

Characterization of Outer Membrane Vesicle Release in *Vibrio vulnificus*.

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The production of outer membrane vesicles (OMVs) by Gram-negative bacteria is an active process that takes place when a section of the outer membrane bulges off to form periplasm-containing vesicles without causing bacterial lysis. OMVs directly influence bacteria-host interaction and pathogenesis through their ability to modulate immune responses, contribute to biofilm formation, and deliver toxins and other virulence factors to host cells [1]. The ability of OMVs to bind to eukaryotic membranes makes them potential delivery vehicles for antibiotics and candidate vaccines against pathogens such as *Vibrio cholerae*, indicating their significant biotechnological potential [2]. OMVs have also been recognized to be important components of bacterial biofilms. *V. vulnificus* is a Gram-negative bacterium which colonizes filter-feeding shellfish. This opportunistic pathogen is responsible for 95% of seafood-related deaths in the United States and is categorized as a category B biodefense priority pathogen by the National Institutes of Allergies and Infectious Diseases [3]. Virulence in *V. vulnificus* is regulated by structural components such as cellular appendages (flagella and pili) and the capsular polysaccharide. Even though their contribution to bacterial pathogenesis is widely recognized, the mechanism of OMV production is not yet understood.

Here, we use multiple TEM imaging techniques to characterize the unusual secretion and spatial arrangement of OMVs. We present further cryo-electron tomography data [4] (Fig. 1) of OMV release and distribution in *Vibrio vulnificus* and several of its mutant forms in order to link these results to known differences in virulence. Because of the low contrast of cryo TEM, the capsular polysaccharide remained obscure. In order to visualize the role of the capsular polysaccharide in providing this unique spatial arrangement we employ a high pressure freezing technique, self pressurized rapid freezing (SPRF) [5]. For cryo EM and cryo ET, *V. vulnificus* was grown in LB broth at 30 degrees C with shaking at 250 rpm overnight, then diluted into fresh LB and grown and additional 4 hours. Bacteria were applied to copper 200 mesh grids with R2/1 Quantifoil carbon supports and plunge frozen into liquid ethane. For SPRF, colonies from a fresh LB plate were scraped into a paste and thinned with enough LB broth to facilitate pipetting into a 16mm long and 0.65 mm o.d. copper tube. After crimping the ends closed, the tube was manually plunged into liquid propane. Tubes were sectioned and freeze-substituted.

Both cryo EM and cryo ET reveal a regular arrangement of OMVs spaced approximately 100 nm from the outer membrane and in successive rings with 200 nm spacing. We do not observe this discrete spacing in reduced virulence mutants lacking the capsular polysaccharide. We were able to visualize the capsular polysaccharide using the SPRF technique, and observe that the capsule extends in a radius around the outer membrane a distance of ~100 nm, and that the individual OMVs retained these 100 nm polysaccharide chains which remain joined at the distal end, creating 200 nm spacing between the outer membrane and OMVs, as well as between rows of OMVs (Fig. 2). We hypothesize that the capsular polysaccharide has a definitive role in OMV distribution and virulence in *V. vulnificus*.

References:

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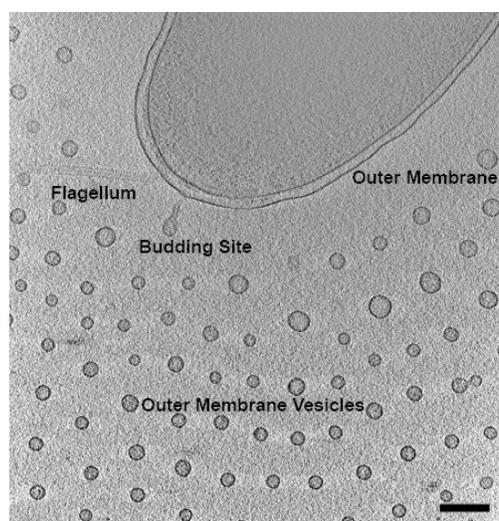


Figure 1. Cryo ET slice of plunge-frozen *V. vulnificus* producing outer membrane vesicles. Scale bar 200 nm.

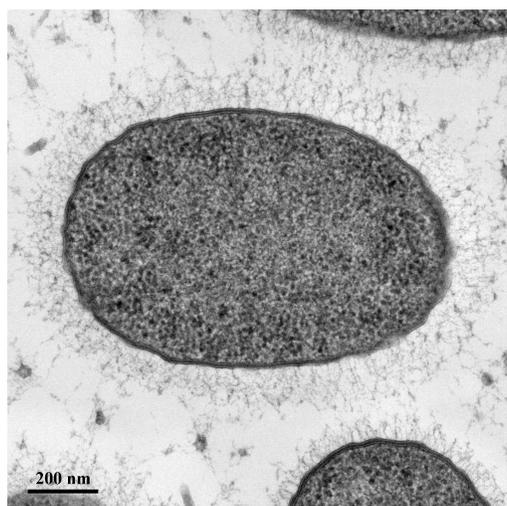


Figure 2. SPRF, freeze-substituted section of *V. vulnificus* reveals the capsular polysaccharide chains extending from the outer membrane and attached to surrounding OMVs. Scale bar 200 nm.