Author's Accepted Manuscript

Evolution of dispersal under variable connectivity

Petteri Karisto, Éva Kisdi



www.elsevier.com/locate/yjtbi

PII:S0022-5193(16)30367-8DOI:http://dx.doi.org/10.1016/j.jtbi.2016.11.007Reference:YJTBI8859

To appear in: Journal of Theoretical Biology

Received date: 21 March 2016 Revised date: 31 October 2016 Accepted date: 8 November 2016

Cite this article as: Petteri Karisto and Éva Kisdi, Evolution of dispersal unde variable connectivity, *Journal of Theoretical Biology* http://dx.doi.org/10.1016/j.jtbi.2016.11.007

This is a PDF file of an unedited manuscript that has been accepted fo publication. As a service to our customers we are providing this early version o the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting galley proof before it is published in its final citable form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain

ACCEPTED MANUSCRIPT

Evolution of dispersal under variable connectivity

Petteri Karisto^{*} & Éva Kisdi

Department of Mathematics and Statistics University of Helsinki

Keywords: adaptive dynamics, dispersal polymorphism, evolutionary branching, kin competition, patch connectivity

Mathematics Subject Classification: 92D15, 92D25, 92D40

Abstract. The pattern of connectivity between local populations or between microsites sup-1 porting individuals within a population is a poorly understood factor affecting the evolution of 2 dispersal. We modify the well-known Hamilton-May model of dispersal evolution to allow for 3 variable connectivity between microsites. For simplicity, we assume that the microsites are ei-4 ther solitary, i.e., weakly connected through costly dispersal, or part of a well-connected cluster 5 of sites with low-cost dispersal within the cluster. We use adaptive dynamics to investigate the 6 evolution of dispersal, obtaining analytic results for monomorphic evolution and numerical re-7 sults for the co-evolution of two dispersal strategies. A monomorphic population always evolves 8 to a unique singular dispersal strategy, which may be an evolutionarily stable strategy or an 9 evolutionary branching point. Evolutionary branching happens if the contrast between connec-10 tivities is sufficiently high and the solitary microsites are common. The dimorphic evolutionary 11 singularity, when it exists, is always evolutionarily and convergence stable. The model exhibits 12 both protected and unprotected dimorphisms of dispersal strategies, but the dimorphic singular-13 ity is always protected. Contrasting connectivities can thus maintain dispersal polymorphisms 14 in temporally stable environments. 15

16

^{*}Corresponding author. Present address: Petteri Karisto, Department of Environmental Systems Science, ETH Zürich, Universitätstrasse 2, 8092 Zürich, Switzerland; email: petteri.karisto@usys.ethz.ch and eva.kisdi@helsinki.fi.

Download English Version:

https://daneshyari.com/en/article/5760123

Download Persian Version:

https://daneshyari.com/article/5760123

Daneshyari.com