

In silico identification and chemical remodelling of tick protein epitopes for vaccine antigen development

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Stepan S. Denisov^{1,2}, M. Amine Jmel³, Wouter Peeters¹, Maya Idema¹, Michalis Kotsyfakis³, Johannes H. Ippel¹, Tilman M. Hackeng¹, Ingrid Dijkgraaf^{1*}

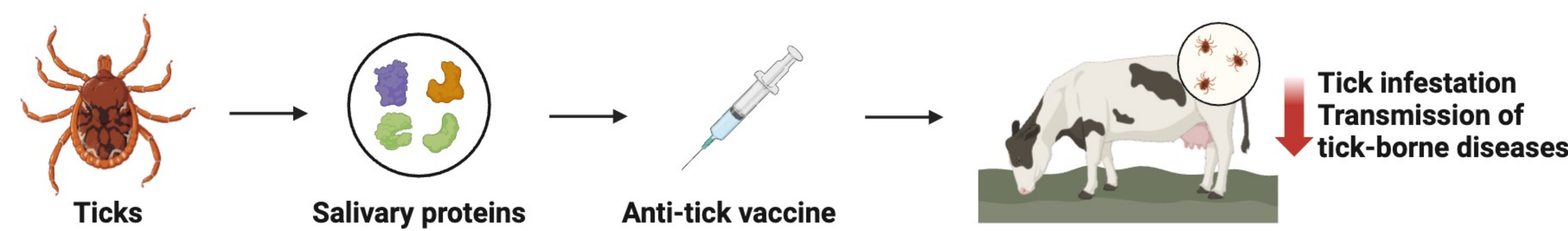
¹Department of Biochemistry, Maastricht University, Cardiovascular Research Institute Maastricht (CARIM), Maastricht, the Netherlands.

²Radcliffe Department of Cardiovascular Medicine, University of Oxford, the United Kingdom

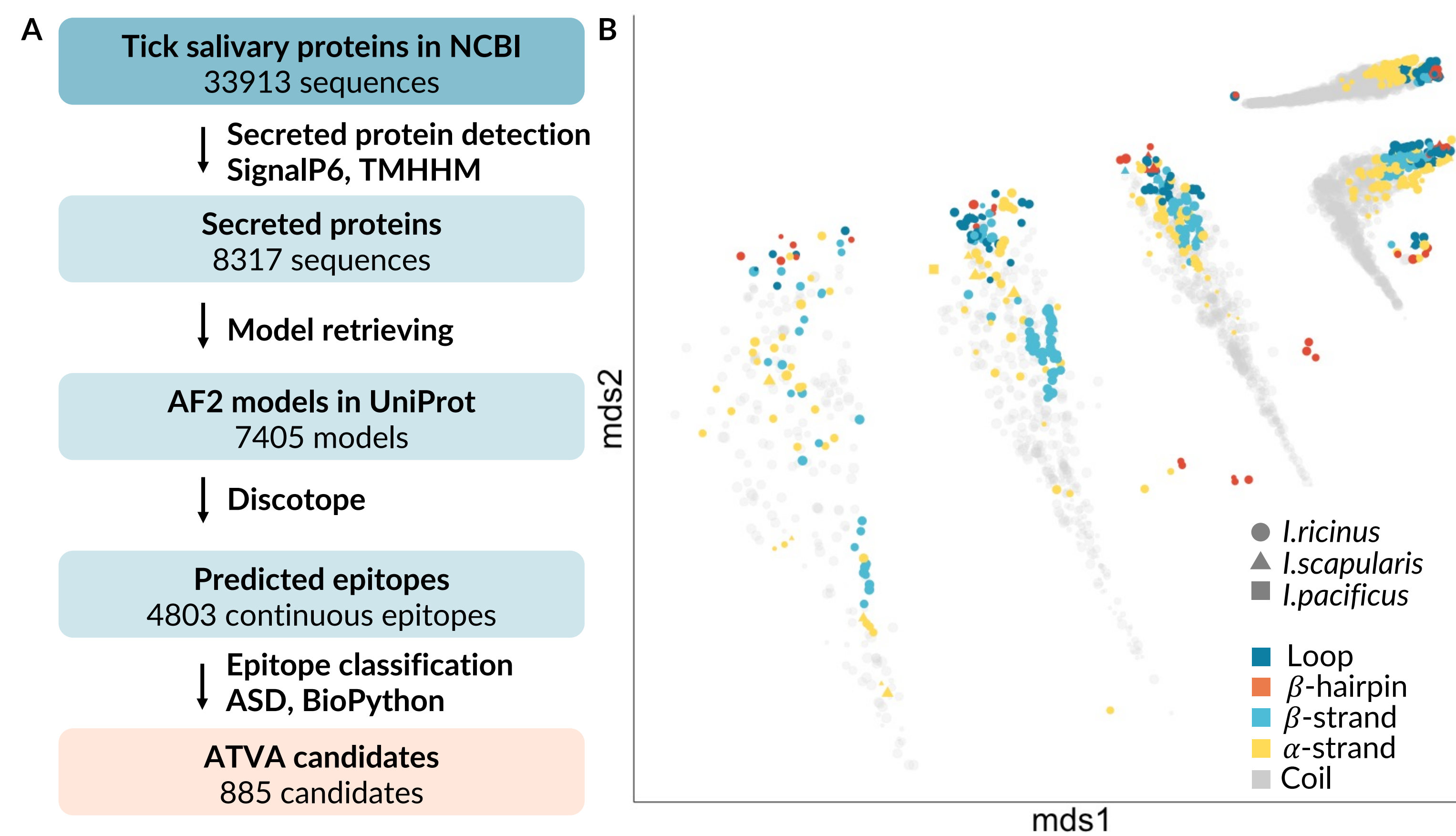
³Institute of Parasitology, Biology Centre, Czech Academy of Sciences, České Budějovice, Czech Republic

Introduction

Ticks and tick-borne diseases are severe burdens for healthcare systems and for animal husbandry amounting to billions of dollars in economic losses worldwide. Due to climate change, ticks' habitats are expanding, which makes the need for novel ways of tick control as pressing as never before. One of the desirable strategies is the development of anti-tick vaccines to elicit acquired tick resistance¹. The challenge in the development of such vaccines often lies in the inherently low immunogenicity and proteolytical stability of tick proteins, which they acquired through millions of years of evolution. Here we present a pipeline for the development of anti-tick vaccine antigens (ATVA) which utilizes AlphaFold2 structure modelling of tick proteins, *in silico* identification of antigenic epitopes by the protrusion-based algorithm², chemical remodelling and multimerization using the tick salivary lectin pathway inhibitor (TSLPI) as a model protein.



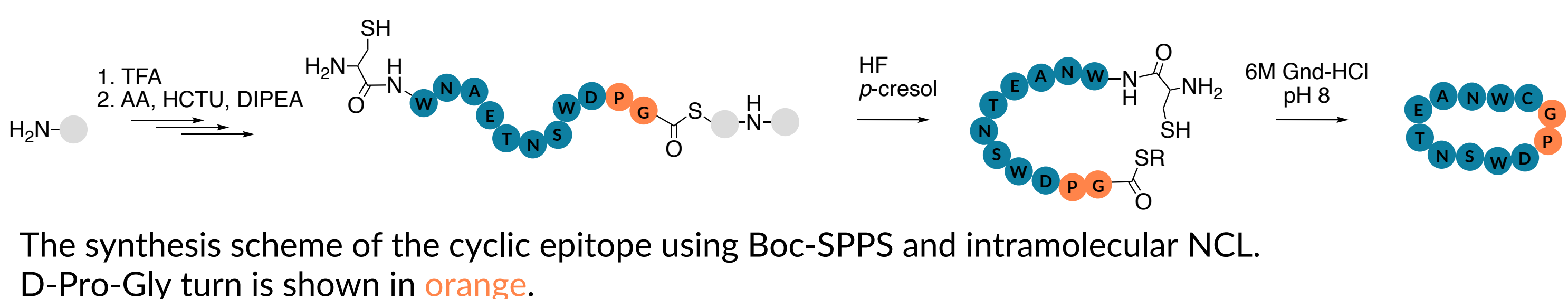
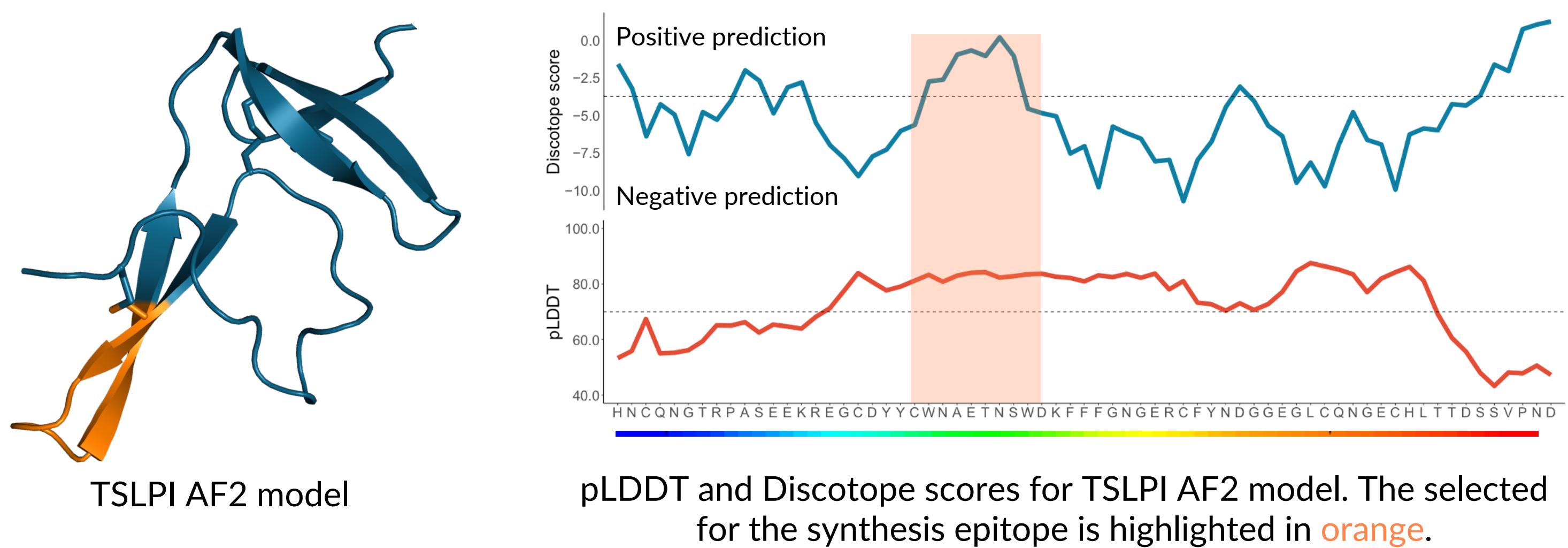
Bioinformatic pipeline allows mining of ATVA candidates



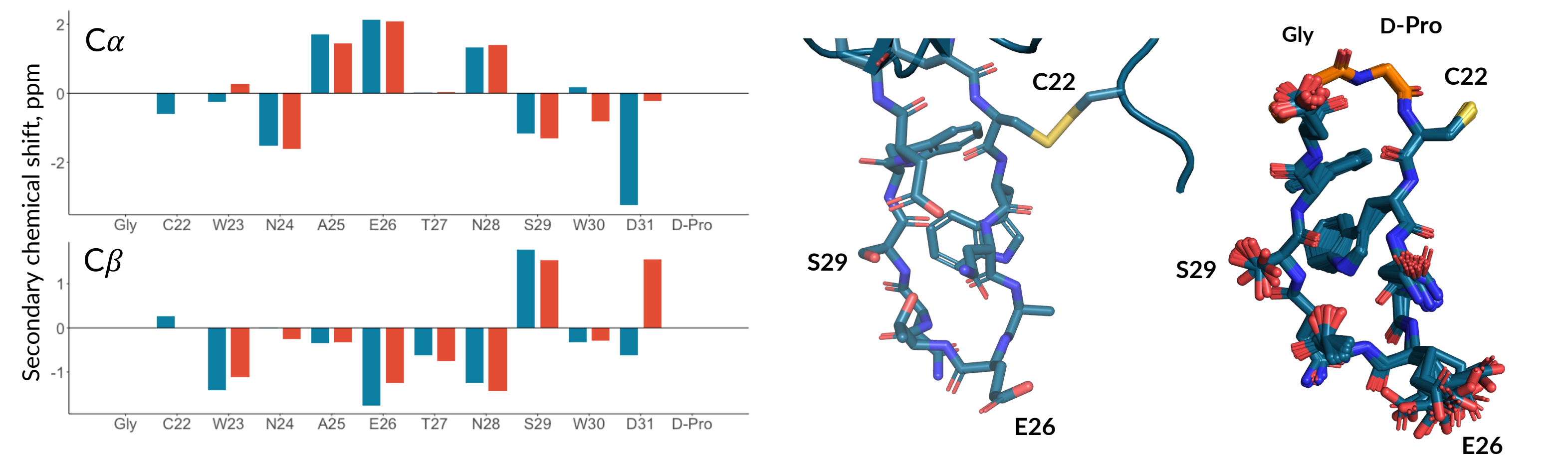
A. The pipeline for ATVA detection. B. The 2D representation of the epitope structural space based on amplitude spectrum distance (ASD)³.

TSLPI contains a prospective epitope for ATVA

Tick salivary Lectin Pathway Inhibitor (TSLPI) from *Ixodes Scapularis* blocks the lectin pathway and facilitates transmission of *Borrelia burgdorferi*⁴.

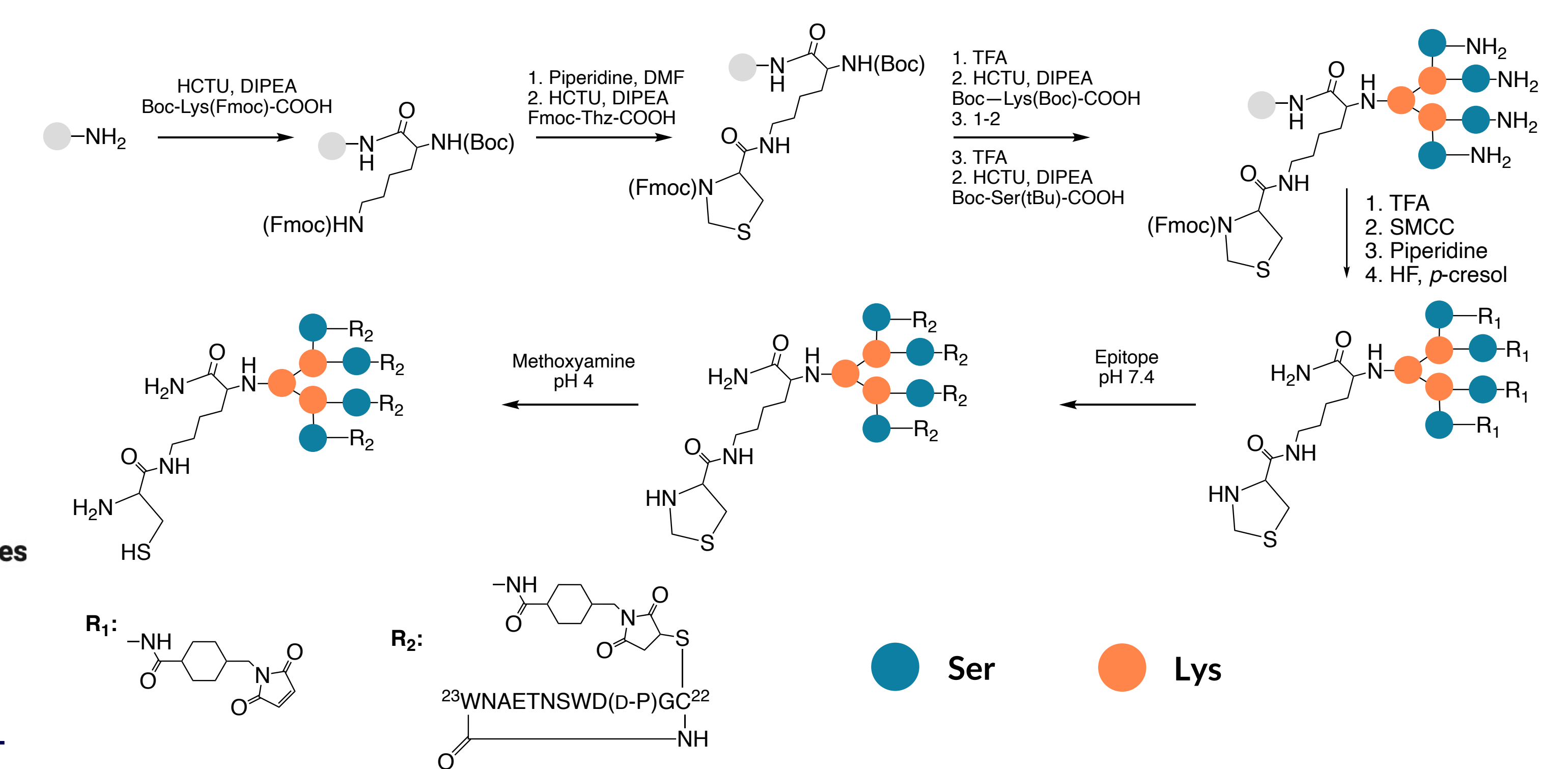


Synthetic epitope mimics the parental TSLPI



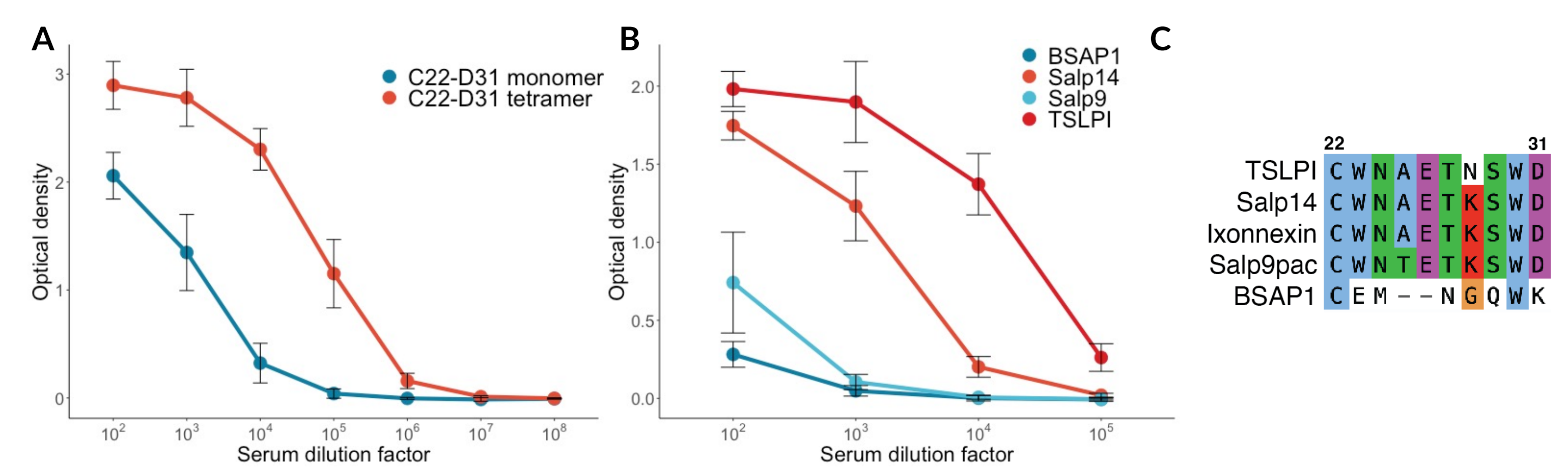
Comparison of secondary chemical shifts for full-length TSLPI and the cyclic epitope. C22-D31 region in the TSLPI AF2 model (left) and overlay of 20 lowest-energy NMR structures of the cyclic epitope (right). D-Pro-Gly turn is shown in orange.

Epitope is multimerized for higher immunogenicity

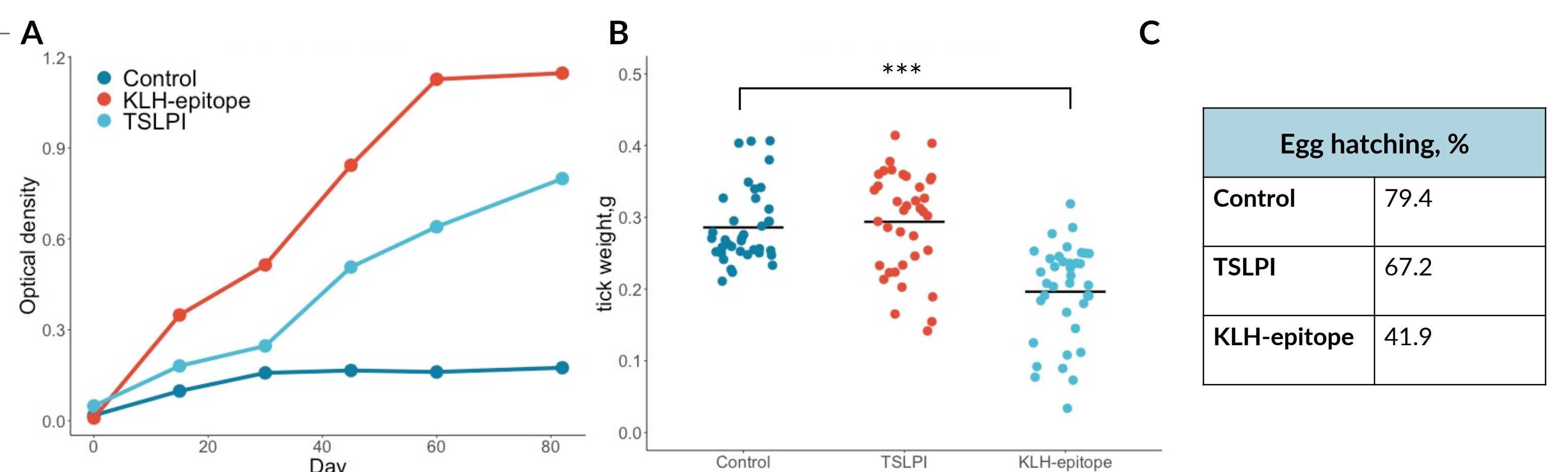


Synthesis of the tetrameric cyclic epitope coupled to a lysine wedge. SMCC, R₁: succinimidyl-4-(N-maleimidomethyl)cyclohexane-1-carboxylate

Synthetic epitopes elicit specific immune response



A. Level of TSLPI-specific antibodies in sera samples of mice vaccinated with KLH-peptide conjugates. B. Cross-reactivity of antibodies against tick proteins in sera samples of mice vaccinated with tetrameric cyclic epitope. C. Section of multiple sequence alignment related to TSLPI proteins from *I. scapularis*.



Results experiments with naïve rabbits (control), immunized with the parent TSLPI and tetrameric-cyclic C22-D31 TSLPI epitope (KLH-epitope) challenged with *I. ricinus* ticks. The level of antibodies in rabbit sera (A), tick weight (B) and egg hatching percentage (C).

Conclusions

- AlphaFold and structure-based epitope prediction algorithms allows fast identification of ATVB
- The predicted epitope from *I. Scapularis* was chemically synthesised and modified to structurally mimic the parental protein
- Tertramered TSLPI epitope elicits higher TSLPI-specific immune response in mice compared to the monomeric epitope
- Immunisation of rabbits with the tetrameric epitope might affect tick feeding and reproduction

References

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