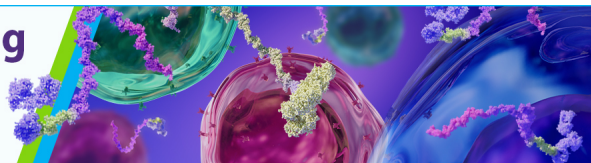


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On the cover: Structural depiction of Ab Fab fragments in complex with the 2009 pandemic influenza HA protein (blue). Two head domain-binding Fabs, 5J8 (orange) and Ab6649 (beige), and one stem domain-binding Fab, CR6261 (green), are shown. The newly discovered anchor Ab, P1-05, is modeled in yellow. Nagashima, K., J. V. Dzimianski, J. Han, N. Abbadi, A. D. Gingerich, F. Royer, S. O'Rourke, G. A. Sautto, T. M. Ross, A. B. Ward, R. M. DuBois, and J. J. Mousa. 2022. The pre-existing human antibody repertoire to computationally optimized influenza H1 hemagglutinin vaccines. *J. Immunol.* 209: 5–15.

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