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Open-Ended Evolution and Innovation in a Deterministic Cellular Automata Universe ALYSSA ADAMS, Arizona State Univ, HECTOR ZENIL, Unit of Computational Medicine, Karolinska Institutet, Sweden, PAUL DAVIES, SARA WALKER, Arizona State Univ — One of the most remarkable features of life on Earth is the apparent open-ended evolution and innovation of the biosphere over its 2.3.5 billion year history. This is also one of the most perplexing features of biological evolution from the perspective of theoretical and computational modeling. Here we show that state-dependent dynamical rules can generate openended evolution for simple cellular automata (CA) organisms coupled to an external environment in a fully deterministic system. We present formal definitions of openended evolution as patterns that are non-repeating within the expected Poincare recurrence time of an isolated organism, and of innovation as trajectories not observed in isolated organisms. We demonstrate that a small subset of CA organisms implementing a state-dependent update rule, which is a function of the organisms current state and rule and the state of the environment, satisfy these minimal requirements for open-endedness and innovation. Our results demonstrate that an additional requirement for open-ended evolution and innovation is to remove the segregation of states and (fixed) dynamic laws characteristic of the physical sciences in attempts to model biological complexity.

> Alyssa Adams Arizona State Univ

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