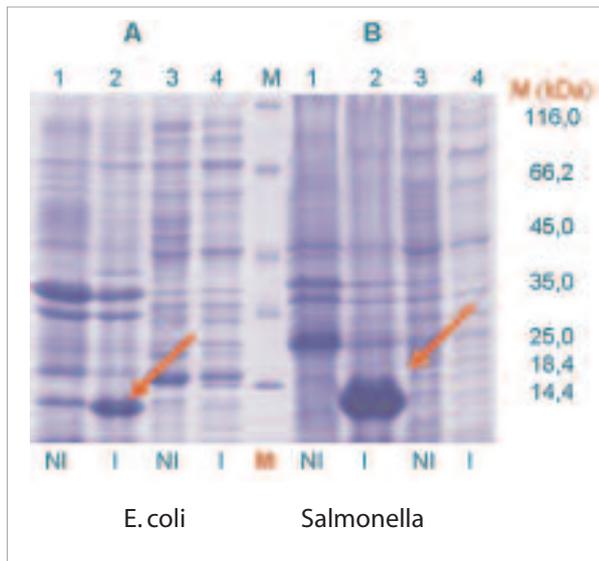


CLONING AND EXPRESSION OF PROTEINS IDENTIFIED IN THE GENOME OF *Leptospira interrogans* serovar *Copenhageni* FOR POTENTIAL VACCINE AND DIAGNOSTIC USE

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Purification of recombinant proteins. SDS-PAGE (15%) of purified recombinant proteins obtained by metal affinity chromatography. Proteins encoded by (1) LIC10009, (2) LIC10191, (3) LIC11947, (4) LIC12730, (5) LIC10494, (6) LIC12906

Leptospirosis is considered the most widespread zoonotic disease caused by the spirochaete from the *Leptospira* genus. In Brazil, 80% of all the diagnosed cases are due to *Leptospira interrogans* serovar *Copenhageni*. Rats are the main reservoir in urban centers and the bacteria are disseminated through animal urines. Around 10,000 cases of leptospirosis are annually notified to FUNASA (Health Ministry, Brazil), epidemically associated to the rainy seasons (<http://www.funasa.gov.br>). A promising way to treat this health problem would be a vaccination program, since the control of rats is a difficult task. Thus, the development of a suitable vaccine against leptospirosis is of seminal importance, since up to date, there is no licensed human vaccine against leptospirosis. Furthermore, the general clinical signs of this disease are often confounded with those of other diseases, like dengue and flu symptoms, reinforcing the need for an efficient diagnose assay for leptospirosis, not yet available in the market. Recently, the genome of the *Leptospira interrogans* serovar *Copenhageni* was described. The information of the genome would help to identify potential *Leptospira* vaccine antigen candidates or antigens for diagnose applications. The present project is divided in 3 subprojects, aiming: 1) cloning of 8 selected hemolysins. Recombinant proteins will be expressed in *E. coli* and further evaluated for their potential use as vaccine antigens and for potential use in diagnose. Biochemical and immunological characterization will also be performed; 2) cloning of 5 selected lipoproteins. Recombinant proteins will be expressed in *E. coli* and further evaluated for their potential use as vaccine antigens and for potential use in diagnose. Biochemical and immunological characterization will also be performed; 3) cloning and expression of 5 of the above antigens in *Salmonella typhimurium*. The recombinant *Salmonella* will be evaluated as live vaccine against *Leptospira*.

SUMMARY OF RESULTS TO DATE AND PERSPECTIVES

We have accumulated data from dozens of potential antigens of *Leptospira*. Though we have initially proposed the study of 8 hemolysins and 10 lipoproteins, being 10 of them also expressed as live vaccine in *Salmonella thyphimurium*, we have expanded this universe. For instance, in the case of the subproject dealing with hemolysins, we started with 8 ORFs, but we ended up with 20 ORFs. All these were cloned and evaluated for expression in *E. coli*. This represents our first step in our experimental approach to uncover new antigens for vaccine or diagnostic kit development for leptospirosis. If these ORFs were expressed in *E. coli*, the study progress to step 2. If not, we try to express them using other *E. coli* strains, conditions. If they are



Leptospira antigen crystals for determining three-dimensional structures

still not expressed, we look for new ORFs to replace them. For those that we were able to express, the proteins are purified and antiserum is produced against each of them.

These sera are immunocharacterized and used to screen a panel of different *Leptospira* serovars to define if they are conserved antigens among them. Those that are conserved, they are selected for the next steps, since we are searching for antigens that would compose a universal vaccine against all the *Leptospira* serovars but not a vaccine against a specific serovar. These ORFs are biologically

characterized, for reactivity against patient sera as well as assayed as protective antigens in a hamster model of immunization and challenge with pathogenic virulent *Leptospira*. At least one antigen we presented in *Salmonella* was able to elicit partial protection in animal model. We were able to identify several conserved antigens among the serovars that would be useful for leptospira diagnosis, new adhesins that would mediate leptospira attachment to host cells, new antigens that mediate the inflammatory response observed during leptospira infection, besides other findings. We can anticipate that the project will at least increase our comprehension of this bacteria and the disease it causes that will allow us in the future to interfere with the progress of the disease.

MAIN PUBLICATIONS

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