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Vede seminář: Prof. RNDr. Michal Otyepka, Ph.D.

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- **2005** V. Schramke et al. "Retraction: RNA-interference-directed chromatin modification coupled to RNA polymerase II transcription" in *Nature* (volume 437, page 1057). Irreproducible results.
- **2005** R. C. Allshire. "Retraction. Hairpin RNAs and retrotransposon LTRs effect RNAi and chromatin-based gene silencing" in *Science* (volume 310, page 49). Irreproducible results.













Retracted papers (wiki)

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Information for Authors of Papers

(Revised January 2011)

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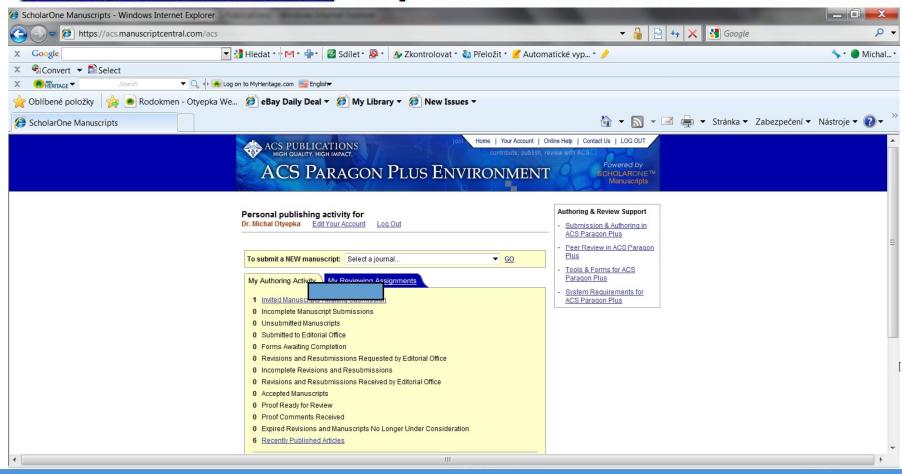




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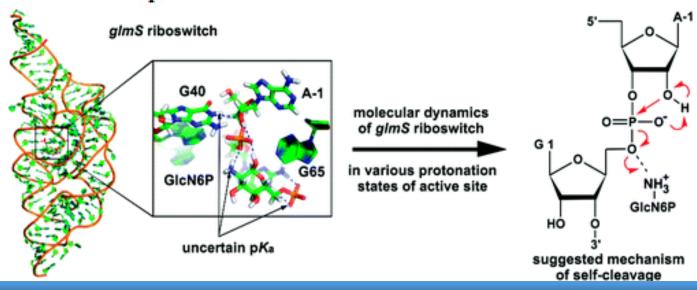




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Ukázka

J. Phys. Chem. B 2010, 114, 8701-8712

8701

Protonation States of the Key Active Site Residues and Structural Dynamics of the glmS Riboswitch As Revealed by Molecular Dynamics

Pavel Banáš, † Nils G. Walter, Jiří Šponer, * † and Michal Otyepka * † †

Department of Physical Chemistry, Faculty of Science, Palacky University, 17. Listopadu 12, 771 46 Olomouc, Czech Republic, Institute of Biophysics, Academy of Sciences of the Czech Republic, Kralovopolska 135, 612 65 Brno, Czech Republic, and Single Molecule Analysis Group, Department of Chemistry, University of Michigan, 930 North University Avenue, Ann Arbor, Michigan 48109-1055

Received: November 18, 2009; Revised Manuscript Received: March 29, 2010

The glmS catalytic riboswitch is part of the 5'-untranslated region of mRNAs encoding glucosamine-6-phosphate (GlcN6P) synthetase (glmS) in numerous Gram-positive bacteria. Binding of the cofactor GlcN6P induces site-specific self-cleavage of the RNA. However, the detailed reaction mechanism as well as the protonation state of the glmS reactive form still remains elusive. To probe the dominant protonation states of key active site residues, we carried out explicit solvent molecular dynamic simulations involving various protonation states of three crucial active site moieties observed in the available crystal structures: (i) guanine G40 (following the Thermoanaerobacter tengcongensis numbering), (ii) the GlcN6P amino/ammonium group, and (iii) the GlcN6P phosphate moiety. We found that a deprotonated G40⁻ seems incompatible with the observed glmS active site architecture. Our data suggest that the canonical form of G40 plays a structural role by stabilizing an in-line attack conformation of the cleavage site A-1(2'-OH) nucleophile, rather than a more direct chemical role. In addition, we observe weakened cofactor binding upon protonation of the GlcN6P phosphate moiety, which explains the experimentally observed increase in K_m with decreasing pH. Finally, we discuss a possible role of cofactor binding and its interaction with the G65 and G1 purines in structural stabilization of the A-1(2'-OH) in-line attack conformation. On the basis of the identified dominant protonation state of the reaction precursor, we propose a hypothesis of the self-cleavage mechanism in which A-1(2'-OH) is activated as a nucleophile by the G1(pro-R₀) nonbridging oxygen of the scissile phosphate, whereas the ammonium group of GlcN6P acts as the general acid protonating the G1(O5') leaving group.

Introduction

Riboswitches are RNA motifs embedded in messenger RNAs (mRNAs) that regulate gene expression in response to binding 5'-OH termini of the reaction products. The same general mechanism is found in all ribozymes classified as "small" with typically less than 100 nucleotides in the catalytic core, yet the details of how the reaction participants are activated differ.













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Ukázka

protonation states of the essential residues in the AS. MD is suitable to suggest the protonation states of the critical nucleobases corresponding to the crystalline conditions. This was previously shown, for example, for protonated cytosines in a HDV ribozyme and a frameshifting pseudoknot. 50,55 The aim of our simulations is to suggest dominant protonation states of AS residues, to describe the cofactor binding, and to study the dynamic behavior of AS to get ideas about the plausible reaction state. In particular, our data indicate that G40 is not deprotonated and likely plays a structural role in stabilizing the in-line attack conformation of the cleavage site A-1(2'-OH). We propose that A-1(2'-OH) could be activated as the nucleophile by the G1(pro-R₀) nonbridging oxygen of the scissile phosphate, whereas the ammonium group of GlcN6P acts as the general acid that neutralizes the leaving group 5'-oxygen (to avoid any confusion, we will further use terms pro-So for O1P and pro-Ro for O2P nonbridging oxygens according to IUPAC terminology). The presented results should be considered within the context of common limitations (mentioned above) of the contemporary simulation methods.

Methods

Preparation of Starting Structures. The starting geometries













Ukázka

Acknowledgment. This study was supported by Grants LC512, LC06030, and MSM6198959216 from the Ministry of Education of the Czech Republic, Grants 203/09/1476 and 203/09/H046 from the Grant Agency of the Czech Republic, Grants IAA400040802 and 1QS500040581 from the Grant Agency of the Academy of Sciences of the Czech Republic, Grants AV0Z50040507 and AV0Z50040702 from the Academy of Sciences of the Czech Republic, and NIH Grant GM62357 (to N.G.W.). We thank S. R. Das for helpful comments and discussions.

Supporting Information Available: The content of the Supporting Information includes force field parameters of nonstandard residues, a detailed analysis of the structural dynamic of the glmS riboswitch active site, details of the structural dynamics of the glmS riboswitch without cofactor, analysis of constant-pH MD simulations using implicit solvent methods, and some other material. This material is available free of charge via the Internet at http://pubs.acs.org.

References and Notes

- Mandal, M.; Boese, B.; Barrick, J. E.; Winkler, W. C.; Breaker, R. R. Cell 2003, 113, 577.
- (2) Tucker, B. J.; Breaker, R. R. Curr. Opin. Struct. Biol. 2005, 15, 342.
- (3) Winkler, W. C.; Breaker, R. R. Annu. Rev. Microbiol. 2005, 59, 487













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 nikdy citovány!







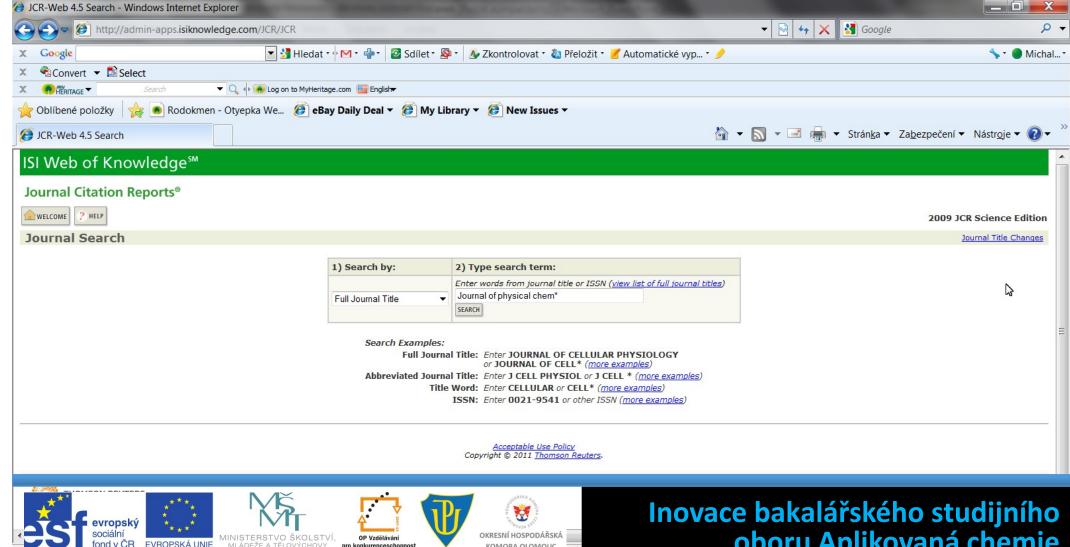




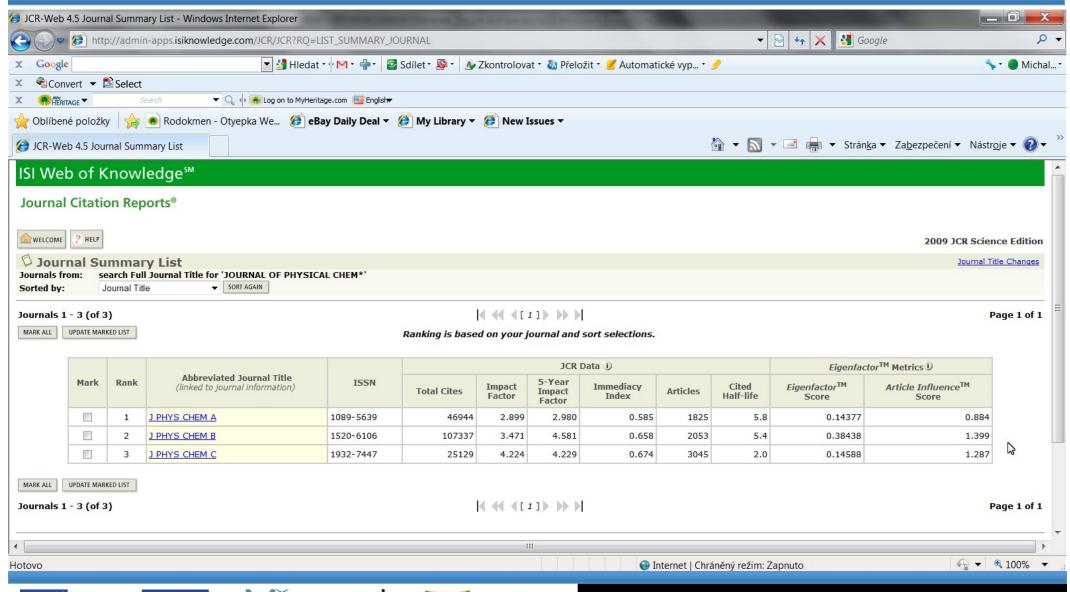


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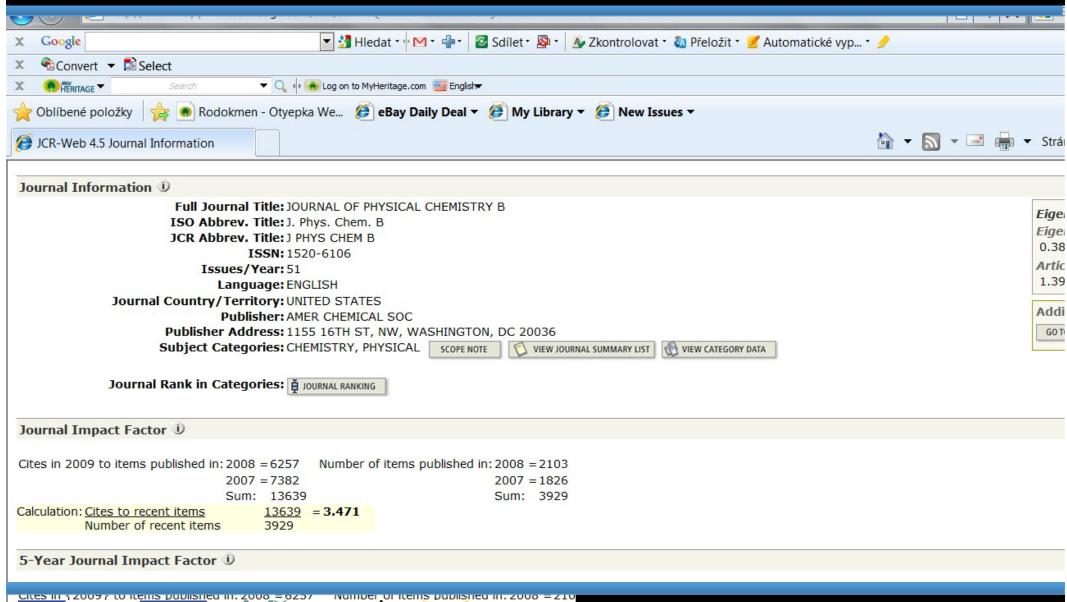






Inovace bakalářského studijního oboru Aplikovaná chemie

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