Accepted refereed manuscript of:

Farkas JZ, Hinow P & Engelstadter J (2013) Corrigendum to "Pathogen evolution in switching environments: A hybrid dynamical system approach" [Math. Biosci. 240 (2012) 70-75], *Mathematical Biosciences*, 241 (1), pp. 147-148.

DOI: <u>10.1016/j.mbs.2012.10.003</u>

© 2015, Elsevier. Licensed under the Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International <u>http://creativecommons.org/licenses/by-nc-nd/4.0/</u>

## Corrigendum to "Pathogen evolution in switching environments: a hybrid dynamical system approach"

József Z. Farkas

Division of Computing Science and Mathematics, University of Stirling, Stirling, FK9 4LA, United Kingdom; jzf@maths.stir.ac.uk

Peter Hinow

Department of Mathematical Sciences, University of Wisconsin – Milwaukee, P.O. Box 413, Milwaukee, WI 53201, USA; hinow@uwm.edu

Jan Engelstädter

Institute of Integrative Biology, Swiss Federal Institute of Technology, Universitätsstrasse 16, 8092 Zürich, Switzerland; j.engelstaedterOuq.edu.au<sup>1</sup>

## Abstract

We provide a corrected proof of Theorems 3.3 and 3.6 in the paper "Pathogen evolution in switching environments: a hybrid dynamical system approach", *Mathematical Biosciences* **240** (2012), p. 70-75.

In the proofs of Theorems 3.3 and 3.6 of our paper we cannot assume a special structure for the generator matrix Q with nearly identical rows (this would mean that the infinitesimal probability to jump into a new environment is independent of the current environment). Nevertheless, the idea of the proof, namely that the Lyapunov function can be written as a sum of functions, is correct.

To fix notation, we consider the switching differential equation

$$\frac{dP_i(t)}{dt} = P_i(t) \left( w_i^k - \sum_{j=1}^m w_j^k P_j(t) \right),\tag{1}$$

Preprint submitted to Mathematical Biosciences

 $<sup>^1\</sup>mathrm{Current}$ address: School of Biological Sciences, University of Queensland, Brisbane QLD 4072, Australia

where  $w_i^k > 0$  is the fitness value of genotype *i* in environment (host species) k and

$$\sum_{j=1}^{m} P_j(t) = 1.$$
 (2)

The switching process  $k = \alpha(t)$  is a Markov process with generator matrix Q(P) whose entries  $q_{kl}(P)$  are defined by

$$\mathbf{P}\{\alpha(t+\Delta t) = l \mid \alpha(t) = k, (P(s), \alpha(s)), s \le t\} = q_{kl}(P(t))\Delta t + o(\Delta t).$$
(3)

The elements  $q_{kl}$  of the generator matrix Q satisfy  $q_{kl} \ge 0$  for all  $k \ne l$  and  $\sum_{l \in \mathcal{M}} q_{kl} = 0$  for every  $k \in \mathcal{M}$ .

**Theorem 1.** Assume that the generator matrix  $Q = (q_{kl})_{k,l=1}^n$  is irreducible and let  $\pi$  be its unique stationary distribution. Let  $P_1$  be the genotype with the highest mean fitness, that is

$$\pi \cdot \mathbf{w}_1 > \pi \cdot \mathbf{w}_i \quad for \ all \ i = 2, \dots, m$$

$$\tag{4}$$

Then the equilibrium  $e_1$  is asymptotically stable in probability and all other equilibria are unstable in probability.

**Proof.** For i = 2, ..., m we set  $a_{i,1}^k = w_i^k - w_1^k$  for the difference of fitness values with respect to genotype 1 and  $\mathbf{a}_{i,1} = (a_{i,1}^1, ..., a_{i,1}^n)$ . Using the constraint (2), we eliminate  $P_1$  and obtain the *reduced systems* 

$$\frac{dP_i(t)}{dt} = a_{i,1}^k P_i(1-P_i) - P_i \sum_{j=2, \, j \neq i}^m a_{i,1}^k P_j,\tag{5}$$

for i = 2, ..., m and k = 1, ..., n. Notice that for fixed environment k the linear part of this system has a diagonal structure. We define

$$\beta_i := -\pi \cdot \mathbf{a}_{i,1} > 0,$$

with the last inequality holding true since genotype 1 has the higher mean fitness compared to every other genotype. For i = 2, ..., m we solve the systems of equations

$$Q\mathbf{c}_i = \mathbf{a}_{i,1} + \beta_i \mathbf{1}$$

for the vector  $\mathbf{c}_i = (c_i^1, \ldots, c_i^n)$  where **1** is the column vector with *n* entries 1. The right hand sides of these equation are orthogonal to the kernel of Qwhich is spanned by **1**, hence there exist solutions. For  $i = 2, \ldots, m$  and  $k = 1, \ldots, n$ , we define

$$V_i(P_i, k) = (1 - \gamma c_i^k) P_i^{\gamma}, \quad P_i > 0,$$

with  $0 < \gamma < 1$  yet to be selected, in such a way that all coefficients are positive. We have

,

$$\mathcal{L}V_{i}(P_{i},k) = \gamma(1-\gamma c_{i}^{k})P_{i}^{\gamma-1}(a_{i,1}^{k}P_{i}+o(1)) + \sum_{j=1}^{n}q_{kj}(1-\gamma c_{i}^{j})P_{i}^{\gamma}$$

$$= \gamma P_{i}^{\gamma}\left((1-\gamma c_{i}^{k})a_{i,1}^{k} - \sum_{j=1}^{n}q_{kj}c_{i}^{j}+o(1)\right)$$

$$= \gamma P_{i}^{\gamma}\left((1-\gamma c_{i}^{k})a_{i,1}^{k} - (a_{i,1}^{k}+\beta_{i})+o(1)\right)$$

$$= \gamma P_{i}^{\gamma}\left(-\gamma c_{i}^{k}a_{i,1}^{k} + \pi \cdot \mathbf{a}_{i,1}+o(1)\right),$$
(6)

where we have made use of the fact that the row sums of Q are zero. In order to make all the factors in parentheses negative, we have to choose  $0 < \gamma < 1$ such that the inequality

$$\pi \cdot \mathbf{a}_{i,1} < \gamma c_i^k a_i^k \tag{7}$$

holds. By assumption (4), the left hand side of inequality (7) is negative. Therefore, for those indices i and k for which  $c_i^k a_{i,1}^k \ge 0$ , no condition arises for  $\gamma$ . If on the other hand  $c_i^k a_{i,1}^k < 0$ , then we can select

$$0 < \gamma < \min_{i=2,\dots,m \atop k=1,\dots,n} \left\{ \frac{\pi \cdot \mathbf{a}_{i,1}}{c_i^k a_{i,1}^k} \, : \, c_i^k a_{i,1}^k < 0 \right\}.$$

Although the  $c_i^k$  are not explicitly known, this is a minimum of finitely many positive numbers. The Lyapunov function is the sum of functions of a single variable

$$V(P_2,\ldots,P_m,k) = \sum_{i=2}^m V_i(P_i,k)$$

and the condition of Proposition 8.6 in [1] follows from the linearity of the operator  $\mathcal{L}$  and the choice of  $\gamma$ .

To prove the unstability in probability of equilibrium  $e_i$  for i > 1, we use the constraint (2) to eliminate  $P_i$ . This results in the reduced systems

$$\frac{dP_i(t)}{dt} = a_{l,i}^k P_l(1 - P_l) - P_l \sum_{j \neq i,l}^m a_{j,i}^k P_j,$$
(8)

for  $l \neq i$  and  $a_{l,i}^k = w_l^k - w_i^k$ . For i = 2, ..., m let  $\mathbf{c}_i = (c_i^1, ..., c_i^n)$  be the solution of

$$Q\mathbf{c}_i = \mathbf{a}_{1,i} - \beta_i \mathbf{1}.$$

We set

$$V(P_1, \dots, P_{i-1}, P_{i+1}, \dots, P_m, k) = V(P_1, k) = (1 - \gamma c_i^k) P_1^{\gamma}, \quad P_1 > 0,$$

where  $0 > \gamma > -1$  has yet to be selected, small enough that all coefficients are positive. With a calculation similar to (6) we obtain

$$\mathcal{L}V(P_1, k) = \gamma (1 - \gamma c_i^k) P_1^{\gamma - 1} (a_{1,i}^k P_1 + o(1)) + \sum_{j=1}^n q_{kj} (1 - \gamma c_i^j) P_1^{\gamma}$$
$$= \gamma P_1^{\gamma} \left( (1 - \gamma c_i^k) a_{1,i}^k - \sum_{j=1}^n q_{kj} c_i^j + o(1) \right)$$
$$= \gamma P_1^{\gamma} \left( (1 - \gamma c_i^k) a_{1,i}^k - (a_{1,i}^k - \beta_i) + o(1) \right)$$
$$= \gamma P_1^{\gamma} \left( -\gamma c_i^k a_{1,i}^k + \pi \cdot \mathbf{a}_{1,i} + o(1) \right).$$

In order to make all the factors in parentheses positive (so that the entire expression becomes negative), we need to have

$$0 > \gamma > \max_{\substack{i=2,\dots m \\ k=1,\dots,n}} \left\{ \frac{\pi \cdot \mathbf{a}_{1,i}}{c_i^k a_{1,i}^k} : c_i^k a_{1,i}^k < 0 \right\}.$$

The expressions whose maximum is taken are all negative since  $\pi \cdot \mathbf{a}_{1,i} > 0$  by assumption (4). The condition of Proposition 8.7 in [1] is thereby verified.  $\Box$ 

## References

 G. G. Yin and C. Zhu, *Hybrid Switching Diffusions*, Springer, New York (2010)