

CHARACTERIZATION OF THE VIRB7-VIRB9 INTERACTION AND INITIAL
NMR STRUCTURE DETERMINATION OF VIRB7 FROM *XANTHOMONAS
AXONOPODIS* PV. *CITRI* CHROMOSOMAL TYPE IV SECRETION SYSTEM

Souza, D.P.¹, Salinas, R.K.¹, Farah, C.S.¹

¹Departamento de Bioquímica, IQ-USP, São Paulo, Brazil

Xanthomonas axonopodis pv. *citri* (*Xac*) is a phytopathogen that causes citrus canker in orange trees. Among the possible virulence determinants in the *Xac* genome is the chromosomal Type IV secretion system (T4SS), an important transenvelope apparatus that secretes proteins and DNA in many plant and animal pathogens. The model T4SS found in *Agrobacterium tumefaciens* is constituted by twelve structural proteins: VirB1-VirB11 and VirD4. We expressed the *Xac* VirB7_{His-24-139} and VirB9₃₄₋₂₅₅ fragments in *Escherichia coli*. The proteins were purified by standard chromatographic methods and characterized by SDS-PAGE, mass spectrometry, fluorescence and circular dichroism. VirB9₃₄₋₂₅₅ was expressed as an insoluble polypeptide, purified in the presence of urea and refolded by dialysis. The VirB7-VirB9 interaction was assayed by fluorescence titration, indicating a K_d of $\sim 4 \times 10^{-8}$ M. For NMR studies, VirB7_{His-24-139} was induced in minimum media for incorporation of ¹⁵N and ¹³C and the His-tag was removed by thrombin proteolysis. Several multidimensional spectra were collected for resonance assignment, including ¹⁵N-HSQC, ¹³C-HSQC, HNC(O), HNCA, HN(CO)CA, HNCACB, CBCA(CO)NH, HBHA(CO)NH, ¹⁵N-Tocsy-HSQC and HC(CO)NH-Tocsy. More than 50% of the backbone and the side chain C β and H β have been assigned. The VirB7-VirB9 interaction was monitored by ¹⁵N-HSQC of ¹⁵N-labelled VirB7 before and after addition of VirB9, indicating that the VirB7 C-terminal remains highly flexible in the VirB7-VirB9 complex.

Supported by: CNPq and Fapesp