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Title:	Cryo-EM elucidates mechanism of action of bacterial pore-forming toxins.
Authors:	Mondal, Anish Kumar (/jspui/browse?type=author&value=Mondal%2C+Anish+Kumar) Lata, Kusum (/jspui/browse?type=author&value=Lata%2C+Kusum) Singh, Mahendra (/jspui/browse?type=author&value=Singh%2C+Mahendra) Chatterjee, Shamaita (/jspui/browse?type=author&value=Chatterjee%2C+Shamaita) Chauhan, Aakanksha (/jspui/browse?type=author&value=Chauhan%2C+Aakanksha) Puravankara, Sindhoora (/jspui/browse?type=author&value=Puravankara%2C+Sindhoora) Chattopadhyay, Kausik (/jspui/browse?type=author&value=Chattopadhyay%2C+Kausik)
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Abstract:	Pore-forming toxins (PFTs) rupture plasma membranes and kill target cells. PFTs are secreted as soluble monomers that undergo drastic structural rearrangements upon interacting with the target membrane and generate transmembrane oligomeric pores. A detailed understanding of the molecular mechanisms of the pore-formation process remains unclear due to limited structural insights regarding the transmembrane oligomeric pore states of the PFTs. However, recent advances in the field of cryo-electron microscopy (cryo-EM) have led to the high-resolution structure determination of the oligomeric pore forms of diverse PFTs. Here, we discuss the pore-forming mechanisms of various PFTs, specifically the mechanistic details contributed by the cryo-EM-based structural studies.
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