

# STRUCTURAL BASIS OF LIPOPROTEIN SIGNAL PEPTIDASE II ACTION AND INHIBITION BY THE ANTIBIOTIC GLOBOMYCIN

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- 背景 (background)
- 方法 (methods) & 结果 (results)
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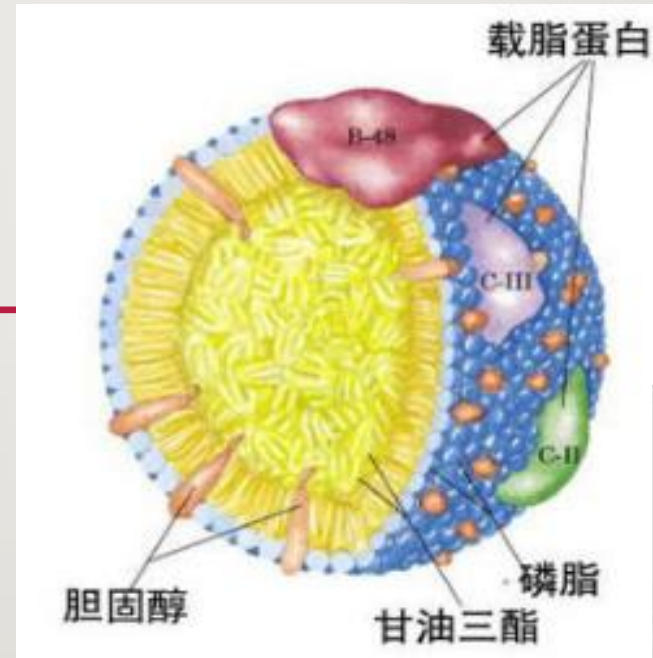
# PART I BACKGROUND

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# BACKGROUND

- Lipoproteins
- Globomycin
- Signal peptide
- Signal peptidase



# BACKGROUND

- The threat of antibiotic resistance is recognized as a serious public health issue.

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- With functions that range from cell envelope structure to signal transduction and transport lipoproteins constitute 2 to 3% of bacterial genomes and play critical roles in bacterial physiology, pathogenicity, and antibiotic resistance.
- LspA, as a key enzyme involved in the posttranslational processing of lipoproteins in *Pseudomonas aeruginosa*, an opportunistic human pathogen, is a target for antibiotic development



# PART II

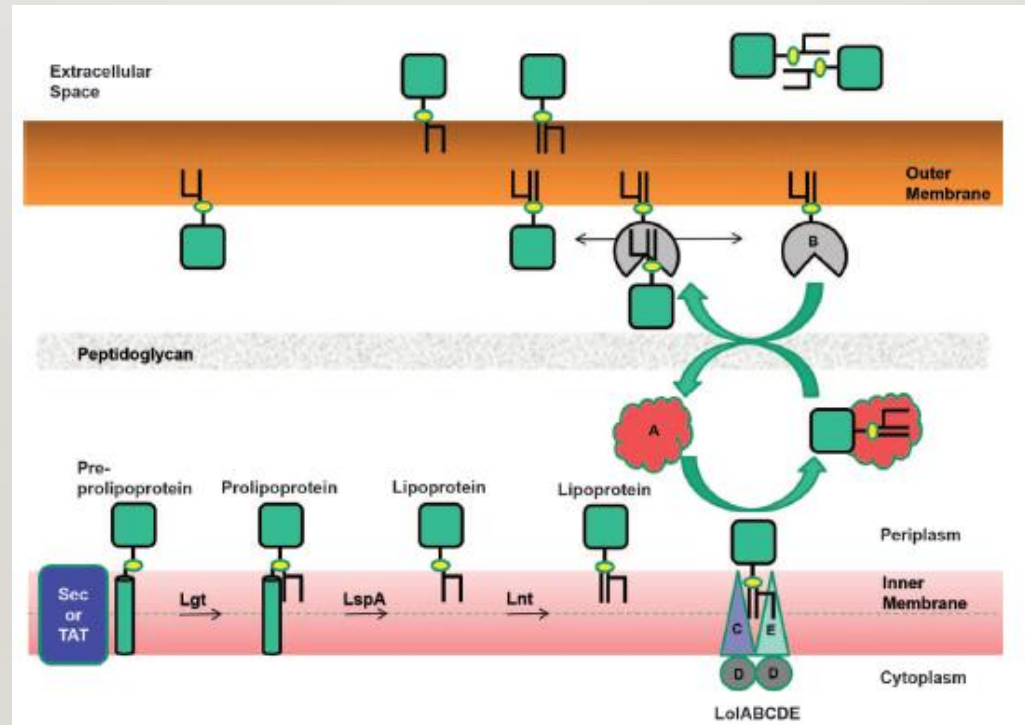
# METHODS & RESULTS

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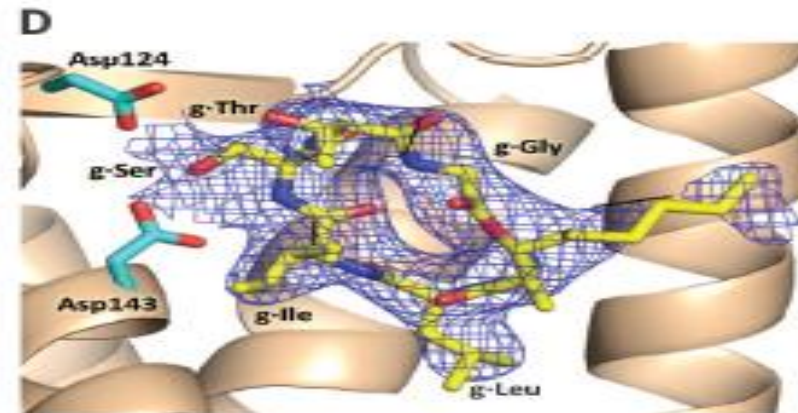
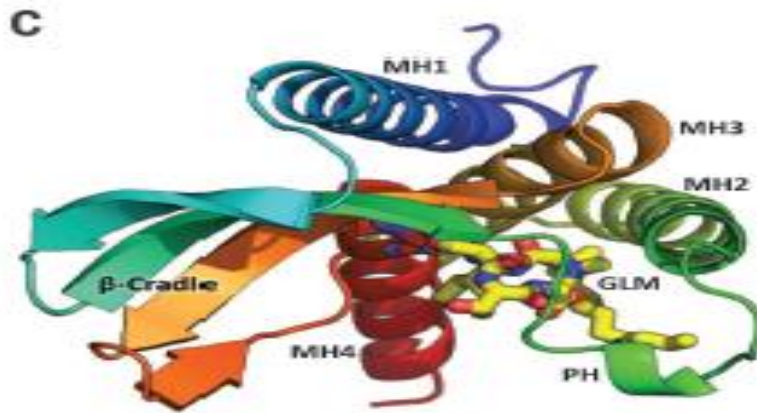
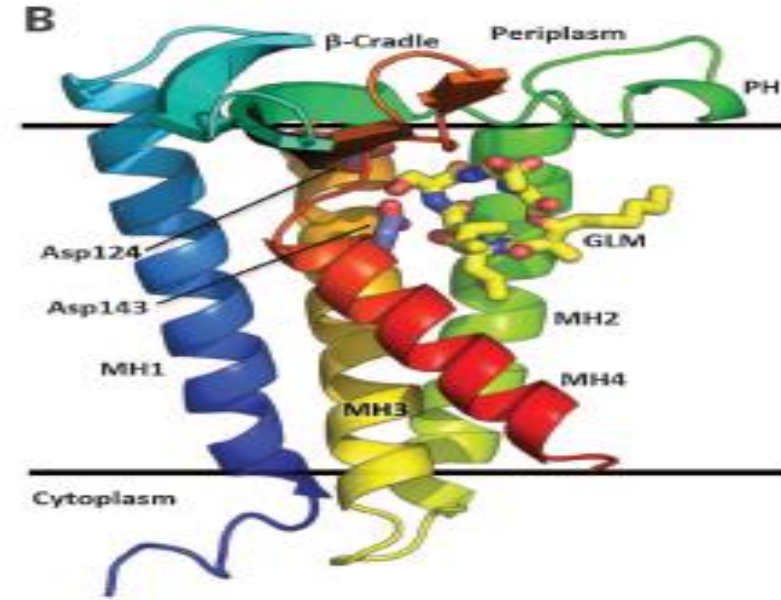
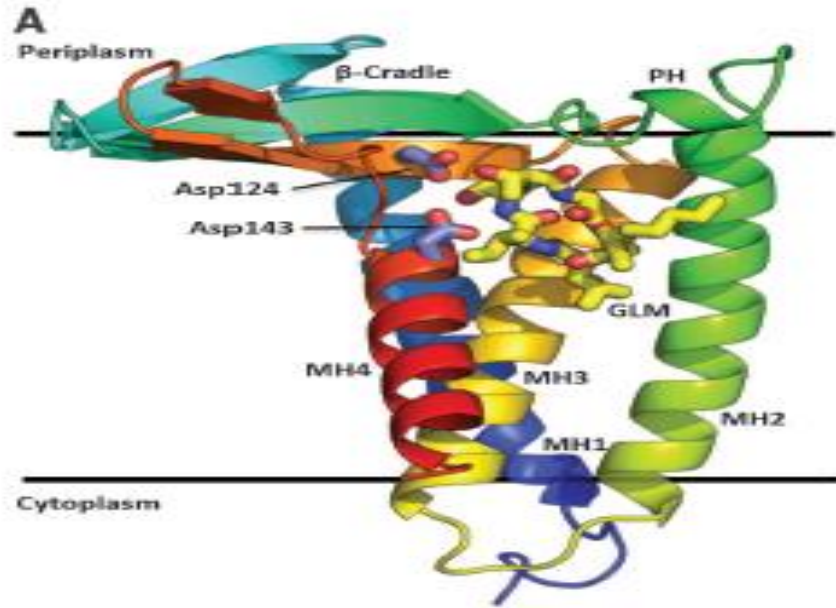
- the crystal structure ———Pseudomonas aeruginosa complexed with the antimicrobial globomycin at 2.8 angstrom resolution
- Crystallization trials—————obtain a crystal structure of LspA
- Mutagenesis studies—————identify LspA as an aspartyl peptidase
- molecular mimicry—————globomycin appears to inhibit by acting as a noncleavable peptide thatsterically blocks the active site.

# POSTTRANSLATIONAL PROCESSING LEADING TO LIPOPROTEIN MATURATION IN GRAM-NEGATIVE BACTERIA.

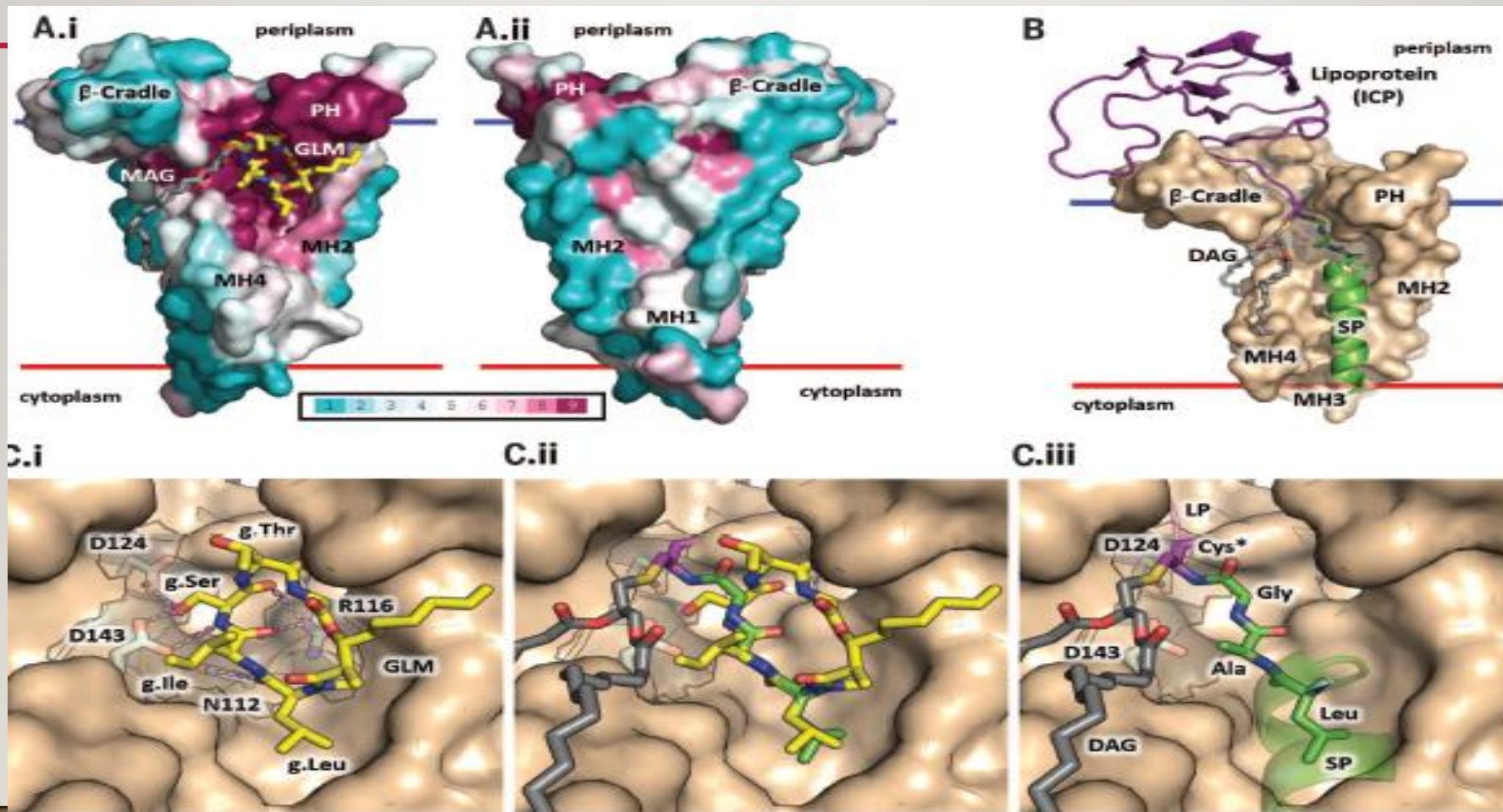




# LSPA-GLOBOMYCIN COMPLEX STRUCTURE.

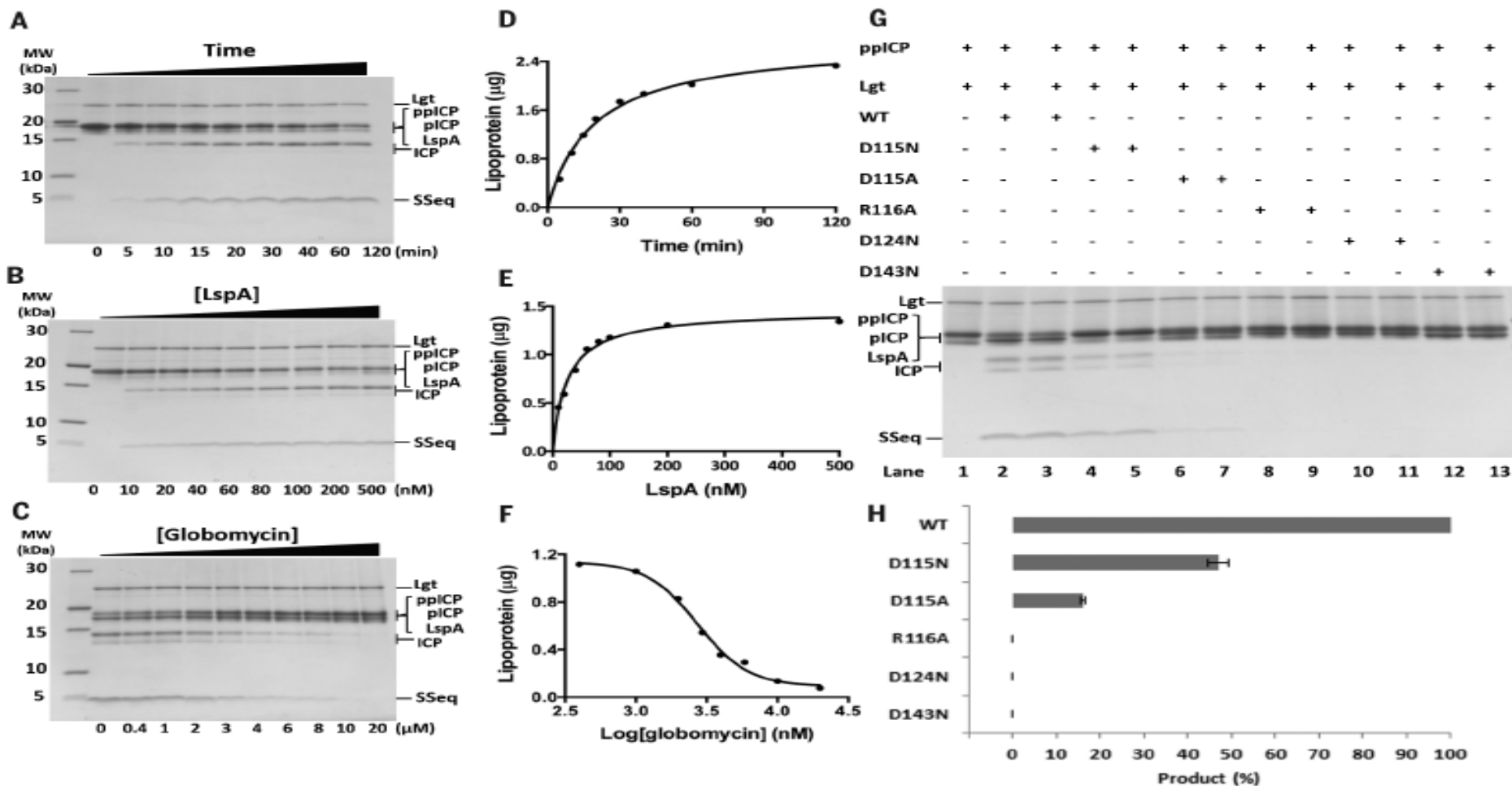


# CONSERVED RESIDUES IN AND GLOBOMYCIN AND PROLIPOPROTEIN BINDING TO LSPA.





# ENDOPEPTIDASE ACTIVITY OF LSPA WITH THE SUBSTRATE PROLIPOPROTEIN, GENERATED IN SITU, BASED ON THE PREPROLIPOPROTEIN ICP.



# PART III

# CONCLUSIONS

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# CONCLUSION

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- LspA is an aspartyl endopeptidase, it is a suitable target with which to explore the thousands of inhibitors developed for other medically relevant aspartyl proteases
- This structure should inform rational antibiotic drug discovery.



# PART IV

# SUMMARIES

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主要结论：

**LSPA** 是天冬氨酸肽酶，可作为抗生素发展的靶标，是常规的抗生素类药物发现过程需要了解的结构。

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- 启发（设计实验思路，方法）
- 实验设计合理（**structure-properties**通过探索结构研究性质）
- 实验方法多样（**MDS, MM, MS**）
- 结果分析清晰（一个实验一个结果，逻辑清晰）
  
- 改进：
- 实验细节省略过多，不利于别人进行重复。