

MiniReview

Type III secretion: The bacteria-eukaryotic cell express

Luís Jaime Mota¹, Isabel Sorg, Guy R. Cornelis^{*}

Biozentrum der Universität Basel, Biozentrum, Klingelbergstrasse, 50-70 CH4051 Basel, Switzerland

Received 29 March 2005; accepted 17 August 2005

First published online 8 September 2005

Edited by I. Henderson

Abstract

Type III secretion (T3S) is an export pathway used by Gram-negative pathogenic bacteria to inject bacterial proteins into the cytosol of eukaryotic host cells. This pathway is characterized by (i) a secretion nanomachine related to the bacterial flagellum, but usually topped by a stiff needle-like structure; (ii) the assembly in the eukaryotic cell membrane of a translocation pore formed by T3S substrates; (iii) a non-cleavable N-terminal secretion signal; (iv) T3S chaperones, assisting the secretion of some substrates; (v) a control mechanism ensuring protein delivery at the right place and time. Here, we review these different aspects focusing in open questions that promise exciting findings in the near future.

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Keywords: Bacterial pathogenesis; EPEC; *Salmonella*; *Shigella*; Type III secretion; *Yersinia*

1. Introduction

Type III secretion (T3S) systems are protein export devices essential for the interaction between Gram-negative bacteria and their eukaryotic hosts. These systems are present in many pathogenic bacteria for animals and plants, but also in endosymbionts. Bacteria use T3S to translocate proteins across lipid membranes and to inject some of them, called “effectors”, into the cytoplasm of host cells. Inside the eukaryotic cell cytoplasm, the effectors thwart or hijack host cell signaling to the benefit of the bacteria, in the case of pathogens, or both organisms, in the case of symbionts.

With the sequencing of bacterial genomes, the list of species known to harbour T3S systems has become too long to be listed exhaustively (Table 1). Among the animal pathogens, T3S have been most intensively studied in *Yersinia* spp., *Salmonella* spp., *Shigella* spp., *Pseudomonas aeruginosa*, and enteropathogenic *Escherichia coli* (EPEC) and enterohaemorrhagic *E. coli* (EHEC). The most studied T3S system from a plant pathogen is probably that of *P. syringae*, but this secretion system has also been identified and studied in several other plant pathogens.

The T3S substrates are synthesized in the bacterial cytoplasm and are secreted by a nano-machine, called injectisome, across the bacterial inner membrane, periplasm and the bacterial outer membrane. Some of these secreted proteins, called “translocators” insert in the host cell plasma membrane and mediate the translocation of the effectors across eukaryotic plasma or vacuolar membranes into the host cell cytosol. Secretion of the effectors across the bacterial membranes and translocation through the eukaryotic membrane is thought

^{*} Corresponding author. Tel.: + 41 61 267 2121; fax: +41 61 267 2118.

E-mail addresses: j.mota@unibas.ch (L.J. Mota), guy.cornelis@unibas.ch (G.R. Cornelis).

¹ Present address: Imperial College London, Centre for Molecular Microbiology and Infection, Armstrong Road, Flowers Building, London SW7 2AZ, United Kingdom.

Table 1
T3S systems of bacterial pathogens for mammals

Bacterium	Diseases	T3S system(s) function(s) ^a
<i>Aeromonas</i> spp.	Opportunistic human pathogen causing gastroenteritis and septicemia (<i>A. hydrophila</i>) Furunculosis in salmonids (<i>A. salmonicida</i>)	<u>Asc</u> Cytotoxicity on fish cells
<i>Bordetella</i> spp.	Respiratory tract infections, whooping cough (<i>B. pertussis</i> , <i>B. parapertussis</i>) Infection of four-legged animals (<i>B. bronchiseptica</i>)	<u>Bsc</u> Establishment of long term infection Downregulation of inflammation
<i>Burkholderia</i> spp.	Melioidosis: pneumonia and skin abscesses (<i>B. pseudomallei</i>)	<u>Bp1</u> (<i>B. pseudomallei</i>) ?
	Glanders (<i>B. mallei</i>)	<u>Bp2</u> (<i>B. pseudomallei</i> ; <i>B. mallei</i> ; <i>B. thailandensis</i>) ?
	Pulmonar infection of cystic fibrosis patients (<i>B. cepacia</i> complex)	<u>Bp3</u> (<i>B. pseudomallei</i> ; <i>B. mallei</i> ; <i>B. thailandensis</i>) Survival and escape from macrophage vacuoles; Invasion of non-phagocytic cells <u>Bcsc</u> (<i>B. cepacia</i> complex) Important for virulence in the murine model
Chlamydiaceae	Sexual transmitted diseases, trachoma (<i>C. trachomatis</i>) Pneumonia, atherosclerosis (<i>C. pneumoniae</i>)	<u>Sct</u> Cell internalization
<i>Chromobacterium violaceum</i>	Sporadic infections in humans and mammals resulting in fulminant septicaemia that resembles melioidosis	<u>Civ</u> Similar to Inv (SPI-1) and Mxi-Spa <u>Csa</u> Similar to Ssa (SPI-2)
<i>Citrobacter rodentium</i>	Murine colonic hyperplasia, attaching/effacing lesions 1	<u>Esc</u> Adhesion to the intestinal epithelium Cytoskeleton rearrangements
<i>Desulfovibrio vulgaris</i>	Implicated in ulcerative colitis	<u>Dsc</u> ?
<i>Edwardsiella tarda</i> <i>Edwardsiella ictaluri</i>	Hemorrhagic septicemia in fishes, diverse infections in humans	<u>Eds</u> Survival and replication in fish phagocytes
Enteropathogenic <i>Escherichia coli</i> (EPEC) and enterohemorrhagic <i>E. coli</i> (EHEC)	Diarrhoea, attachment/effacement lesion (EPEC)	<u>Esc</u>
	Diarrhoea, attachment/effacement lesion, haemorrhagic colitis (EHEC)	Colonization Cell attachment Actin remodelling Pedestal formation
<i>Pseudomonas aeruginosa</i>	Pneumonias and chronic bronchopneumonia in patients with cystic fibrosis; ulcerative keratitis	<u>Psc</u> Cytotoxicity Block phagocytosis

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