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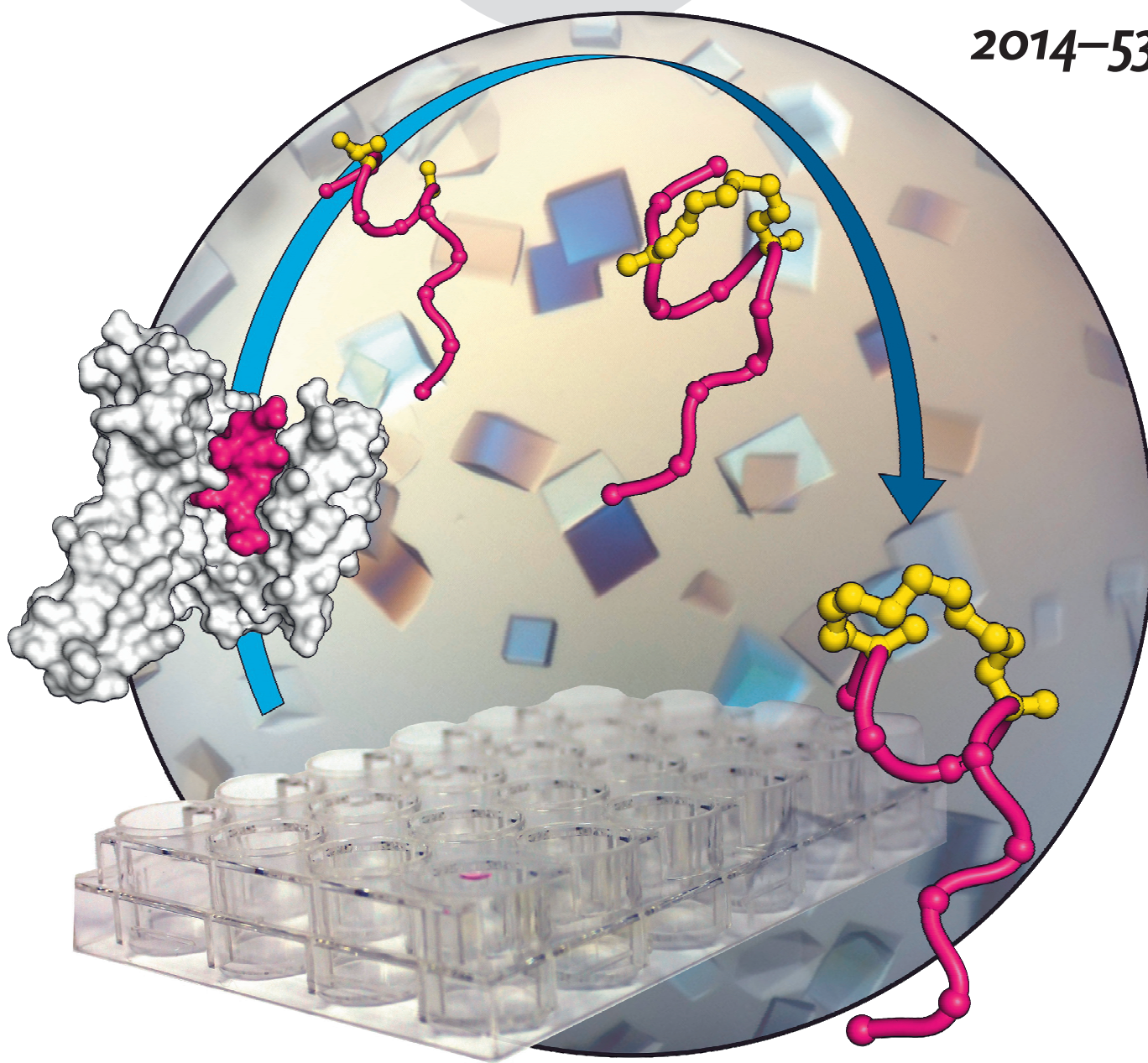
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Target-adapted hydrophobic cross-links ...

... were used to conformationally constrain a peptide (see picture, pink). In their Communication on page 2489 ff., T. N. Grossmann et al. present the iterative and rational evolution of these cross-links (yellow) based on X-ray crystal structures. The resulting macrocyclic peptides inhibit the protein–protein interaction between bacterial virulence factor ExoS and human protein 14-3-3.

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