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On the cover: Modeled pocket 6 of HLA-DRB1*0901 (shown with p6Lys in foreground), uniquely accommodates a wide range of residues (acidic, aliphatic, polar, and basic), due to the arrangement of three acidic and two basic residues (shown in stick form). B9Lys swings into this pocket although normally found pointing into neighboring pocket 9. James, E. A., A. K. Moustakas, J. Bui, R. Nouv, G. K. Papadopoulos, and W. W. Kwok. 2009. The binding of antigenic peptides to HLA-DR is influenced by interactions between pocket 6 and pocket 9. *J. Immunol.* 183: 3249–3258.
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