

Modelling the Dynamic Structure of Biological State-Based Systems

I. Stamatopoulou^{a,*} P. Kefalas^b M. Gheorghe^c

^a*South-East European Research Centre, 17 Mitropoleos Str., Thessaloniki, 54624, Greece*

^b*Dept. of Computer Science, CITY College, 13 Tsimiski Str., Thessaloniki, 54624, Greece*

^c*Dept. of Computer Science, University of Sheffield, Regent Court, 211 Portobello Str., Sheffield S1 4DP, UK*

Abstract

The paper discusses the modelling aspects of systems with dynamic processes and dynamic structure. A combination of models bringing together the benefits of two paradigms, Population P Systems and Communicating X-machines, is introduced. A simple case study is used in order to illustrate the potential of the combined use of the two methods.

1 Introduction

Biological systems exhibit highly dynamic processes within a very dynamic environment. The individual components which they consist of demand modelling of evolving data structures and modelling of the control over their internal changing states. In addition, the systems overall imply modelling of their configuration, including the ability to exchange messages between individual components as well as the ability to re-structure their formation over time. Examples of such systems include colonies of ants or bees, flocks of birds, cell tissues etc. [4], [8], [21].

* Corresponding author. Tel.; ++30-2310-253-477; fax. ++30-2310-253-478.

Email addresses: istamatopoulou@seerc.info (I. Stamatopoulou), kefalas@city.academic.gr (P. Kefalas), M.Gheorghe@dcs.shef.ac.uk (M. Gheorghe).