

DNA minor groove sensing and widening by the CCAAT-binding complex.

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Abstract

The CCAAT box is a frequent element of eukaryotic promoters, and its specific recognition by the conserved heterotrimeric CCAAT-binding complex (CBC) constitutes a key step in promoter organization and regulation of transcription. Here, we report the crystal structures of the CBC from *Aspergillus nidulans* in the absence and in complex with double-stranded DNA at 1.8 Å resolution. The histone-like subunits HapC and HapE induce nucleosome-like DNA bending by interacting with the sugar-phosphate backbone. Minor groove sensing and widening by subunit HapB tightly anchor the CBC to the CCAAT box, as shown by structural and biochemical data. Furthermore, crucial interactions of the DNA duplex with subunit HapB provide an explanation for the sequence specificity of the CBC. The herein-described mode of transcription factor binding answers the question of how histone proteins gained sequence specificity for the CCAAT box.

Beteiligte Forschungseinheiten

[Molekulare und Angewandte Mikrobiologie Axel Brakhage](#) [Mehr erfahren](#)

Leibniz-HKI-Autor*innen



Axel A. Brakhage

[Details](#)



Peter Hortschansky

[Details](#)



Daniel H. Scharf

[Details](#)

Identifier

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