

Library Indian Institute of Science Education and Research Mohali



DSpace@IISERMohali (/jspui/)

- / Thesis & Dissertation (/jspui/handle/123456789/1)
- / Doctor of Philosophy (PhD) (/jspui/handle/123456789/268)
- / PhD-2013 (/jspui/handle/123456789/269)

Please use this identifier to cite or link to this item: http://hdl.handle.net/123456789/1550

Title: Role of the bacterial nucleoid associated protein "HU" in cell-cell and cell-DNA interactions through

the binding of HU with eDNA and Lipopolysaccharide

Authors: Bhishem (/jspui/browse?type=author&value=Bhishem)

Keywords: DNA interactions

Bacterial nucleoid associated protein

Lipopolysaccharide

Issue Date: Sep-2019

Publisher: IISER Mohali

Abstract:

Bacteria in biofilms are embedded within a matrix of extracellular DNA (e-DNA) derived from the lysis of bacterial cells, or cellular secretions. In biofilms, negative charges decorate surfaces of both bacteria and DNA, creating scope for repulsive interactions. An abundant and non- sequencespecific DNA-binding protein such as HU, which is decorated with positive charges could potentially bind to both DNA and to bacteria, to function as a charge-neutralizing glue. HU is already known to be present in bacterial biofilms (in association with e-DNA) and limiting for biofilm formation (with anti-HU antibodies disrupting biofilms). The work in this thesis demonstrates: (1) that HU binds to free lipopolysaccharide (fLPS) as well as to the surfaces of bacterial cells [i.e., to cellular LPS (cLPS) present in bacterial outer membranes]; (2) that binding of HU to fLPS or cLPS can involve either (a) HU's canonical DNA-binding site, or (b) HU's non-canonical DNA-binding site; (3) that addition of micellar fLPS to free HU (fHU) generates large molecular assemblies; (4) that addition of fHU to cells bearing cLPS generates large cellular assemblies (bacterial clumps); (5) that the charged head-group of the lipid A component of LPS contains two hexose-linked sugarphosphate moieties that bind to lysine/arginine residues on fHU's DNA-binding sites in specific geometries. Further, the thesis (6) examines the stability of HU's dimeric interface, and (7) constructs a protein-engineered (HU simulacrum) construct containing both types of DNA-binding sites, with other regions removed.

URI: http://hdl.handle.net/123456789/1550 (http://hdl.handle.net/123456789/1550)

Appears in PhD-2013 (/jspui/handle/123456789/269) Collections:

Files in This Item:

File Description Size Format

PH13030.pdf
(/jspui/bitstream/123456789/1550/3/PH13030.pdf)

Show full item record (/jspui/handle/123456789/1550?mode=full)

Items in DSpace are protected by copyright, with all rights reserved, unless otherwise indicated.

Admin Tools

Edit...

Export Item

Export (migrate) Item

Export metadata