

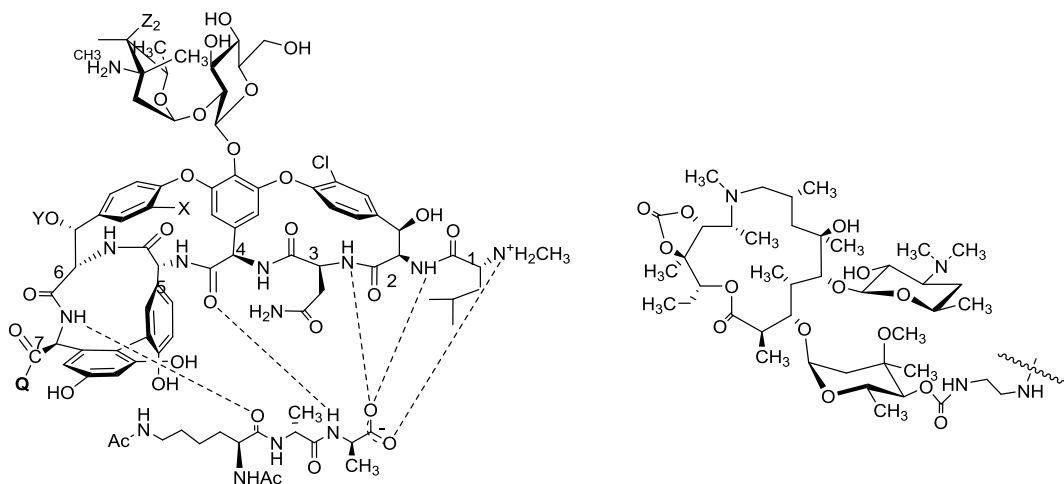
The study of the correlation of the binding energy of peptide ligand complexes with hybrid antibiotics vancomycin-azithromycin and eremomycin-azithromycin with antibacterial activity

Bykov E.E., Mirchink E.P., Isakova E.B., Bychkova E.N., Olsufyeva E.N., Tevyashova A.N.

Gause Institute of New Antibiotics, B. Pirogovskaya str., 11, Moscow 119021, Russia e-mail: evgenbykov@yandex.ru

One of modern strategies that can solve the problem of antibacterial resistance is the development of dual-acting hybrid antibiotics – structures that contain two covalently linked antimicrobial drugs that interact with different targets in a bacterial cell [1].

Antibacterial activity of hybrid antibiotics vancomycin-azithromycin (C11, C12-carbonate) and eremomycin-azithromycin (C11, C12-carbonate) was evaluated. Quantum chemical calculations of energy these hybrid antibiotics with a model tripeptide ligand α, ϵ -di-Ac-L-Lys-D-Ala-D-Ala by the semiempirical PM6 method using a software package Spartan-10[2] provided data on geometrical parameters of these complexes along with the energy of their formation and the influence of protonation of the NHCH_3 group. A correlation between the energy of formation of antibiotics-ligand complexes and antibacterial activity of hybrid antibiotics against Gram-positive bacterial strains was found



A) α, ϵ -di-Ac-L-Lys-D-Ala-D-Ala (LAA-)

Vancomycin⁺ + LAA- (1a)

R=OH; X=Cl; Y=H;
Y=эремозаминил- α
Z₁=OH; Z₂=H; Q=OH

Eremomycin⁺ + LAA- (2a)

R=OH; X=H;
Z₁=H; Z₂=OH; Q=OH

Vanco-azithro⁺ + LAA- (5a) Q =

Eremo-azithro⁺ + LAA- (6a) Q =

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References

[1] A N Tevyashova, E N Olsufyeva, M N Preobrazhenskaya, "Design of dual action antibiotics as an approach to search for new promising drugs", *RUSS CHEM REV*, 2015, **84** (1), 61–97

[2] <https://www.wavefun.com/>