





Gene silencing in pathogenic *Leptospira* by CRISPR-interference

CRISPR/Cas9 targeting genomic Cells using specific specif

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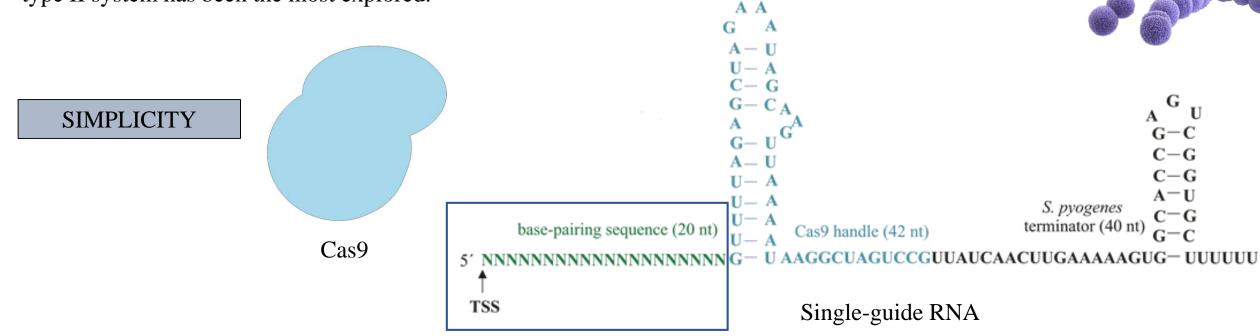
INTRODUCTION

CRISPR/Cas system

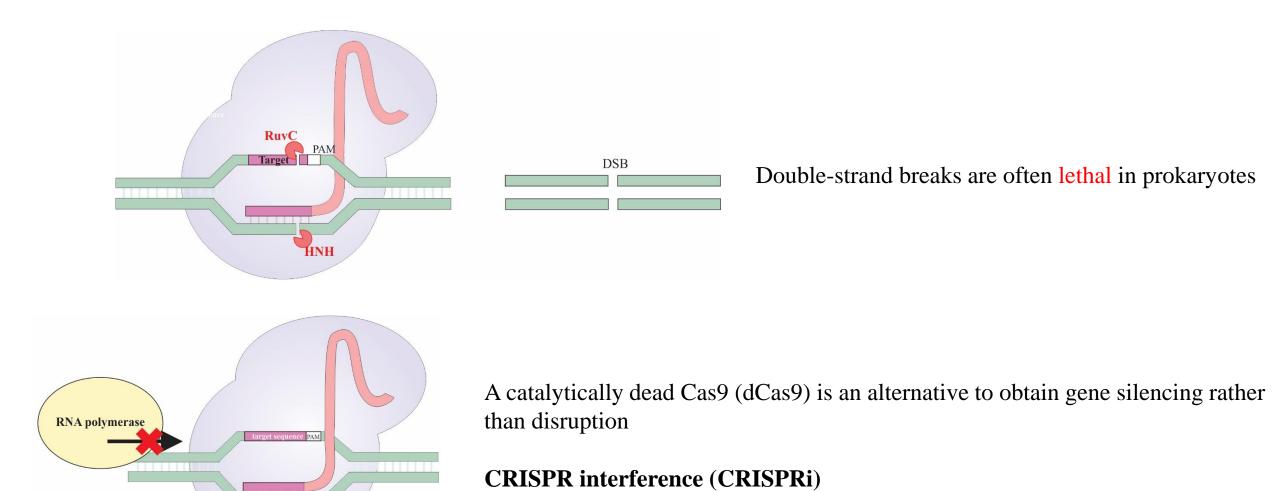
This system has been found in the genomes of most Archaea and Bacteria, playing an important role in their immunity against phages and invading plasmids

Clustered Regularly Interspaced Short Palindromic Repeats

To date, there are several CRISPR systems characterized, and among them, the *Streptococcus pyogenes* type II system has been the most explored.

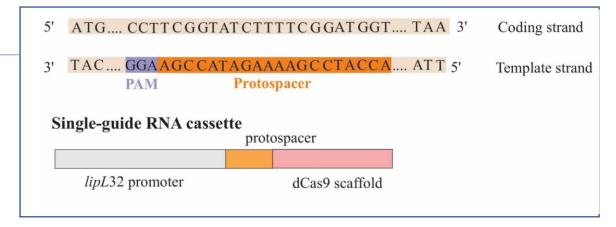


INTRODUCTION





METHODS



sgRNA cassette

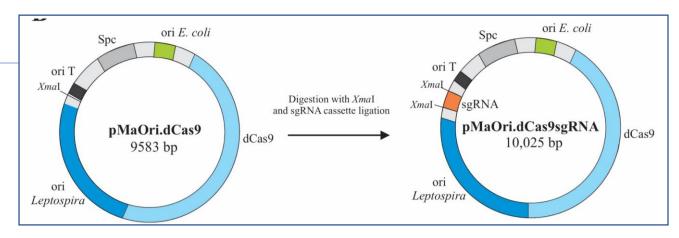
METHODS

5' ATG.... CCTTCGGTATCTTTTCGGATGGT.... TAA 3' Coding strand
3' TAC.... GGAAGCCATAGAAAAGCCTACCA.... ATT 5' Template strand
PAM Protospacer

Single-guide RNA cassette

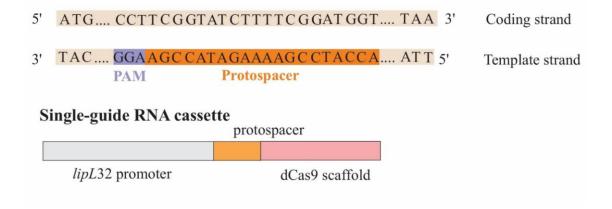
protospacer

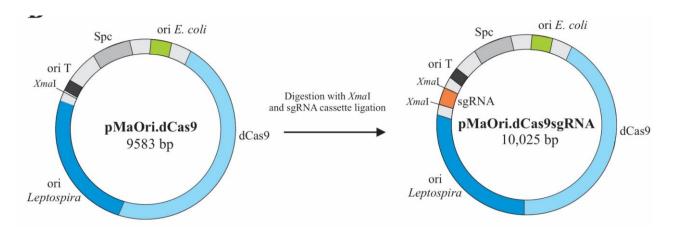
lipL32 promoter dCas9 scaffold



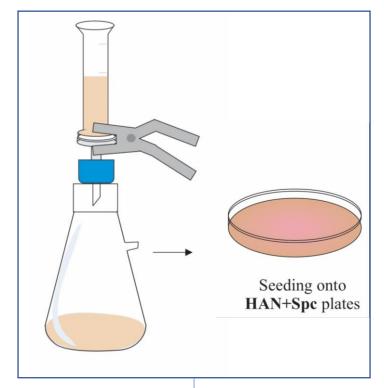
Plasmid construction

METHODS



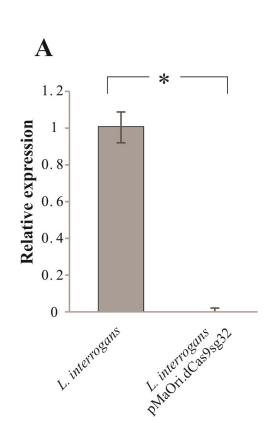


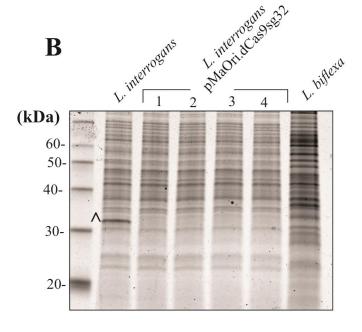
Leptospira spp. transformation by conjugation

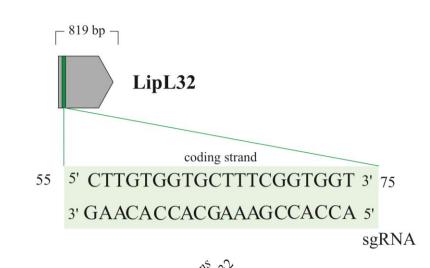


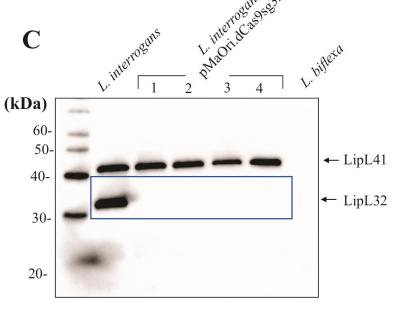
Proof of concept: silencing of the highly transcribed *lipL32* gene

The application of CRISPRi to *L. interrogans* resulted in a **complete silencing** of the major LipL32 protein



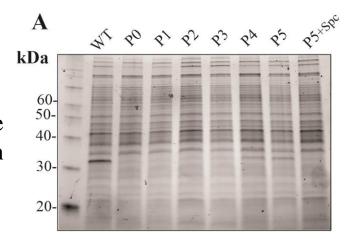


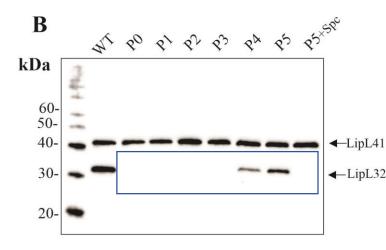




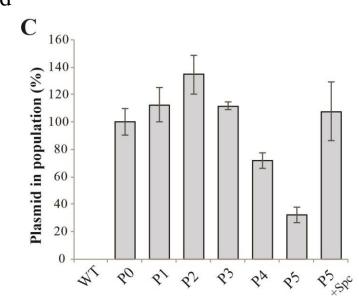
Stability of gene silencing

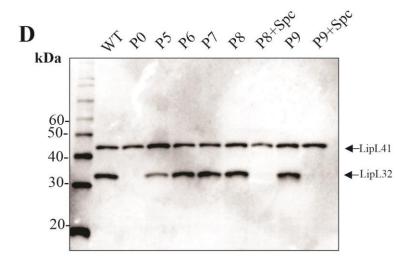
Even though CRISPRi tool is episomal, stable silencing could be observed until passage 3 in HAN media with no antibiotic





Growth in the presence of spectinomycin could completely restore silencing.

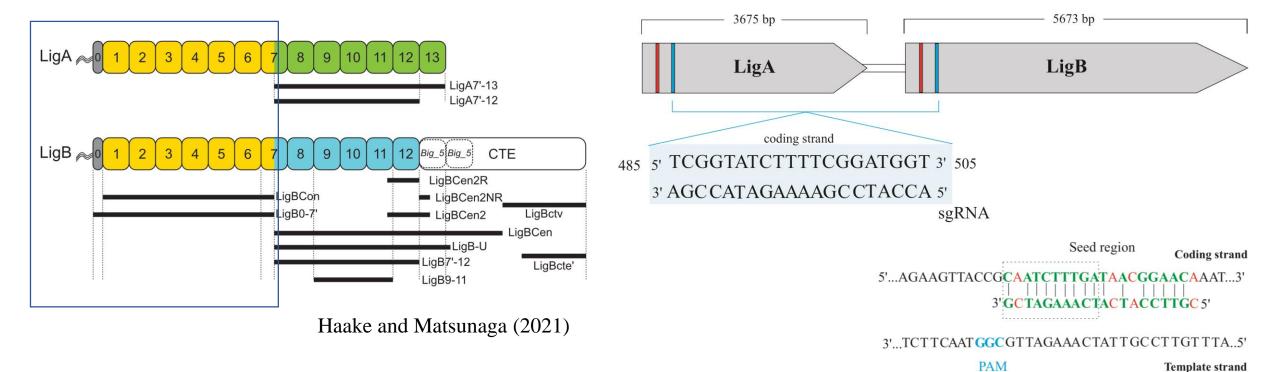




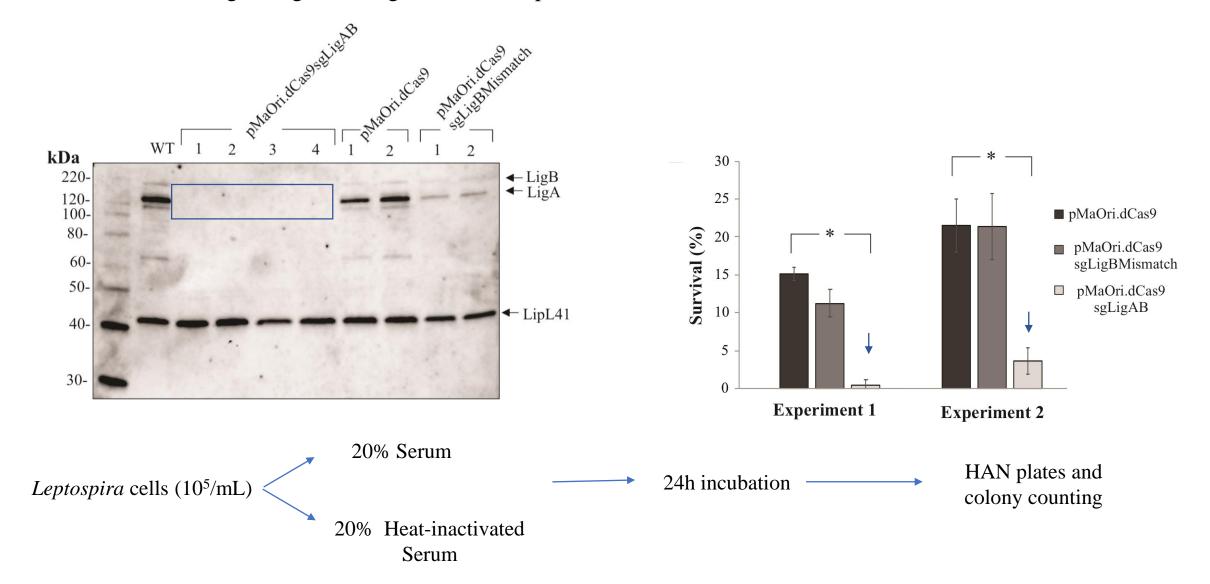
In vitro studies have demonstrated the ability of LigA and LigB protein to interact with host complement molecules

These proteins display overlapping properties and could be responsible for serum resistance displayed by pathogenic *Leptospira* strains.

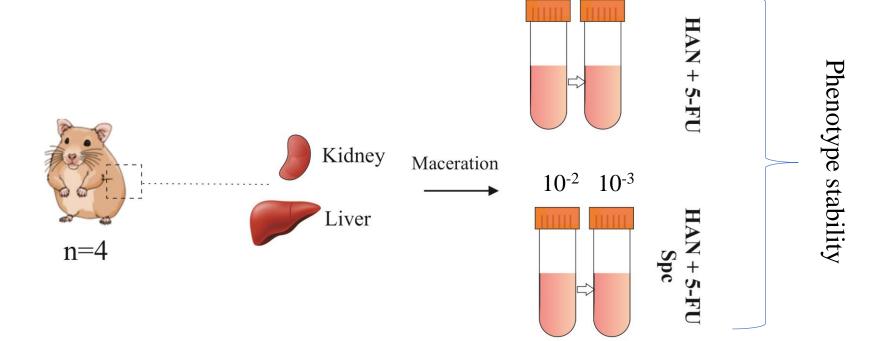
Conserved N-terminal region



Concomitant silencing of LigA and LigB and effect upon serum resistance



Virulence assays in Hamster model of infection



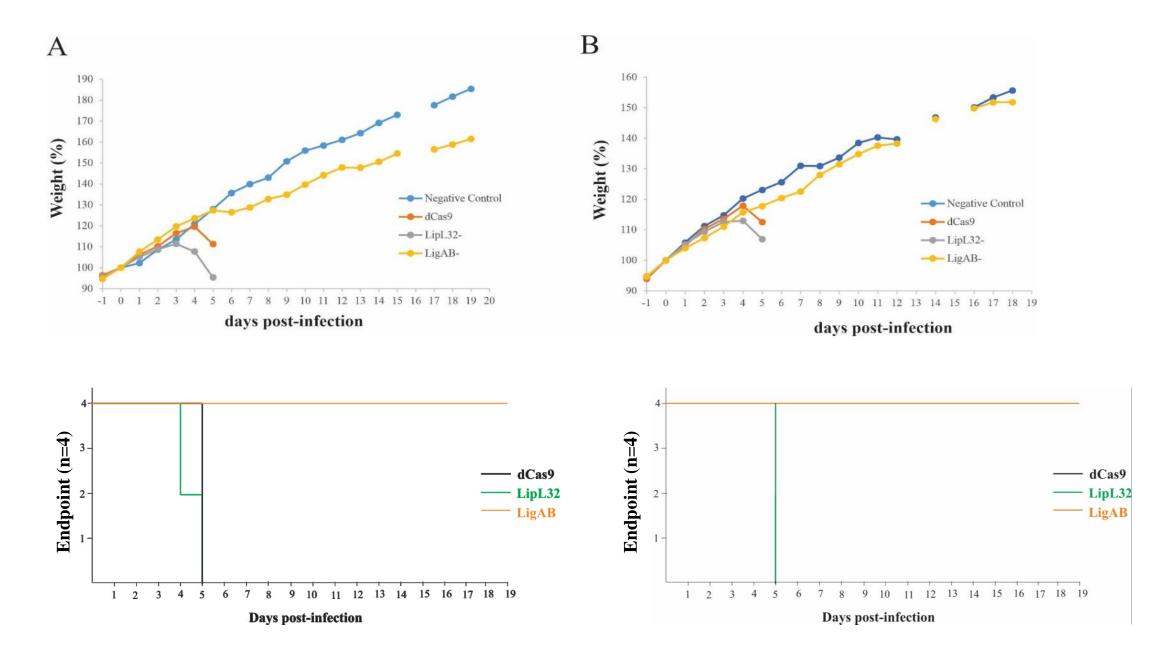
 10^{-2} 10^{-3}

Animals sacrificed at **endpoint:** Weight lost and clinical signs

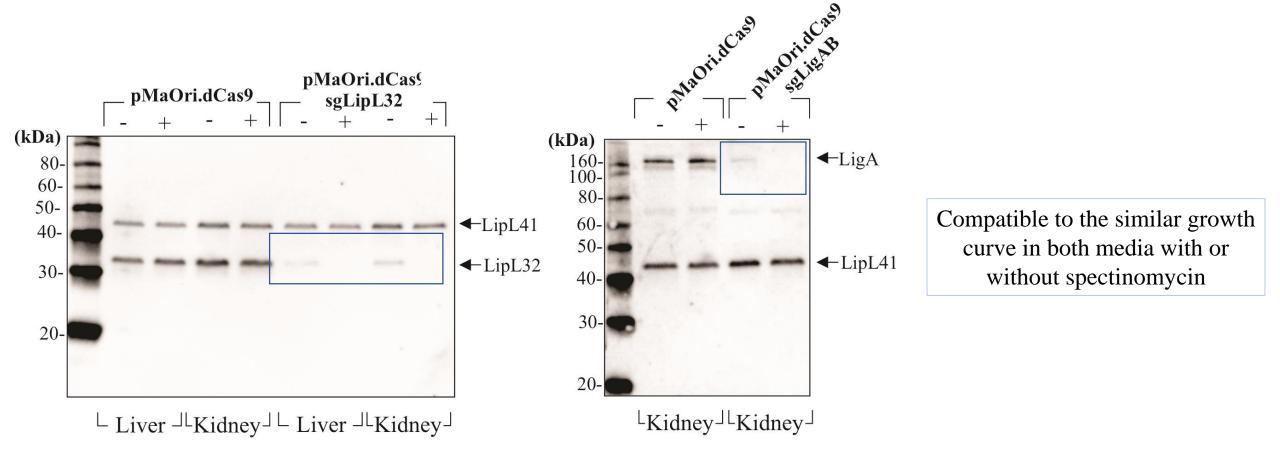
n=4

Recovered mutants were

Recovered mutants were used for another round of infection

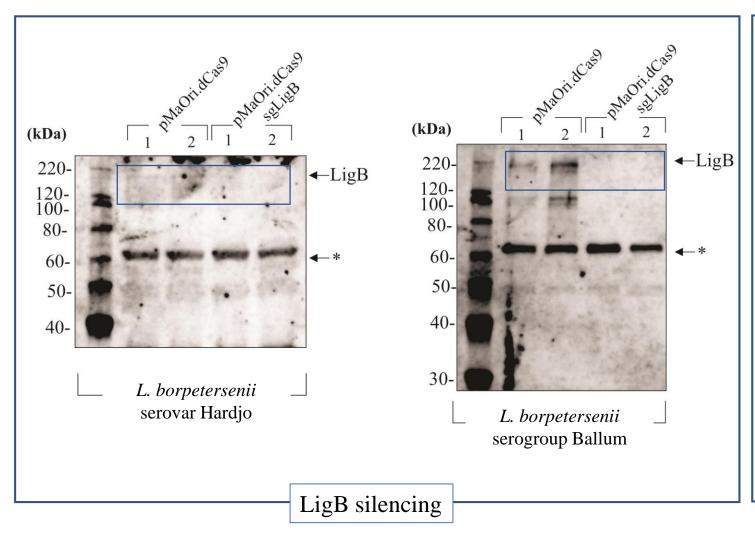


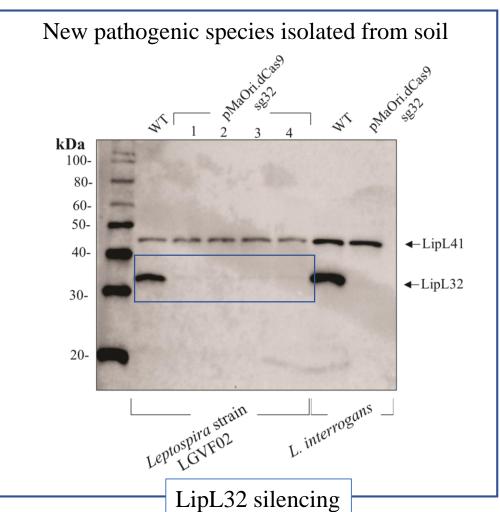
Detection of proteins in recovered leptospires



LigAB double mutante caused chronic disease

Application of CRISPRi to other *Leptospira* species





CONCLUSIONS

- CRISPRi was successfully applied to distinct pathogenic *Leptospira*
- Complete and stable gene silencing was observed, suitable for animal experiments
- The loss of virulence of the double LigA/LigB mutant was confirmed, probably due to reduced serum resistance

scientific reports

Check for updates

OPEN Genetic manipulation of pathogenic Leptospira: CRISPR interference (CRISPRi)-mediated gene silencing and rapid mutant recovery at 37 °C

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Vivarium small animal care team

THANK YOU FOR YOUR ATTENTION!



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