Wojciech Bartoszek<sup>(1)</sup> and Małgorzata Pułka<sup>(2)</sup> DEPARTMENT OF MATHEMATICS, GDAŃSK UNIVERSITY OF TECHNOLOGY, UL. NARUTOWICZA 11/12, 80-233 GDAŃSK, POLAND e-mail: bartowk@mifgate.mif.pg.gda.pl<sup>(1)</sup> e-mail: mpulka@mif.pg.gda.pl<sup>(2)</sup>

## On dynamics of quadratic stochastic processes and their applications in biology

A quadratic stochastic operator  $\mathbf{Q} : \mathfrak{X} \to \mathfrak{X}$  is defined by a cubic (finite or infinite) array of nonnegative real numbers  $[q_{ij,k}]_{i,j,k\geq 1}$  which satisfy

(1)  $0 \le q_{ij,k} = q_{ji,k} \le 1$  for all  $i, j, k \ge 1$ ,

(2)  $\sum_{k=1} q_{ij,k} = 1$  for any pair (i, j),

where  $\mathfrak{X}$  is  $\ell^1$  or  $\ell^1_d$  equipped with a standard norm. The family of all quadratic stochastic operators is denoted by  $\mathfrak{Q}$ . Any quadratic stochastic operator (process)  $\mathbf{Q}$  may be viewed as a bilinear mapping  $\mathbf{Q} : \mathfrak{X} \times \mathfrak{X} \to \mathfrak{X}$  if we set  $\mathbf{Q}(\underline{x}, \underline{y})(k) = \sum_{i=1,j=1}^{n} x_i y_j q_{ij,k}$ . Clearly  $\mathbf{Q}$  is monotone (i.e.  $\mathbf{Q}(\underline{x}, \underline{y}) \ge \mathbf{Q}(\underline{u}, \underline{w})$  whenever  $\underline{x} \ge \underline{u} \ge 0$  and  $\underline{y} \ge \underline{w} \ge 0$ ) and is bounded as  $\sup_{\|\underline{x}\|_1, \|\underline{y}\|_1 \le 1} \|\mathbf{Q}(\underline{x}, \underline{y})\|_1 = 1$ . It follows that  $\mathbf{Q}$  may also be viewed as a mapping  $\mathbf{Q} : \mathcal{D} \times \mathcal{D} \to \mathcal{D}$ , where  $\mathcal{D}$  stands for the simplex of probability vectors. In population genetics a special attention is paid to a nonlinear mapping  $\mathcal{D} \ni \underline{p} \to \mathbb{Q}(\underline{p}) = \mathbf{Q}(\underline{p}, \underline{p})$ . Here  $\mathbb{Q} : \mathcal{D} \to \mathcal{D}$ . Roughly speaking  $\mathbb{Q}(\underline{p})$  represents a distribution of genes in the next generation if parent's gens have a distribution  $\underline{p}$ . In this simplified model the iterates  $\mathbb{Q}^k(\underline{p})$ , where  $k = 0, 1, \ldots$ , describe the evolution of a genom. Given an initial distribution  $\underline{p} \in \mathcal{D}$  one may ask about asymptotic behaviour of the trajectory (i.e. the sequence  $(\mathbb{Q}^n(\underline{p}))_{n\geq 0}$ ). Because of nonlinearity, the trajectories enjoy several unexpected features (as it was conjectured by S. Ulam). In this talk we discuss some generic properties in the set  $\mathfrak{Q}$ . We also present conditions for asymptotic stability of  $\mathbf{Q} \in \mathfrak{Q}$ .