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Адрес

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Образование

1993–1997	Москва	аспирантура ИБХ РАН
1987–1993	Долгопрудный	ФФХБ МФТИ

Работа

2002–наст.вр.	Москва	ИБХ РАН	старший научный сотрудник
1999–2002	Москва	ИБХ РАН	научный сотрудник

Работа в ИБХ

2024–наст.вр.	Главный научный сотрудник
2002–2024	Старший научный сотрудник

Навыки

Методы гетероядерной ЯМР-спектроскопии высокого разрешения и молекулярного моделирования для исследования структуры и динамики низкомолекулярных соединений и биологических макромолекул.

Владение языками

русский, английский, немецкий

Научные интересы

Научные интересы относятся к области структурно-динамических исследований белков и их комплексов с биомолекулами (липидами, ДНК, РНК и т.д.) методами гетероядерной ЯМР-спектроскопии высокого разрешения в интеграции с другими методами структурной биологии и биофизики. Разработка новых методик ЯМР-спектроскопии для изучения подвижных белковых комплексов.

Основные работы посвящены исследованиям молекулярных механизмов биологической активности мембранных и мембраноактивных белков в норме и при патогенезе различных заболеваний человека (в том числе, онкологических и нейродегенеративных). В последние годы основные исследования направлены на установление биофизических принципов проведения сигнала рецепторами через мембрану клетки, а также функционирования мембранных белков в макромолекулярных комплексах. Особое внимание уделяется изучению аллостерических конформационных перестроек, белок-белковых и белок-липидных взаимодействий компонент цитохром Р450-зависимых систем, рецепторных тирозинкиназ (РТК) и РТК-подобных рецепторов, белка-предшественника β -амилоида (мутации в которых связаны с онкогенезом и болезнью Альцгеймера) для разработки перспективных лекарственных соединений. В то же время, часть работ направлена на исследования структурно-динамических свойств растворимых белков и их комплексов, в том числе рибосомных белков, цитокинов, цитохромов, гистонотипных НУ-белков из патогенных бактерий, пептидных и белковых токсинов из растений и животных.

Реализация научных Проектов в качестве Руководителя:

- 2001 г., РФФИ 01-04-48651-а и 01-04-06130-мас по теме «Пространственная структура и динамика рибосомного белка L7/L12 из *Escherichia coli* в растворе»;
- 2001-2002 г.г., РФФИ 01-04-06130-мас и 02-04-06650-мас «Исследование фемто-микросекундной внутримолекулярной динамики белков с использованием новых методов гетероядерной спектроскопии ЯМР»;
- 2003 г., РФФИ 03-04-06408-мас «Исследование взаимосвязи пространственной структуры и внутримолекулярной подвижности биологически активных полипептидов с их функцией»;
- 2003-2005 г.г., РФФИ 03-04-49011-а «Учет конформационной подвижности при описании электростатических взаимодействий в белках с помощью новых методов гетероядерной спектроскопии ЯМР и молекулярной динамики»;
- 2006-2008 г.г., РФФИ 06-04-49740-а «Структурно-динамические исследования трансмембранных доменов рецепторных протеинкиназ современными методами гетероядерной спектроскопии ЯМР и молекулярной динамики»;
- 2009-2011 г.г., РФФИ 09-04-00551-а «Структурно-динамический анализ специфических взаимодействий трансмембранных доменов рецепторных тирозинкиназ в норме и патологии»;
- 2012-2014 г.г., РФФИ 12-04-01816-а «Структурные, кинетические и термодинамические аспекты специфических взаимодействий трансмембранных доменов рецепторных тирозинкиназ в норме и патологии»;
- 2015-2017 г.г., РФФИ 15-04-07983-а «Установление молекулярных механизмов влияния патогенных мутаций в трансмембранных доменах на активность рецепторных тирозинкиназ»;
- 2018-2020 г.г., РФФИ 18-04-01289-а «Структурные аспекты воздействия онкогенных мутаций в трансмембранных доменах рецепторных тирозинкиназ на проведение сигнала через мембрану клетки»;
- 2020-2022 г.г., РФФИ 20-54-00041-Бел-а «Анализ взаимодействия мембран-связанных компонентов цитохром Р450-зависимых систем человека с использованием методов интегративной структурной биологии»;
- 2020-2022 г.г., РФФИ 20-64-46027 «Структурно-динамические исследования внутри- и межмолекулярных взаимодействий амилоидогенных пептидов и их предшественников для установления молекулярных механизмов патогенеза и разработки таргетных методов терапии нейродегенеративных заболеваний»;
- 2023-2025 г.г., РФФИ 23-44-10021 (международный, совместно с БРФФИ) «Конформационная динамика в образовании и функционировании биомакромолекулярных комплексов для биомедицинских и фармацевтических разработок».

Членство в сообществах

Российское общество Биохимиков и Молекулярных Биологов, Biophysical society, FEBS.

Степени и звания

2024	Доктор наук (Физико-математические науки, 1.5.2. — Биофизика)
1998	Кандидат наук (Химические науки, 02.00.10 — Биоорганическая химия)

Ссылки и контакты

ResearcherID: [R-5231-2016](#), Scopus: [7004085574](#), ORCID: [0000-0002-3635-1609](#), [Google Scholar](#)

Гранты и проекты

2024– наст.вр.	Онкотераностика и проблемы резистентности к противоопухолевым и антибактериальным препаратам
2011– 2014	Структурно-динамические модели белков высокой точности, верифицированные при помощи расширенного массива данных ЯМР-спектроскопии

2023– наст.вр.	Конформационная динамика в образовании и функционировании биомакромолекулярных комплексов для биомедицинских и фармацевтических разработок
2020– 2023	Бел: Анализ взаимодействия мембран-связанных компонентов цитохром P450-зависимых систем человека с использованием методов интегративной структурной биологии
2018– 2020	Структурные аспекты воздействия онкогенных мутаций в трансмембранных доменах рецепторных тирозинкиназ на проведение сигнала через мембрану клетки
2019– 2022	Структурная биология мембранных белков для создания новых лекарственных и диагностических средств
2014– 2018	Структурные основы молекулярных механизмов передачи сигнала интегральными мембранными белками I типа

Публикации

1. Lushpa VA, Lin C, Talyzina IA, Goncharuk MV, **Bocharov EV**, Arseniev AS, Wang X, Goncharuk SA, Mineev KS (2025). The intracellular domain of TLR2 is capable of high-affinity Zn binding: possible outcomes for the receptor activation. *FEBS Lett* , , [10.1002/1873-3468.70026](#)
2. Oparin P, Khokhlova O, Cherkashin A, Nadezhdin K, Palikov V, Palikova Y, Korolkova Y, Mosharova I, Rogachevskaja O, Baranov M, Shaidullova K, Ermakova E, Lushpa V, Bruter A, Deykin A, Ivanova E, Silaeva Y, Dyachenko I, **Bocharov E**, Sitdikova G, Andreev-Andrievskiy A, Poteryaev D, Shuster A, Murashev A, Kolesnikov S, Stepanenko V, Grishin E, Vassilevski AA (2025). Potent painkiller from spider venom antagonizes P2X3 receptors without dysgeusia. *Mol Ther* 33 (2), 771–785, [10.1016/j.ymthe.2024.12.036](#)
3. Volynsky PE, Urban AS, Pavlov KV, Bershatsky YV, Bocharova OV, Kryuchkova AK, Zlobina VV, Gavrilenkova AA, Dolotova SM, Kamynina AV, Zangieva OT, Taldaev A, Batishchev OV, Okhrimenko IS, Rakitina TV, Efremov RG, **Bocharov EV** (2025). Diverse Interactions of Sterols with Amyloid Precursor Protein Transmembrane Domain Can Shift Distribution Between Alternative Amyloid- β Production Cascades in Manner Dependent on Local Lipid Environment. *Int J Mol Sci* 26 (2), 553, [10.3390/ijms26020553](#)
4. Bedanokova DR, Goncharuk MV, Shabalkina AV, Lushpa VA, Arseniev AS, **Bocharov EV**, Mineev KS, Goncharuk SA (2024). Production and Refolding of the Ligand-Binding Domain of TrkA Receptor with the Extracellular Juxtamembrane Region. *Russ. J. Bioorganic Chem.* 50 (6), 2589–2595, [10.1134/S1068162024060232](#)
5. Motov VV, Kot EF, Kislova SO, **Bocharov EV**, Arseniev AS, Boldyrev IA, Goncharuk SA, Mineev KS (2024). On the Properties of Styrene–Maleic Acid Copolymer–Lipid Nanoparticles: A Solution NMR Perspective. *Polymers (Basel)* 16 (21), 3009, [10.3390/polym16213009](#)
6. Konshina AG, **Bocharov EV**, Konovalova EV, Schulga AA, Tolmachev V, Deyev SM, Efremov RG (2024). Structural Basis of Activity of HER2-Targeting Construct Composed of DARPIn G3 and Albumin-Binding Domains. *Int J Mol Sci* 25 (21), 11370, [10.3390/ijms252111370](#)
7. Petoukhov MV, Rakitina TV, Agapova YK, Petrenko DE, Podshivalov DD, Timofeev VI, Peters GS, Gaponov YA, **Bocharov EV**, Shtykova EV (2024). Molecular Dynamics and Small-Angle X-ray Scattering: Comparison of Computational and Experimental Approaches to Studying Structures of Biological Complexes. *Cryst. Rep* 69 (5), 674–681, [10.1134/S1063774524601540](#)
8. Lushpa VA, Goncharuk MV, Talyzina IA, Arseniev AS, **Bocharov EV**, Mineev KS, Goncharuk SA (2024). TIR domains of TLR family-from the cell culture to the protein sample for structural studies. *PLoS One* 19 (7), e0304997, [10.1371/journal.pone.0304997](#)
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10. Smirnova EV, Timofeev VI, Rakitina TV, Petrenko DE, Elmeeva OS, Saratov GA, Kudriaeva AA, **Bocharov EV**, Belogurov AA (2024). Myelin Basic Protein Attenuates Furin-Mediated Bri2 Cleavage and Postpones Its Membrane Trafficking. *Int J Mol Sci* 25 (5), 2608, [10.3390/ijms25052608](#)
11. Pham DD, Pham TH, Bui TH, Britikova EV, Britikov VV, **Bocharov EV**, Usanov SA, Phan VC, Le TBT (2024).

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21. Britikov VV, **Bocharov EV**, Britikova EV, Dergousova NI, Kulikova OG, Solovieva AY, Shipkov NS, Varfolomeeva LA, Tikhonova TV, Timofeev VI, Shtykova EV, Altukhov DA, Usanov SA, Arseniev AS, Rakitina TV, Popov VO (2022). Unusual Cytochrome c552 from *Thioalkalivibrio paradoxus*: Solution NMR Structure and Interaction with Thiocyanate Dehydrogenase. *Int J Mol Sci* 23 (17), , [10.3390/ijms23179969](https://doi.org/10.3390/ijms23179969)
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27. Bozin TN, Chukhontseva KN, Lesovoy DM, Filatov VV, Kozlovskiy VI, Demidyuk IV, **Bocharov EV** (2021). NMR assignments and secondary structure distribution of emfourin, a novel proteinaceous protease inhibitor. *Biomol NMR Assign* 15 (2), 361–366, [10.1007/s12104-021-10030-x](https://doi.org/10.1007/s12104-021-10030-x)
28. Urban AS, Pavlov KV, Kamynina AV, Okhrimenko IS, Arseniev AS, **Bocharov EV** (2021). Structural Studies Providing Insights into Production and Conformational Behavior of Amyloid- β Peptide Associated with Alzheimer's Disease Development. *Molecules* 26 (10), , [10.3390/molecules26102897](https://doi.org/10.3390/molecules26102897)
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31. Albrecht C, Kuznetsov AS, Appert-Collin A, Dhaideh Z, Callewaert M, Bershatsky YV, Urban AS, **Bocharov EV**, Bagnard D, Baud S, Blaise S, Romier-Crouzet B, Efremov RG, Dauchez M, Duca L, Gueroult M, Maurice P, Bennasroune A (2020). Transmembrane Peptides as a New Strategy to Inhibit Neuraminidase-1 Activation. *Front Cell Dev Biol* 8, 611121, [10.3389/fcell.2020.611121](https://doi.org/10.3389/fcell.2020.611121)
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