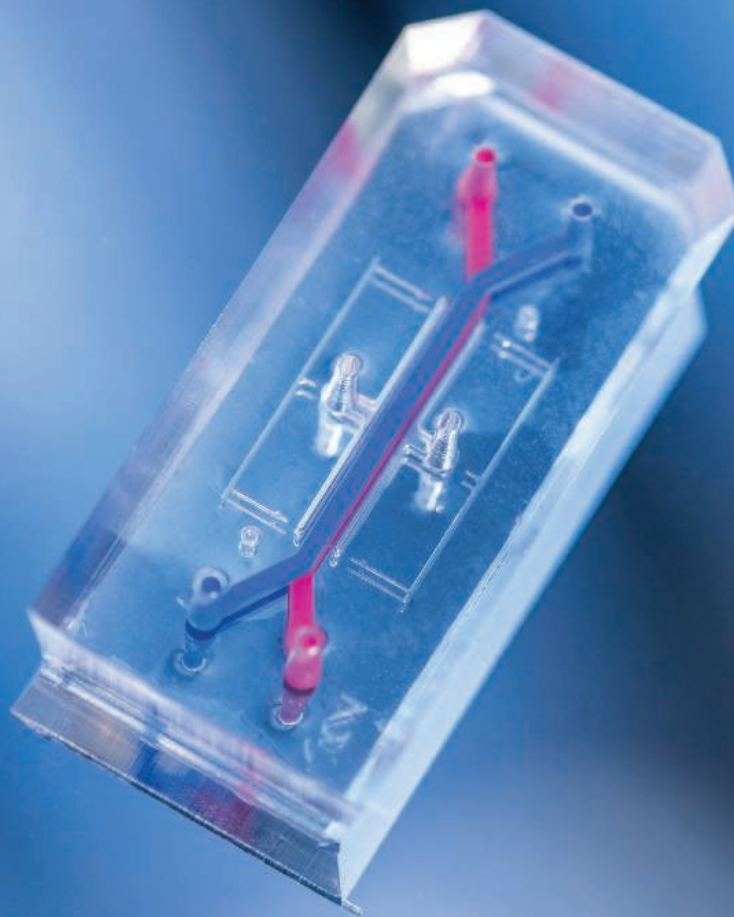


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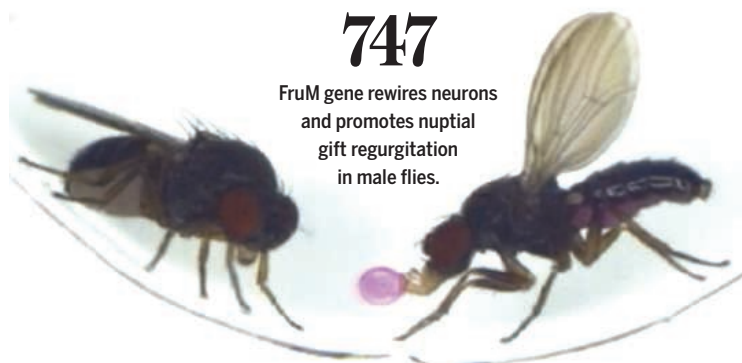


Predicting how biomolecules function requires understanding not just their sequence and structure but also their dynamic motions; yet, scalable methods for exploring these motions have been lacking. BioEmu is a deep learning system that rapidly generates diverse protein conformations, enabling fast, accurate insights into protein flexibility and function. This image shows an overlay of protein structures of a cyclic di-guanosine monophosphate receptor module sampled by BioEmu. See page 700. Illustration: N. Burgess/*Science*; Data: S. Lewis *et al.*, *Science* **389**, eadv9817 (2025).



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