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Cover photograph (Copyright © 2013, American Society for Microbiology. All Rights Reserved.): Residues in the outer membrane porin PorB of *Neisseria gonorrhoeae* strain FA1090 that are involved in binding to the complement regulatory factors C4bp and factor H. The structure of the FA1090 PorB monomer was predicted using I-TASSER and visualized using DeepView Swiss-PdbViewer (v. 4.1). Extracellular loops 4 through 7 are labeled. (Left) Regions of the protein involved in C4bp binding are labeled in red in the context of the predicted structure. (Right) Regions of the protein involved in factor H binding are labeled in blue, and regions labeled in purple show an intermediate factor H binding phenotype when mutated. (See related article on p. 4383.)