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Cover picture: The universe of all possible binding curve line shapes for proteins containing four ligand-binding sites can be represented by the values of three shape parameters \(a\), \(b\), and \(c\). The points in the region of the \((a, b, c)\) space enclosed by the blue, red, and black sheets correspond to binding curves for which accurate estimates of binding parameters are possible (see Research Article by Middendorf and Aldrich, 121–147).