**Explicit Language and the Next Gen. Conversations: Albany at Tech.**

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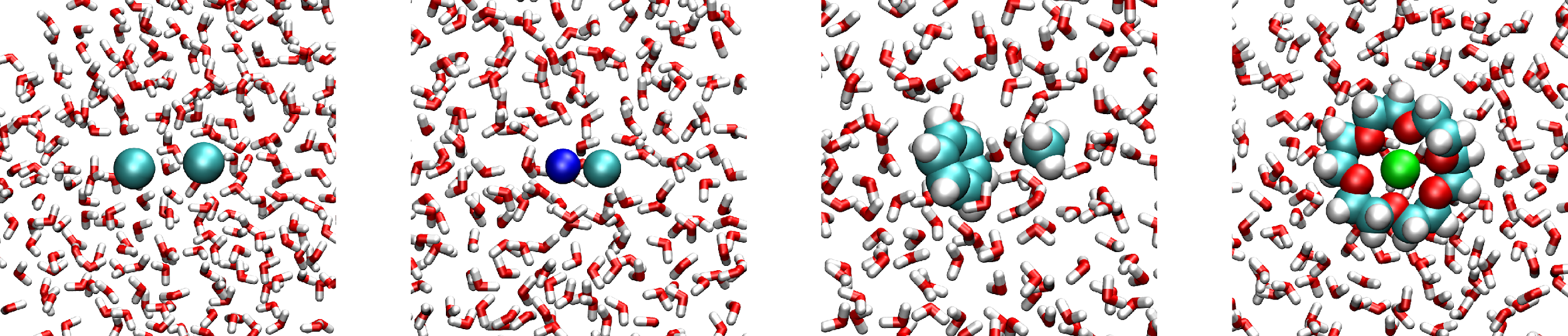


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**References**

K. A. Henzler-Wildman, K. A. & Kern. D. (2007). Dynamic personalities of proteins.*Nature* *450*, 964-972.

G. A. Huber, G. A. & Kim, S. (1996). Weighted-ensemble Brownian dynamics simulations for protein association reactions. *Biophys J* *70*, 97-110.

M. C. Zwier, M. C. & Chong, L. T. (2010). Reaching biological timescales with all-atom molecular dynamics simulations. *Curr. Opin. Pharmacol.* *10*, 745-752.