}$ as the weak limit of $\nu_\Lambda^{\beta,\phi}$. The main technique is based on cluster expansion method cf. [6,5]. Note that

$$\int_0^{+\infty} \Phi_e(\xi_\tau(\hat{\gamma})) d\tau = \sum_{\{x,y\} \subseteq \gamma} \phi(x;y) \min(s_x, s_y), \quad \hat{\gamma} = (\gamma, s(\gamma)).$$

To use the appropriate convergence theorems we have to make some assumptions on ϕ and ψ , where

$$\psi(\hat{x}, \hat{y}) := \phi(x;y) \min(s_x, s_y), \quad \hat{x} = (x, s_x), \quad \hat{y} = (y, s_y).$$

(S) Stability of ϕ : $\exists B \geq 0$ such that $\forall \gamma \in \Gamma_0(X)$

$$\sum_{\{x,y\} \subseteq \gamma} \phi(x;y) \geq -B|\gamma|. \quad (14)$$

(I $_\psi$) Integrability of ψ

$$C(\beta, h) := \operatorname{esssup}_{y \in X, t \in \mathbb{R}_+} \int_X \int_0^{+\infty} |e^{-\beta\psi((x,s),(y,t))} - 1| e^{2\beta Bs - hs} ds \sigma(dx) < \infty. \quad (15)$$

1 Consequence of stability assumption:

Lemma 2.5. *Let ϕ fulfill (S). Then $\forall \hat{\gamma} = (\gamma, s(\gamma)) \in \hat{\Gamma}_0(X, \mathbb{R}_+)$*

$$\sum_{\{x,y\} \subseteq \gamma} \phi(x;y) \min(s_x, s_y) \geq -B \sum_{x \in \gamma} s_x(\gamma). \quad (16)$$

and $\forall \hat{\gamma} = (\gamma, s(\gamma)) \in \hat{\Gamma}_0(X, \mathbb{R}_+)$ there exists $x_0 \in \gamma$ such that

$$\sum_{x \in \gamma \setminus \{x_0\}} \phi(x; x_0) \min(s_x, s_{x_0}) \geq -2Bs_{x_0}. \quad (17)$$

2 2.3. Cluster expansion

By the definition of $d\nu_\Lambda^{\beta, \phi}$

$$d\nu_\Lambda^{\beta, \phi}(\hat{\gamma}_\Lambda) = \frac{1}{Z_{\beta, \Lambda}} \exp \left\{ -\beta \sum_{\{\hat{x}, \hat{y}\} \subseteq \hat{\gamma}_\Lambda} \psi(\hat{x}; \hat{y}) \right\} d\nu_\Lambda^h(\hat{\gamma}_\Lambda). \quad (18)$$

Denote by $\hat{\sigma}(dx, ds) = e^{-sh(x)} \sigma(dx) ds$. Theorem 2.2 says that ν_Λ^h is the Poisson measure on $\hat{\Gamma}(\Lambda, \mathbb{R}_+)$ with intensity $\hat{\sigma}(dx, ds)$. By the definition of Poisson and the Lebesgue-Poisson measure

$$d\nu_\Lambda^h = \exp\{-\hat{\sigma}(\Lambda \times [0, +\infty))\} d\lambda_{\hat{\sigma}}.$$

Then (18) can be written as

$$d\mu_\Lambda^{\beta, \phi}(\hat{\gamma}_\Lambda) = \frac{1}{\hat{Z}_{\beta, \Lambda}} \exp \left\{ -\beta \sum_{\{\hat{x}, \hat{y}\} \subseteq \hat{\gamma}_\Lambda} \psi(\hat{x}; \hat{y}) \right\} d\lambda_{\hat{\sigma}}(\gamma_\Lambda),$$

3 where $\hat{Z}_{\beta, \Lambda} = Z_{\beta, \Lambda} \cdot \exp\{\hat{\sigma}(\Lambda \times [0, +\infty))\}$.

Cluster expansion is a tool which is used to effectively estimate the Gibbs factor $e^{-\beta E(\gamma)}$ for small parameters, see e.g. [7]. Here we follow the presentation given in [5,6]. There the cluster expansion was generalized to a general metric space, i.e. no translation invariant structure is present. In our case the factor which we are going to decompose is

$$p_{\Lambda, \beta}(\hat{\gamma}_\Lambda) := \exp \left\{ -\beta \sum_{\{\hat{x}, \hat{y}\} \subseteq \hat{\gamma}_\Lambda} \psi(\hat{x}; \hat{y}) \right\}. \quad (19)$$

From [5,6] we know that the cluster decomposition of (19) is as follows:

$$p_{\Lambda, \beta}(\hat{\gamma}_\Lambda) = \sum_{(\hat{\gamma}_1, \hat{\gamma}_2, \dots, \hat{\gamma}_m)}^{(\hat{\gamma}_\Lambda)} k(\hat{\gamma}_1) k(\hat{\gamma}_2) \dots k(\hat{\gamma}_m),$$

$p_{\Lambda, \beta}(\emptyset) = 1$. Here $\sum_{(\hat{\gamma}_1, \hat{\gamma}_2, \dots, \hat{\gamma}_m)}^{(\hat{\gamma}_\Lambda)}$ means the summation over all partitions of the configuration $\hat{\gamma}_\Lambda$ into non-empty subconfigurations $\hat{\gamma}_i \subseteq \hat{\gamma}_\Lambda$, i.e. over all non-ordered sets $\{\hat{\gamma}_1, \hat{\gamma}_2, \dots, \hat{\gamma}_m\}$, $m = 1, 2, \dots, |\hat{\gamma}_\Lambda|$ of subconfigurations of $\hat{\gamma}_\Lambda$ with pair-disjoint supports $\gamma_i \subseteq \gamma_\Lambda$ such that $\cup_{i=1}^m \gamma_i = \gamma_\Lambda$. The values of $k(\hat{\gamma})$ are given for a finite non-empty configuration $\hat{\gamma}$ by

$$k(\hat{\gamma}) = \sum_{G \in \mathcal{G}(\hat{\gamma})} \prod_{\{x, y\} \in G} (e^{-\beta\phi(x; y) \min\{s_x, s_y\}} - 1),$$

$k(\hat{\gamma}) = 1$ if $|\hat{\gamma}| = 1$. By $\mathcal{G}(\hat{\gamma})$ we denote the set of all connected graphs with the set of vertices γ , and the product $\prod_{\{x, y\} \in G}$ is taken over all edges of the graph G . For $\gamma \in \Omega_0 \setminus \emptyset$, $\hat{x} = (x, s_x) \in \hat{\gamma}$ define \bar{k} as

$$\bar{k}(\{\hat{x}\}, \hat{\gamma} \setminus \{\hat{x}\}) := k(\hat{\gamma}). \quad (20)$$

The general idea of cluster expansion is to find a function Q dominating \bar{k} . One can show that

$$Q(\hat{\gamma}, \hat{\zeta}) = \prod_{y \in \gamma \cup \zeta} \exp\{2\beta B s_y\} \sum_{T \in \mathcal{T}(\hat{\gamma} \cup \hat{\zeta})} \prod_{\{y, y'\} \in T} |e^{-\beta\phi(y, y') \min(s_y, s_{y'})} - 1| \quad (21)$$

gives such an upper bound. Using the function Q we prove the following fact.

Theorem 2.6. *Let $\Lambda \in \mathcal{B}_c(\hat{\Gamma}(X, \mathbb{R}_+))$ be given. Then for any parameters β and h such that $2\beta B - h < 0$ and*

$$C(\beta, h) < \frac{1}{2e}, \quad (22)$$

where $C(\beta, h)$ is given by the integrability condition (15), we have

$$\int_{\hat{\Gamma}(\Lambda, \mathbb{R}_+) \setminus \{\emptyset\}} \int_{\hat{\Gamma}_0(X, \mathbb{R}_+)} |k(\hat{\gamma} \cup \hat{\eta})| \lambda_{\hat{\sigma}}(d\hat{\gamma}) \lambda_{\hat{\sigma}}(d\hat{\eta}) < \infty. \quad (23)$$

From this theorem there follows our main result, similarly to Theorem 3.3.23 [5].

Theorem 2.7. *Let conditions (S), (I $_\psi$) be fulfilled, $2\beta B - h < 0$, and*

$$C(\beta, h) < \frac{1}{2e}. \quad (24)$$

Then there exists a weak limit $\nu_\Lambda^{\beta, \phi} \rightarrow \nu^{\beta, \phi}$, $\Lambda \uparrow X$.

We intend to find some sufficient conditions on ϕ such that conditions of Theorem 2.7 are fulfilled. First, we derive another expression for $C(\beta, h)$.

$$C(\beta, h) = \operatorname{esssup}_{y \in X, t \in \mathbb{R}_+} \int_X \frac{\beta\phi(x, y) |1 - e^{(2\beta B - h - \beta\phi(x, y))t}|}{(2\beta B - h)(2\beta B - h - \beta\phi(x, y))} \sigma(dx).$$

For applications in genetics it seems reasonable to assume that

$$\phi(x, y) \geq 0, \quad \forall x, y \in X.$$

In this case, the stability condition (14) is fulfilled for $B = 0$ and

$$C(\beta, h) = \operatorname{esssup}_{y \in X, t \in \mathbb{R}_+} \int_X \frac{\beta\phi(x, y)}{h(h + \beta\phi(x, y))} (1 - e^{-(h + \beta\phi(x, y))t}) \sigma(dx).$$

From now on we assume for simplicity that $h(x) \equiv \text{const}$. Then, we have

$$C(\beta, h) = \operatorname{esssup}_{y \in X} \int_X \frac{\beta\phi(x, y)}{h(h + \beta\phi(x, y))} \sigma(dx).$$

We reformulate Theorem 2.7 for nonnegative $\phi(x, y)$.

Theorem 2.8. *Let $\phi(x, y)$ be nonnegative, and*

$$\operatorname{esssup}_{y \in X} \int_X \frac{\beta \phi(x, y)}{h(h + \beta \phi(x, y))} \sigma(dx) \leq \frac{1}{2e}. \quad (25)$$

1 *Then the weak limit $\nu_{\Lambda}^{\beta, \phi} \rightarrow \nu^{\beta, \phi}$, $\Lambda \uparrow X$ exists.*

2 **References**

- 3 1. Buerger R. The Mathematical Theory of Selection, Recombination, and Mutation. John Wiley, 2000.
- 4 2. Georgii H.-O., Zessin H. Large deviations and the maximum entropy principle for marked point random
- 5 fields. Probab. Theory Related Fields, 1993, **96**(2), 177–204.
- 6 3. Kondratiev Yu.G., Minlos R., Pirogov S. Generalized mutation-selection model with epistatics. Preprint,
- 7 2007.
- 8 4. Kondratiev Yu.G., Minlos R., Zhizhina E. Self-organized birth-and-death random systems. SFB–701,
- 9 Preprint 07–037, 2007.
- 10 5. Kuna T. Studies in configuration space analysis and applications. Bonn Mathematical Publications, 324.
- 11 Universität Bonn Mathematisches Institut, Dissertation, Rheinische Friedrich-Wilhelms-Universität
- 12 Bonn, Bonn, 1999.
- 13 6. Kuna T., Kondratiev Yu.G., Da Silva J.L. Marked Gibbs measures via cluster expansion. Methods
- 14 Funct. Anal. Topology, 1998, **4**(4), 50–81.
- 15 7. Malyshev V.A., Minlos R. Gibbs random fields. Mathematics and its Applications (Soviet Series),
- 16 volume 44. Kluwer Academic Publishers Group, Dordrecht, 1991; Cluster expansions, Translated from
- 17 the Russian by R. Kotecký and P. Holický.
- 18 8. Steinsaltz D., Evans S.N., Wachter K.W. A generalized model of mutation-selection balance with ap-
- 19 plications to aging. Adv. Appl. Math., 2005, **35**(1), 16–33.

Моделі селекційно-мутаційного балансу з епістатичною селекцією

20 **Ю.Кондратьев, Т.Куна, Н.Олеріх**

Університет Білефельда, Білефельд, Німеччина

Отримано 31 січня 2008 р.

21 Ми представляємо застосування процесів народження-знищення на конфігураційних просторах до

22 узагальненої моделі селекційно-мутаційного балансу. Модель описує старіння популяції як процес

23 накопичення мутацій в генотипі. В математично строгому підході мутації відповідають точкам у аб-

24 абстрактному просторі. Наша модель описує нескінченно-популяційну модель з безмежною кількістю

25 точок у континуумі. Динамічне рівняння, що описує систему, є типу Кімури-Маруяма. Проблема мо-

26 же бути поставлена в термінах еволюції станів (диференціальні рівняння) або, що є еквівалентно, за

27 допомогою формули Фейнмана-Каца. Досліджується питання існування розв'язку, його асимптоти-

28 чної поведінки, властивості граничного стану. У неепістатичному випадку проблема була поставлена

29 і розв'язана у [Steinsaltz D., Evans S.N., Wachter K.W., Adv. Appl. Math., 2005, 35(1)]. В нашій моделі ми

30 розглядаємо топологічний простір X як простір позицій мутацій та вплив на епістатичний потенціал.

31 **Ключові слова:**

32 **PACS:** 02.50.Ga

PROOFS