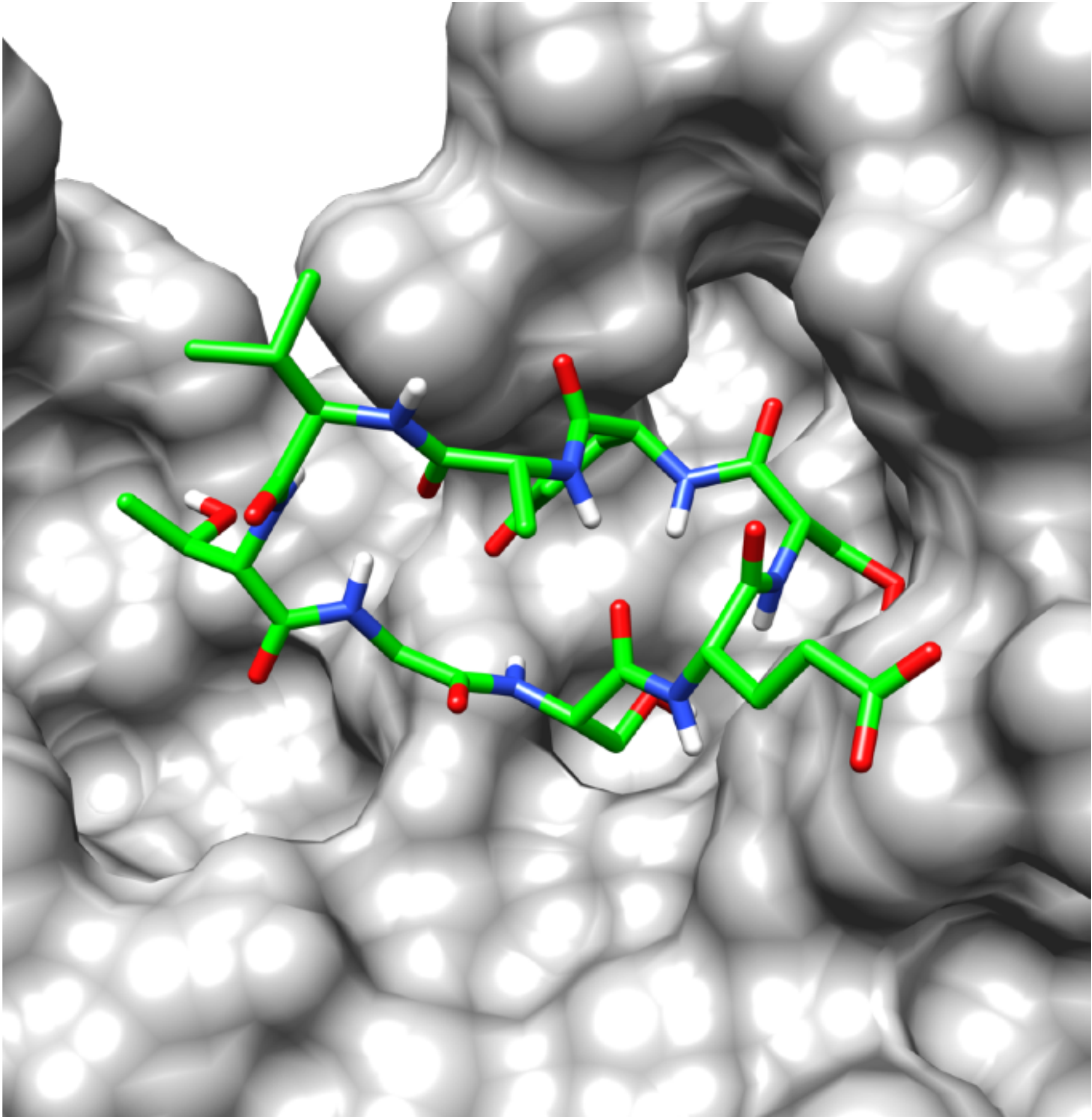
- Designed cyclic peptides that are predicted to be well structured.

- Structural predictions of cyclo-(SESEavTG) align well with NMR data.

- Cell-based studies show little inhibitory effect of cyclo-(SESEavTG) and cycle-(SESEGvvTG).

- Verify our inhibitor design and experimental methods.




cyclo-(SESEavTG)


Skp2

Conclusion and next steps

(SESEavTG) align well with NMR

inhibitory effect of cyclo-(SESEavTG)

experimental methods.

- Verify our inhibitor design and

and cycle-(SESEGvvTG).

- Structural predictions of cyclo-


data.

predicted to be well structured.

- Designed cyclic peptides that are

- Cell-based studies show little

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