

ERRATA

P. Juvan, T. Režen, D. Rozman, K. Monostory, J. Pascussi and A. Belič, *Towards Identification of Gene Interaction Networks of Human Cholesterol Biosynthesis* ACSi 2008, 55, 396–407

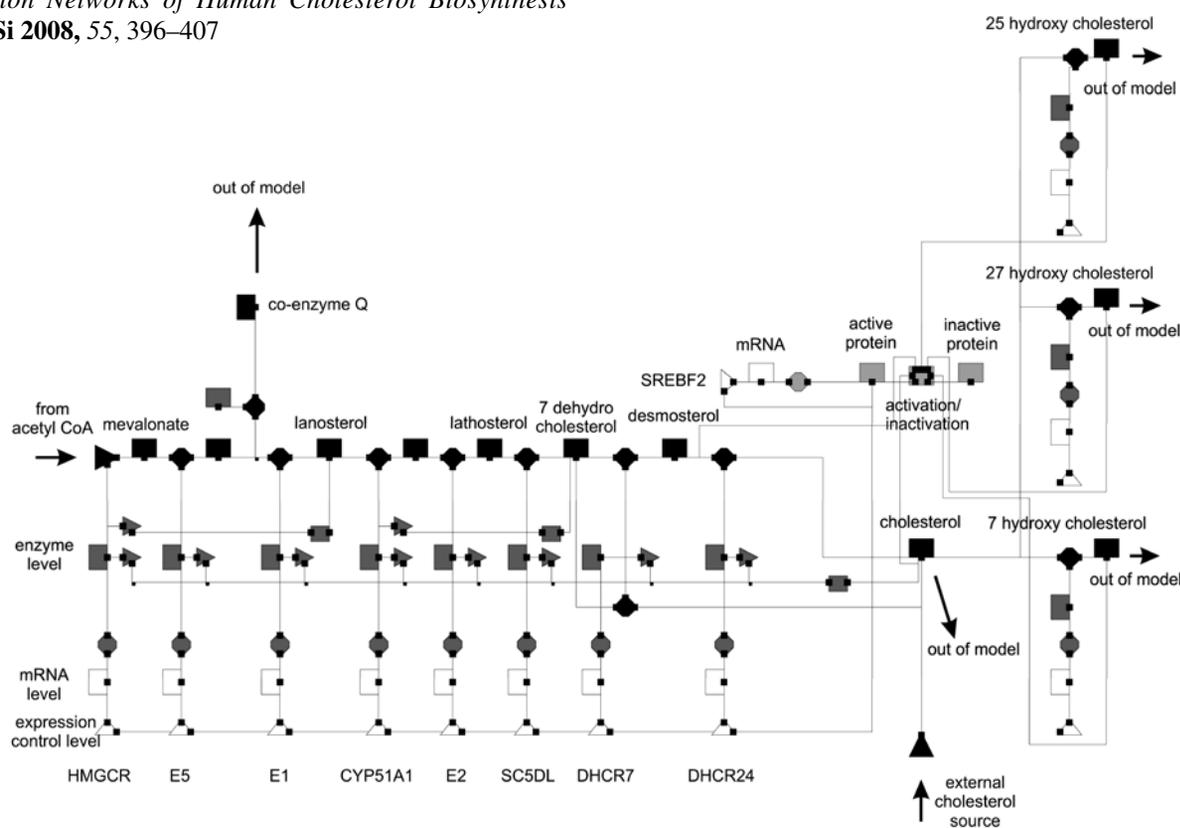


Figure 2. A simplified mathematical model of cholesterol biosynthesis. Black blocks represent metabolites (squares), enzyme reactions (circles) and metabolite sources (triangles). Dark grey blocks represent enzymes (large squares), their biochemical control (triangles and small squares), and enzyme production regarding gene expression (circles). White blocks represent gene expressions (squares) and their control (triangles). Light grey blocks represent transcription factors (squares), their production regarding gene expression (circles), and their activation/inactivation (squares). Gene names are listed in Table 1 in Supplement.