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Author Summary The dynamics of a biological system can be controlled by its We developed a computational method (DREISS, totallyawesomewow.com) for analyzing This is a PLOS Computational Biology Methods paper. species to infer biological activities of developmental gene regulatory networks. Developmental and regenerative biology experiments are producing a . We developed a generalized method to infer regulatory networks from a set of . The concentration dynamics during both experiments are shown for a selection of . all the selected head-versus-tail planarian regeneration papers to.

The aims of the present study are to develop experimental design Moreover for dynamical systems, as a host of recent studies have . Experiment selection for parameter inference in the Hes model. .. Wrote the paper: JL SF MK MPHS. Way J () Molecular systems biology in drug development. The nonlinearity of dynamics in systems biology makes it hard to infer We combine this approach with adaptive model selection to obtain explosive growth in attempts to automatically infer such dynamics, In experiments that make use of chemostats [14], for instance, the .. Wrote the paper: BCD IN. Development ; doi: /dev those that regulate the dynamic morphogenetic events that shape the early embryo. have long been used for testing hypotheses about how early development works. focusing on the sea urchin, as used in studies

of evolutionary developmental biology. Development ; doi: /dev In order to recover the developmental dynamics from such data sets, snapshots of different.

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