

Резюме: Лебедев Юрий Борисович

Адрес

Федеральное государственное бюджетное учреждение науки Институт биоорганической химии им. академиков М.М. Шемякина и Ю.А. Овчинникова Российской академии наук, Москва, Россия

Контакты

<https://www.ibch.ru/ru/users/25>

Образование

1973–1978

Москва

МГУ

Работа в ИБХ

2018–наст.вр.

Главный научный сотрудник

2026–2024

Профессор

Членство в советах и комиссиях ИБХ

Диссертационный совет

Ученый совет

Научные интересы

Руководитель лаборатории доктор биологических наук Юрий Борисович Лебедев является одним из ведущих отечественных специалистов в области изучения структуры, функционирования и эволюции генома млекопитающих. Исследования в этой области начаты Ю. Б. Лебедевым в начале 1990-х, когда он с группой молодых ученых лаборатории академика Е. Д. Свердлова вошли в число первых участников Российской программы «Геном Человека». Как руководитель лаборатории Ю. Б. Лебедев продолжает разрабатывать оригинальное направление эволюционной геномики, связанное с исследованием функциональных последствий распространения и активации ретропозонов в геномах высших приматов.

Членство в сообществах

В 1996 г. Лебедев Ю. Б. был принят в члены HUGO.

Степени и звания

Профессор

Доктор наук (Биологические науки, 03.00.03 — Молекулярная биология)

Гранты и проекты

2020–2022 [Взаимодействие генотипа и репертуара Т-лимфоцитов при формировании противовирусного иммунного ответа](#)

2015–2019 [Изучение динамики системы клеточного адаптивного иммунитета при флавивиральной инфекции](#)

2017–2020 [Анализ клонального спектра малых субпопуляций Т-лимфоцитов при формировании иммунного ответа у человека](#)

Публикации

1. Salnikova MA, Tuchynskaya KK, Minervina AA, Pogorelyy MV, Okhezin EV, Karganova GG, Mamedov IZ, **Lebedev YB** (2025). Tick-borne encephalitis virus variants drive distinct TCR repertoire alterations. *Front Immunol* 16, 1663781, [10.3389/fimmu.2025.1663781](https://doi.org/10.3389/fimmu.2025.1663781)
2. Salnikova MA, **Lebedev YB** (2024). Longitudinal tracking of T-cell repertoire reveals long-lasting CD4⁺ yellow fever specific clone cluster. *Russian Journal of Infection and Immunity* 14 (3), 539–543, [10.15789/2220-7619-LTO-16665](https://doi.org/10.15789/2220-7619-LTO-16665)
3. Smirnova AO, Miroshnichenkova AM, Belyaeva LD, Kelmanson IV, **Lebedev YB**, Mamedov IZ, Chudakov DM, Komkov AY (2023). Novel bimodal TRBD1-TRBD2 rearrangements with dual or absent D-region contribute to TRB V-(D)-J combinatorial diversity. *Front Immunol* 14, 1245175, [10.3389/fimmu.2023.1245175](https://doi.org/10.3389/fimmu.2023.1245175)
4. Smirnova AO, Miroshnichenkova AM, Olshanskaya YV, Maschan MA, **Lebedev YB**, Chudakov DM, Mamedov IZ, Komkov A (2023). The use of non-functional clonotypes as a natural calibrator for quantitative bias correction in adaptive immune receptor repertoire profiling. *Elife* 12, , [10.7554/eLife.69157](https://doi.org/10.7554/eLife.69157)
5. Komech EA, Koltakova AD, Barinova AA, Minervina AA, Salnikova MA, Shmidt EI, Korotaeva TV, Loginova EY, Erdes SF, Bogdanova EA, Shugay M, Lukyanov S, **Lebedev YB**, Zvyagin IV (2022). TCR repertoire profiling revealed antigen-driven CD8⁺ T cell clonal groups shared in synovial fluid of patients with spondyloarthritis. *Front Immunol* 13, 973243, [10.3389/fimmu.2022.973243](https://doi.org/10.3389/fimmu.2022.973243)
6. Sycheva AL, Komech EA, Pogorelyy MV, Minervina AA, Urazbakhtin SZ, Salnikova MA, Vorovitch MF, Kopantzev EP, Zvyagin IV, Komkov AY, Mamedov IZ, **Lebedev YB** (2022). Inactivated tick-borne encephalitis vaccine elicits several overlapping waves of T cell response. *Front Immunol* 13, 970285, [10.3389/fimmu.2022.970285](https://doi.org/10.3389/fimmu.2022.970285)
7. Urazbakhtin S, Smirnova A, Volakhava A, Zerkalnikova E, Salyutina M, Doubek M, Jelinkova H, Khudainazarova N, Volchkov E, Belyaeva L, Komech E, Pavlova S, **Lebedev Y**, Plevova K, Olshanskaya Y, Komkov A, Mamedov I (2022). The Absence of Retroelement Activity Is Characteristic for Childhood Acute Leukemias and Adult Acute Lymphoblastic Leukemia. *Int J Mol Sci* 23 (3), , [10.3390/ijms23031756](https://doi.org/10.3390/ijms23031756)
8. **(конференция)** Sycheva AL, Pogorelyy MV, Komech EA, Urazbakhtin SZ, Minervina AA, Kopancev EP, Vorovitch MF, Zvyagin IV, Mamedov IZ, **Lebedev YB** (2021). Features of T-cell immune response to tick-borne encephalitis vaccine. *Eur J Immunol* 51 (S1), 1–448, <https://doi.org/10.1002/eji.202170200>
9. **(конференция)** Zvyagin IV, Blagov S, Fomchenkova V, Fadeeva M, Komech EA, Zhogov V, Barinova AA, Mikelov AI, Sycheva AL, **Lebedev YB**, Maschan MA (2021). T cell repertoire sequencing to study the contribution of different donor T cell subsets to patient repertoire at the early stage after $\alpha\beta$ T/CD19-depleted allogeneic hematopoietic stem cell transplantation. *Eur J Immunol* 51 (S1), 1–448, <https://doi.org/10.1002/eji.202170200>
10. **(конференция)** Комеч ЕА, Звягин ИВ, **Лебедев ЮБ**, Сальникова МА, Минервина АА (2021). T-CELL REPERTOIRE OF SYNOVIAL FLUID IN SPONDYLOARTHROPATHIES EXHIBITS HALLMARKS OF HLA-DEPENDENT CLONAL EXPANSIONS AND REMAINS STABLE OVER 1.5 YEARS. *Ann Rheum Dis* (80), 204, [10.1136/annrheumdis-2021-eular.3498](https://doi.org/10.1136/annrheumdis-2021-eular.3498)
11. Kovalenko EI, Zvyagin IV, Streltsova MA, Mikelov AI, Erokhina SA, Telford G, Sapozhnikov AM, **Lebedev YB** (2021). Surface NKG2C identifies differentiated $\alpha\beta$ T-cell clones expanded in peripheral blood. *Front Immunol* 11, 613882, [10.3389/fimmu.2020.613882](https://doi.org/10.3389/fimmu.2020.613882)
12. Minervina AA, Komech EA, Titov A, Koraichi MB, Rosati E, Mamedov IZ, Franke A, Efimov GA, Chudakov DM, Mora T, Walczak AM, **Lebedev YB**, Pogorelyy MV (2021). Longitudinal high-throughput TCR repertoire profiling reveals the dynamics of T-cell memory formation after mild COVID-19 infection. *Elife* 10, 1–17, [10.7554/eLife.63502](https://doi.org/10.7554/eLife.63502)
13. Komkov AY, Urazbakhtin SZ, Saliutina MV, Komech EA, Shelygin YA, Nugmanov GA, Shubin VP, Smirnova AO, Bobrov MY, Tsukanov AS, Snezhkina AV, Kudryavtseva AV, **Lebedev YB**, Mamedov IZ (2020). SeqURE – a new copy-capture based method for sequencing of unknown Retroinsertion events. *Mob DNA* 11 (1), 33, [10.1186/s13100-020-00228-6](https://doi.org/10.1186/s13100-020-00228-6)
14. **(конференция)** Mikelov AI, Komech EA, **Lebedev YB**, Zvyagin IV (2020). In- and off- season peripheral blood T cell repertoire profiling of patients with birch pollen allergy. *Allergy* 75 (S109), 188, [10.1111/all.14506](https://doi.org/10.1111/all.14506)
15. **(конференция)** Комков АЮ, Мамедов ИЗ, **Лебедев ЮБ**, Атапина Е (2020). A cost-effective quasi single-cell assay for deciphering of clonal architecture of leukemic cells. *Klin Padiatr* 232 (3), e7, [10.1055/s-0040-](https://doi.org/10.1055/s-0040-)

[1709799](#)

16. Minervina AA, Pogorelyy MV, Komech EA, Karnaukhov VK, Bacher P, Rosati E, Franke A, Chudakov D, Mamedov IZ, **Lebedev YB**, Mora T, Walczak AM (2020). Primary and secondary anti-viral response captured by the dynamics and phenotype of individual T cell clones. *Elife* 9, , [10.7554/eLife.53704](#)
17. Rosati E, Pogorelyy MV, Dowds CM, Moller FT, Sorensen SB, **Lebedev YB**, Frey N, Schreiber S, Spehlmann ME, Andersen V, Mamedov IZ, Franke A (2019). Identification of disease-associated traits and clonotypes in the T-cell receptor repertoire of monozygotic twins affected by inflammatory bowel diseases. *J Crohns Colitis* 14 (6), 778–790, [10.1093/ecco-jcc/jjz179](#)
18. Микелов АИ, Староверов ДБ, Комеч ЕА, **Лебедев ЮБ**, Чудаков ДМ, Zvyagin IV (2019). Correlated dynamics of serum IGE and IGE+ clonotype count with allergen air level in seasonal allergic rhinitis. *Bulletin of Russian State Medical University* 5 (5), 13–22, [10.24075/brsmu.2019.072](#)
19. Komkov A, Miroshnichenkova A, Nugmanov G, Popov A, Pogorelyy M, Zapletalova E, Jelinkova H, Pospisilova S, **Lebedev Y**, Chudakov D, Olshanskaya Y, Plevova K, Maschan M, Mamedov I (2019). High-throughput sequencing of T-cell receptor alpha chain clonal rearrangements at the DNA level in lymphoid malignancies. *Br J Haematol* 188 (5), 723–731, [10.1111/bjh.16230](#)
20. **(конференция)** Mikelov AI, Turchaninova MA, Komech EA, Staroverov DB, Shvets SM, **Lebedev YB**, Chudakov DM, Zvyagin IV (2019). Longitudinal profiling of immunoglobulin heavy-chain repertoires in memory B-cells, plasmablasts and plasma cells from peripheral blood of individuals with birch pollen allergy. *Allergy* 74 (S106), 174.
21. Pogorelyy MV, Minervina AA, Shugay M, Chudakov DM, **Lebedev YB**, Mora T, Walczak AM (2019). Detecting T cell receptors involved in immune responses from single repertoire snapshots. *PLoS Biol* 17 (6), e3000314, [10.1371/journal.pbio.3000314](#)
22. Nugmanov GA, Komkov AY, Saliutina MV, Minervina AA, **Lebedev YB**, Mamedov IZ (2019). [A Pipeline for the Error-free Identification of Somatic Alu Insertions in High-throughput Sequencing Data]. *Mol Biol (Mosk)* 53 (1), 154–165, [10.1134/S0026898419010117](#)
23. Nugmanov GA, Komkov AY, Saliutina MV, Minervina AA, **Lebedev YB**, Mamedov IZ (2019). A Pipeline for the Error-Free Identification of Somatic Alu Insertions in High-Throughput Sequencing Data. *Mol Biol* 53 (1), 138–146, [10.1134/S0026893319010114](#)
24. Pogorelyy MV, Minervina AA, Touzel MP, Sycheva AL, Komech EA, Kovalenko EI, Karganova GG, Egorov ES, Komkov AY, Chudakov DM, Mamedov IZ, Mora T, Walczak AM, **Lebedev YB** (2018). Precise tracking of vaccine-responding T cell clones reveals convergent and personalized response in identical twins. *Proc Natl Acad Sci U S A* 115 (50), 12704–12709, [10.1073/pnas.1809642115](#)
25. Komkov AY, Minervina AA, Nugmanov GA, Saliutina MV, **Lebedev YB**, Mamedov IZ, Khodosevich KV (2018). An advanced enrichment method for rare somatic retroelement insertions sequencing. *Mob DNA* 9 (1), 31, [10.1186/s13100-018-0136-1](#)
26. **(конференция)** Комеч ЕА, Колтакова АД, Мясоутова АА, Коротаева ТВ, Шмидт НИ, Шостак НА, **Лебедев ЮБ**, Звягин ИВ (2018). Клональная характеристика Т-лимфоцитов очага воспаления у больных со спондилоартропатиями. *Nauchno-Prakticheskaya Revmatologiya* 56 (3), 91.
27. **(конференция)** Звягин ИВ, **Лебедев ЮБ**, Мясоутова АА, Комеч ЕА (2018). TCRbeta CDR3 motif is detected in synovial fluid of patients with different spondyloarthropathies. *FEBS Open Bio* 8, 489.
28. **(конференция)** Fomchenkova E, Komech A, Blagov , Sycheva L, **Lebedev B**, Chudakov M, Maschan A, Zvyagin V (2018). T cell repertoire profiling after hematopoietic stem cell transplantation with CD19/αβT cell depletion and donor lymphocyte infusion. *FEBS Open Bio* 8 (S1), 281: P.09–230–Tue.
29. Komech EA, Pogorelyy MV, Egorov ES, Britanova OV, Rebrikov DV, Bochkova AG, Shmidt EI, Shostak NA, Shugay M, Lukyanov S, Mamedov IZ, **Lebedev YB**, Chudakov DM, Zvyagin IV (2018). CD8+T cells with characteristic T cell receptor beta motif are detected in blood and expanded in synovial fluid of ankylosing spondylitis patients. *Rheumatology (Oxford)* 57 (6), 1097–1104, [10.1093/rheumatology/kex517](#)
30. Komech EA, Zvyagin IV, Pogorelyy MV, Mamedov IZ, Fedorenko DA, **Lebedev YB** (2018). Characterization of the T-cell repertoire after autologous HSCT in patients with ankylosing spondylitis. *Acta Naturae* 10 (2), 48–57, [10.32607/2075851-2018-10-2-48-57](#)
31. Sycheva AL, Pogorelyy MV, Komech EA, Minervina AA, Zvyagin IV, Staroverov DB, Chudakov DM, **Lebedev YB**, Mamedov IZ (2018). Quantitative profiling reveals minor changes of T cell receptor repertoire in response

- to subunit inactivated influenza vaccine. *Vaccine* 36 (12), 1599–1605, [10.1016/j.vaccine.2018.02.027](https://doi.org/10.1016/j.vaccine.2018.02.027)
32. Pogorelyy MV, Minervina AA, Chudakov DM, Mamedov IZ, **Lebedev YB**, Mora T, Walczak AM (2018). Method for identification of condition-associated public antigen receptor sequences. *Elife* 7, , [10.7554/eLife.33050](https://doi.org/10.7554/eLife.33050)
 33. Komech EA, **Lebedev YB**, Koshenkova AV, Syrko DS, Musatkina EA, Lukyanov SA, Chudakov DM, Zvyagin IV (2018). A study of the repertoire of activated T-cell clones obtained from a patient with ankylosing spondylitis. *Bulletin of Russian State Medical University* 7 (1), 65–73, [10.24075/brsmu.2018.001](https://doi.org/10.24075/brsmu.2018.001)
 34. Komech EA, Zvyagin IV, Pogorelyy MV, Mamedov IZ, Fedorenko DA, **Lebedev YB** (2018). Characterization of the T-cell Repertoire after Autologous HSCT in Patients with Ankylosing Spondylitis. *Acta Naturae* 10 (2), 48–57, [10.32607/20758251-2018-10-2-48-57](https://doi.org/10.32607/20758251-2018-10-2-48-57)
 35. Pogorelyy MV, Elhanati Y, Marcou Q, Sycheva AL, Komech EA, Nazarov VI, Britanova OV, Chudakov DM, Mamedov IZ, **Lebedev YB**, Mora T, Walczak AM (2017). Persisting fetal clonotypes influence the structure and overlap of adult human T cell receptor repertoires. *PLoS Comput Biol* 13 (7), e1005572, [10.1371/journal.pcbi.1005572](https://doi.org/10.1371/journal.pcbi.1005572)
 36. Komkov A, Miroshnichenkova A, Minervina A, Nugmanov G, **Lebedev Y**, Mamedov I, Olshanskaya Y, Maschan M (2017). High-throughput sequencing for diagnostics of minimal residual disease in acute lymphoblastic leukemia. *Klin Padiatr* , , [10.1055/s-0037-1602224](https://doi.org/10.1055/s-0037-1602224)
 37. **(конференция)** Pogorelyy M, PuelmaTouzer M, Minervina AA, Sycheva AL, Chudakov DM, Mamedov IZ, Mora T, Walczak AM, **Lebedev YB** (2017). High throughput sequencing of identical twins TCR repertoires after yellow fever vaccination. , 60.
 38. Zvyagin IV, Mamedov IZ, Tatarinova OV, Komech EA, Kurnikova EE, Boyakova EV, Brilliantova V, Shelikhova LN, Balashov DN, Shugay M, Sycheva AL, Kasatskaya SA, **Lebedev YB**, Maschan AA, Maschan MA, Chudakov DM (2017). Tracking T-cell immune reconstitution after TCR $\alpha\beta$ /CD19-depleted hematopoietic cells transplantation in children. *Leukemia* 31 (5), 1145–1153, [10.1038/leu.2016.321](https://doi.org/10.1038/leu.2016.321)
 39. **(конференция)** Сычева АЛ, Погорельый МВ, Комеч ЕА, Звягин ИВ, Мамедов ИЗ, **Лебедев ЮБ** (2017). Изучение малых субпопуляций активированных Т-лимфоцитов из крови доноров, вакцинированных против вируса жёлтой лихорадки. , 98.
 40. **(конференция)** Zvyagin I, Tatarinova O, Mamedov I, Komech E, Maschan A, Shelikhova L, Kurnikova E, Boyakova E, **Lebedev Y**, Maschan M, Chudakov D (2016). T Cell Repertoire after Alpha/Beta-T Cell Depleted Allogeneic Hematopoietic Stem Cell Transplantation in Pediatric Patients. *Blood* (128), 4582.
 41. Nazarov VI, Minervina AA, Komkov AY, Pogorelyy MV, Maschan MA, Olshanskaya YV, Zvyagin IV, Chudakov DM, **Lebedev YB**, Mamedov IZ (2016). Reliability of immune receptor rearrangements as genetic markers for minimal residual disease monitoring. *Bone Marrow Transplant* 51 (10), 1408–1410, [10.1038/bmt.2016.148](https://doi.org/10.1038/bmt.2016.148)
 42. **(конференция)** Komkov AY, Minervina AA, Pogorelyy MV, Zvyagin IV, Panferova A, Olshanskaya Y, Chudakov DM, Maschan M, Mamedov IZ, **Lebedev YB** (2016). Next generation sequencing based approach for monitoring of minimal residual disease in acute lymphoblastic leukemia. *FEBS J* 283 (S1), 376, [10.1111/febs.13808](https://doi.org/10.1111/febs.13808)
 43. **(конференция)** Погорельый МВ, Сычева АЛ, Мамедов ИЗ, Мора Т, Вальзак АМ, **Лебедев ЮБ** (2016). Клоны Т-клеток пуповинной крови обнаруживаются в репертуарах Т-клеточных рецепторов взрослых доноров. , 23.
 44. Minervina AA, Komkov AY, Mamedov IZ, **Lebedev YB** (2016). Advanced lymphoblastic clones detection in T-cell leukemia. *Dokl Biochem Biophys* 467 (1), 85–88, [10.1134/S1607672916020022](https://doi.org/10.1134/S1607672916020022)
 45. **(конференция)** Сычева АЛ, Погорельый МВ, Комеч ЕА, Мамедов ИЗ, **Лебедев ЮБ** (2016). Динамика Т-клеточного репертуара человека в ходе противогриппозной вакцинации и ревакцинации. , 27.
 46. Komkov AY, Miroshnichenkova AM, Olshanskaya YV, Myakova NV, Diakonova YY, Minervina AL, Mamedov IZ, **Lebedev YB**, Maschan AA, Maschan MA (2016). Detection of immunoglobulin genes rearrangements in patients with acute lymphoblastic leukemia using highthroughput next generation sequencing. *Probl Gematol Pereliv Krovi* 61 (4), 200–204, [10.18821/0234-5730-2016-61-4-200-204](https://doi.org/10.18821/0234-5730-2016-61-4-200-204)
 47. Nazarov VI, Pogorelyy MV, Komech EA, Zvyagin IV, Bolotin DA, Shugay M, Chudakov DM, **Lebedev YB**, Mamedov IZ (2015). tcR: An R package for T cell receptor repertoire advanced data analysis. *BMC Bioinformatics* 16 (1), 175, [10.1186/s12859-015-0613-1](https://doi.org/10.1186/s12859-015-0613-1)

48. (конференция) Pogorelyy MV, Sycheva AL, Komech EA, Marcou Q, Elhanati Y, Mora T, Walczak A, Mamedov IZ, **Lebedev YB** (2015). Deep TCR repertoire profiling after seasonal influenza vaccination. , 432.
49. Kurnosov AA, Ustyugova SV, Nazarov VI, Minervina AA, Komkov AY, Shugay M, Pogorelyy MV, Khodosevich KV, Mamedov IZ, **Lebedev YB** (2015). The evidence for increased L1 activity in the site of human adult brain neurogenesis. *PLoS One* 10 (2), e0117854, [10.1371/journal.pone.0117854](https://doi.org/10.1371/journal.pone.0117854)
50. Egorov ES, Merzlyak EM, Shelentov AA, Britanova OV, Sharonov GV, Staroverov DB, Bolotin DA, Davydov AN, Barsova E, **Lebedev YB**, Shugay M, Chudakov DM (2015). Quantitative profiling of immune repertoires for minor lymphocyte counts using unique molecular identifiers. *J Immunol* 194 (12), 6155–6163, [10.4049/jimmunol.1500215](https://doi.org/10.4049/jimmunol.1500215)
51. Zvyagin IV, Pogorelyy MV, Ivanova ME, Komech EA, Shugay M, Bolotin DA, Shelentov AA, Kurnosov AA, Staroverov DB, Chudakov DM, **Lebedev YB**, Mamedov IZ (2014). Distinctive properties of identical twins' TCR repertoires revealed by high-throughput sequencing. *Proc Natl Acad Sci U S A* 111 (16), 5980–5985, [10.1073/pnas.1319389111](https://doi.org/10.1073/pnas.1319389111)
52. Britanova OV, Putintseva EV, Shugay M, Merzlyak EM, Turchaninova MA, Staroverov DB, Bolotin DA, Lukyanov S, Bogdanova EA, Mamedov IZ, **Lebedev YB**, Chudakov DM (2014). Age-Related decrease in TCR repertoire diversity measured with deep and normalized sequence profiling. *J Immunol* 192 (6), 2689–2698, [10.4049/jimmunol.1302064](https://doi.org/10.4049/jimmunol.1302064)
53. Mamedov IZ, Britanova OV, Zvyagin IV, Turchaninova MA, Bolotin DA, Putintseva EV, **Lebedev YB**, Chudakov DM (2013). Preparing unbiased T-cell receptor and antibody cDNA libraries for the deep next generation sequencing profiling. *Front Immunol* 4 (DEC), 456, [10.3389/fimmu.2013.00456](https://doi.org/10.3389/fimmu.2013.00456)
54. Putintseva EV, Britanova OV, Staroverov DB, Merzlyak EM, Turchaninova MA, Shugay M, Bolotin DA, Pogorelyy MV, Mamedov IZ, Bobrynina V, Maschan M, **Lebedev YB**, Chudakov DM (2013). Mother and child T cell receptor repertoires: Deep profiling study. *Front Immunol* 4 (DEC), 463, [10.3389/fimmu.2013.00463](https://doi.org/10.3389/fimmu.2013.00463)
55. Kurnosov AA, Ustyugova SV, Pogorelyy MV, Komkov AY, Bolotin DA, Khodosevich KV, Mamedov IZ, **Lebedev YB** (2013). A novel approach to identification of somatic retroelements' insertions in human genome. *Russ. J. Bioorganic Chem.* 39 (4), 417–425, [10.1134/S1068162013040110](https://doi.org/10.1134/S1068162013040110)
56. Bolotin DA, Mamedov IZ, Britanova OV, Zvyagin IV, Shagin D, Ustyugova SV, Turchaninova MA, Lukyanov S, **Lebedev YB**, Chudakov DM (2012). Next generation sequencing for TCR repertoire profiling: Platform-specific features and correction algorithms. *Eur J Immunol* 42 (11), 3073–3083, [10.1002/eji.201242517](https://doi.org/10.1002/eji.201242517)
57. Britanova OV, Bochkova AG, Staroverov DB, Fedorenko DA, Bolotin DA, Mamedov IZ, Turchaninova MA, Putintseva EV, Kotlobay AA, Lukyanov S, Novik AA, **Lebedev YB**, Chudakov DM (2012). First autologous hematopoietic SCT for ankylosing spondylitis: A case report and clues to understanding the therapy. *Bone Marrow Transplant* 47 (11), 1479–1481, [10.1038/bmt.2012.44](https://doi.org/10.1038/bmt.2012.44)
58. Komkov AY, Maschan MA, Shvets VI, **Lebedev YB** (2012). Functional analysis of polymorphic insertions of alu retroelements in acute lymphoblastic leukemia patients. *Russ. J. Bioorganic Chem.* 38 (3), 306–318, [10.1134/S1068162012030089](https://doi.org/10.1134/S1068162012030089)
59. Mamedov IZ, Ustyugova SV, Amosova FL, **Lebedev YB** (2011). Retroelement insertion polymorphism and modulation of human gene activity. , 203–236.
60. Mamedov IZ, Britanova OV, Bolotin DA, Chkalina AV, Staroverov DB, Zvyagin IV, Kotlobay AA, Turchaninova MA, Fedorenko DA, Novik AA, Sharonov GV, Lukyanov S, Chudakov DM, **Lebedev YB** (2011). Quantitative tracking of T cell clones after haematopoietic stem cell transplantation. *EMBO Mol Med* 3 (4), 201–207, [10.1002/emmm.201100129](https://doi.org/10.1002/emmm.201100129)
61. Zvyagin IV, Mamedov IZ, Britanova OV, Staroverov DB, Nasonov EL, Bochkova AG, Chkalina AV, Kotlobay AA, Korostin DO, Rebrikov DV, Lukyanov S, **Lebedev YB**, Chudakov DM (2010). Contribution of functional KIR3DL1 to ankylosing spondylitis. *Cell Mol Immunol* 7 (6), 471–476, [10.1038/cmi.2010.42](https://doi.org/10.1038/cmi.2010.42)
62. Zvyagin IV, Dorodnykh VY, Mamedov IZ, Staroverov DB, Bochkova AG, Rebrikov DV, **Lebedev YB** (2010). Association of ERAP1 Allelic Variants with Risk of Ankylosing Spondylitis. *Acta Naturae* 2 (3), 72–7.
63. Mamedov IZ, Shagina IA, Kurnikova MA, Novozhilov SN, Shagin DA, **Lebedev YB** (2010). A new set of markers for human identification based on 32 polymorphic Alu insertions. *Eur J Hum Genet* 18 (7), 808–814, [10.1038/ejhg.2010.22](https://doi.org/10.1038/ejhg.2010.22)
64. Chkalina AV, Zvyagin IV, Mamedov IZ, Britanova OV, Staroverov DB, **Lebedev YB** (2010). The oligoclonal expansion of T cells: The investigation of its stability over time. *Russ. J. Bioorganic Chem.* 36 (2), 191–198,

[10.1134/S1068162010020081](https://doi.org/10.1134/S1068162010020081)

65. Shagina I, Bogdanova E, Mamedov IZ, **Lebedev Y**, Lukyanov S, Shagin D (2010). Normalization of genomic DNA using duplex-specific nuclease. *Biotechniques* 48 (6), 455–459, [10.2144/000113422](https://doi.org/10.2144/000113422)
66. Amosova AL, Komkov AY, Ustyugova SV, Mamedov IZ, **Lebedev YB** (2009). Retroposons in modern human genome evolution. *Russ. J. Bioorganic Chem.* 35 (6), 702–710, [10.1134/S1068162009060053](https://doi.org/10.1134/S1068162009060053)
67. Mamedov IZ, Britanova OV, Chkalina AV, Staroverov DB, Amosova AL, Mishin AS, Kurnikova MA, Zvyagin IV, Mutovina ZY, Gordeev AV, Khaidukov SV, Sharonov GV, Shagin DA, Chudakov DM, **Lebedev YB** (2009). Individual characterization of stably expanded T cell clones in ankylosing spondylitis patients. *Autoimmunity* 42 (6), 525–536, [10.1080/08916930902960362](https://doi.org/10.1080/08916930902960362)
68. Panchin AY, Spirin SA, Lukyanov SA, **Lebedev YB**, Panchin YV (2008). Human trash ESTs - Sequences from cDNA collection that are not aligned to genome assembly. *J Bioinform Comput Biol* 6 (4), 759–773, [10.1142/S0219720008003709](https://doi.org/10.1142/S0219720008003709)
69. Mamedov IZ, Amosova AL, Fisunov GY, **Lebedev YB** (2008). A new polymorphic retroelement database (PRED) for the human genome. *Mol Biol* 42 (4), 641–646, [10.1134/S0026893308040213](https://doi.org/10.1134/S0026893308040213)
70. **Lebedev YB**, Amosova AL, Mamedov IZ, Fisunov GY, Sverdlov ED (2007). Most recent AluY insertions in human gene introns reduce the content of the primary transcripts in a cell type specific manner. *Gene* 390 (12), 122–129, [10.1016/j.gene.2006.09.031](https://doi.org/10.1016/j.gene.2006.09.031)
71. Ustyugova SV, **Lebedev YB**, Sverdlov ED (2006). Long L1 insertions in human gene introns specifically reduce the content of corresponding primary transcripts. *Genetika* 128 (13), 261–272, [10.1007/s10709-005-5967-2](https://doi.org/10.1007/s10709-005-5967-2)
72. Kutuev I, Khusainova R, Karunas A, Yunusbayev B, Fedorova S, **Lebedev Y**, Hunsmann G, Khusnutdinova E (2006). From east to west: Patterns of genetic diversity of populations living in four Eurasian regions. *Hum Hered* 61 (1), 1–9, [10.1159/000091309](https://doi.org/10.1159/000091309)
73. Ustyugova SV, Amosova AL, **Lebedev YB**, Sverdlov ED (2006). A tissue-specific decrease in the pre-mRNA level of L1- and Alu-containing alleles of human genes. *Russ. J. Bioorganic Chem.* 32 (1), 93–95, [10.1134/S1068162006010110](https://doi.org/10.1134/S1068162006010110)
74. Buzdin A, Vinogradova T, **Lebedev Y**, Sverdlov E (2005). Genome-wide experimental identification and functional analysis of human specific retroelements. *Cytogenet Genome Res* 110 (14), 468–474, [10.1159/000084980](https://doi.org/10.1159/000084980)
75. Mamedov IZ, Arzumanyan ES, Amosova AL, **Lebedev YB**, Sverdlov ED (2005). Whole-genome experimental identification of insertion/deletion polymorphisms of interspersed repeats by a new general approach. *Nucleic Acids Res* 33 (2), e16, [10.1093/nar/gni018](https://doi.org/10.1093/nar/gni018)
76. Ustyugova SV, Amosova AL, **Lebedev YB**, Sverdlov ED (2005). Cell line fingerprinting using retroelement insertion polymorphism. *Biotechniques* 38 (4), 561–565, [10.2144/05384ST02](https://doi.org/10.2144/05384ST02)
77. Khusainova RI, Akhmetova VL, Kutuyev IA, Salimova AZ, Korshunova TY, **Lebedev YB**, Khusnutdinova EK (2004). Genetic structure of the volga-ural and Central Asian populations inferred from the data on Alu polymorphism. *Genetika* 40 (4), 552–559.
78. Khodosevich KV, **Lebedev YB**, Sverdlov ED (2004). The tissue-specific methylation of human-specific endogenous retroviral LTRs. *Russ. J. Bioorganic Chem.* 30 (5), 441–445, [10.1023/B:RUJBI.0000043787.07628.2a](https://doi.org/10.1023/B:RUJBI.0000043787.07628.2a)
79. Mamedov I, **Lebedev Y**, Hunsmann G, Khusnutdinova E, Sverdlov E (2004). A rare event of insertion polymorphism of a HERV-K LTR in the human genome. *Genomics* 84 (3), 596–599, [10.1016/j.ygeno.2004.04.010](https://doi.org/10.1016/j.ygeno.2004.04.010)
80. Mamedov IZ, **Lebedev YB**, Sverdlov ED (2004). Unusually long target site duplications flanking some of the long terminal repeats of human endogenous retrovirus K in the human genome. *J Gen Virol* 85 (6), 1485–1488, [10.1099/vir.0.19717-0](https://doi.org/10.1099/vir.0.19717-0)
81. Khusainova RI, Akhmetova VL, Kutuyev IA, Salimova AZ, Korshunova TY, **Lebedev YB**, Khusnutdinova EK (2004). Genetic structure of the Volga-Ural and Central Asian populations inferred from the data on Alu polymorphism. *Russ J Genet* 40 (4), 443–449, [10.1023/B:RUJGE.0000024983.75183.8b](https://doi.org/10.1023/B:RUJGE.0000024983.75183.8b)
82. Khodosevich K, **Lebedev Y**, Sverdlov ED (2004). Large-scale determination of the methylation status of retrotransposons in different tissues using a methylation tags approach. *Nucleic Acids Res* 32 (3), e31, [10.1093/nar/gnh035](https://doi.org/10.1093/nar/gnh035)

83. Glazkova DV, Nadezhdin EV, Vinogradova TV, **Lebedev YB**, Bronholdt D, Grzeschik KX, Arman IP, Sverdlov ED (2003). Nucleotide sequences of long terminal repeats of the human endogenous retrovirus (LTR HERV-K) on the short arm of chromosome 7: Identification, analysis, and evaluation of transcriptional activity. *Genetika* 39 (5), 702–708.
84. Glazkova DV, Nadezhdin EV, Vinogradova TV, **Lebedev YB**, Bornholdt D, Grzeschik KX, Arman IP, Sverdlov ED (2003). Nucleotide sequences of long terminal repeats of the human endogenous retroviruses (LTR HERV-K) on the short arm of chromosome 7: Identification, map locations, and transcriptional activity. *Russ J Genet* 39 (5), 578–583, [10.1023/A:1023744019585](https://doi.org/10.1023/A:1023744019585)
85. Buzdin A, Ustyugova S, Gogvadze E, **Lebedev Y**, Hunsmann G, Sverdlov E (2003). Genome-wide targeted search for human specific and polymorphic L1 integrations. *Hum Genet* 112 (56), 527–533, [10.1007/s00439-002-0904-2](https://doi.org/10.1007/s00439-002-0904-2)
86. Buzdin A, Ustyugova S, Khodosevich K, Mamedov I, **Lebedev Y**, Hunsmann G, Sverdlov E (2003). Human-specific subfamilies of HERV-K (HML-2) long terminal repeats: Three master genes were active simultaneously during branching of hominoid lineages. *Genomics* 81 (2), 149–156, [10.1016/S0888-7543\(02\)00027-7](https://doi.org/10.1016/S0888-7543(02)00027-7)
87. Buzdin AA, **Lebedev YB**, Sverdlov ED (2003). Human-Specific HERV-K Intron LTRs Have Nonaccidental Opposite Orientation Relative to the Direction of Gene Transcription and Might Be Involved in the Antisense Regulation of Gene Expression. *Russ. J. Bioorganic Chem.* 29 (1), 91–93, [10.1023/A:1022294906202](https://doi.org/10.1023/A:1022294906202)
88. Domansky AN, Akopov SB, **Lebedev YB**, Nikolaev LG, Sverdlov ED (2002). Enhancer activity of solitary long terminal repeat of human endogenous retrovirus K. *Russ. J. Bioorganic Chem.* 28 (4), 308–311, [10.1023/A:1019543808495](https://doi.org/10.1023/A:1019543808495)
89. Khodosevich K, **Lebedev Y**, Sverdlov E (2002). Endogenous retroviruses and human evolution. *Comp Funct Genomics* 3 (6), 494–498, [10.1002/cfg.216](https://doi.org/10.1002/cfg.216)
90. **Lebedev YB**, Bolorma B, Kzhyskowska JG, Ostashkin AS, Ilyin KV, Myandina GI, Pyagai PE, Itkes AV (2002). Integration of the type D Mason-Pfizer monkey virus into the human chromosome. *Mol Biol* 36 (6), 821–822, [10.1023/A:1021681909390](https://doi.org/10.1023/A:1021681909390)
91. Mamedov I, Batrak A, Buzdin A, Arzumanyan E, **Lebedev Y**, Sverdlov ED (2002). Genome-wide comparison of differences in the integration sites of interspersed repeats between closely related genomes. *Nucleic Acids Res* 30 (14), e71, [10.1093/nar/gnf071](https://doi.org/10.1093/nar/gnf071)
92. Buzdin A, Ustyugova S, Gogvadze E, Vinogradova T, **Lebedev Y**, Sverdlov E (2002). A new family of chimeric retrotranscripts formed by a full copy of U6 small nuclear RNA fused to the 3' terminus of L1. *Genomics* 80 (4), 402–406, [10.1006/geno.2002.6843](https://doi.org/10.1006/geno.2002.6843)
93. Domansky AN, Akopov SB, **Lebedev YB**, Nikolaev LG, Sverdlov ED (2002). Enhancer activity of solitary long terminal repeat of human endogenous retrovirus K. *Bioorg Khim* 28 (4), 345.
94. Buzdin A, Khodosevich K, Mamedov I, Vinogradova T, **Lebedev Y**, Hunsmann G, Sverdlov E (2002). A technique for genome-wide identification of differences in the interspersed repeats integrations between closely related genomes and its application to detection of human-specific integrations of HERV-K LTRs. *Genomics* 79 (3), 413–422, [10.1006/geno.2002.6705](https://doi.org/10.1006/geno.2002.6705)
95. Nadezhdin EV, **Lebedev YB**, Glazkova DV, Bornholdt D, Arman IP, Grzeschik KH, Hunsmann G, Sverdlov ED (2001). Identification of paralogous HERV-K LTRs on human chromosomes 3, 4, 7 and 11 in regions containing clusters of olfactory receptor genes. *Mol Genet Genomics* 265 (5), 820–825, [10.1007/s004380100476](https://doi.org/10.1007/s004380100476)
96. Kurdyukov SG, **Lebedev YB**, Artamonova II, Gorodentseva TN, Batrak AV, Mamedov IZ, Azhikina TL, Legchilina SP, Efimenko IG, Gardiner K, Sverdlov ED (2001). Full-sized HERV-K (HML-2) human endogenous retroviral LTR sequences on human chromosome 21: Map locations and evolutionary history. *Gene* 273 (1), 51–61, [10.1016/S0378-1119\(01\)00570-4](https://doi.org/10.1016/S0378-1119(01)00570-4)
97. Domansky AN, Kopantzev EP, Snezhkov EV, **Lebedev YB**, Leib-Mosch C, Sverdlov ED (2000). Solitary HERV-K LTRs possess bi-directional promoter activity and contain a negative regulatory element in the U5 region. *FEBS Lett* 472 (23), 191–195, [10.1016/S0014-5793\(00\)01460-5](https://doi.org/10.1016/S0014-5793(00)01460-5)
98. **Lebedev YB**, Belonovitch OS, Zybrowa NV, Khil PP, Kurdyukov SG, Vinogradova TV, Hunsmann G, Sverdlov ED (2000). Differences in HERV-K LTR insertions in orthologous loci of humans and great apes. *Gene* 247 (12), 265–277, [10.1016/S0378-1119\(00\)00062-7](https://doi.org/10.1016/S0378-1119(00)00062-7)

99. **Lebedev YB** (2000). Endogenous retroviruses: A possible role in human cell function. *Mol Biol* 34 (4), 544–553, [10.1007/BF02759563](https://doi.org/10.1007/BF02759563)
100. Bogush ML, Velikodvorskaya TV, **Lebedev YB**, Nikolaev LG, Lukyanov SA, Fradkov AF, Pliyev BK, Boichenko MN, Usatova GN, Vorobiev AA, Andersen GL, Sverdlov ED (1999). Identification and localization of differences between Escherichia coli and Salmonella typhimurium genomes by suppressive subtractive hybridization. *Mol Gen Genet* 262 (45), 721–729, [10.1007/s004380051134](https://doi.org/10.1007/s004380051134)
101. Belyaeva OV, Balanovsky OP, Ashworth LK, **Lebedev YB**, Spitsyn VA, Guseva NA, Erdes S, Mikulich AI, Khusnutdinova EK, Limborska SA (1999). Fine mapping of a polymorphic CA repeat marker on human chromosome 19 and its use in population studies. *Gene* 230 (2), 259–266, [10.1016/S0378-1119\(99\)00056-6](https://doi.org/10.1016/S0378-1119(99)00056-6)
102. Lavrentieva I, Broude NE, **Lebedev Y**, Gottesman II, Lukyanov SA, Smith CL, Sverdlov ED (1999). High polymorphism level of genomic sequences flanking insertion sites of human endogenous retroviral long terminal repeats. *FEBS Lett* 443 (3), 341–347, [10.1016/S0014-5793\(99\)00004-6](https://doi.org/10.1016/S0014-5793(99)00004-6)
103. Lapuk AV, Khil PP, Lavrentieva IV, **Lebedev YB**, Sverdlov ED (1999). A human endogenous retrovirus-like (HERV) LTR formed more than 10 million years ago due to an insertion of HERV-H LTR into the 5' LTR of HERV-K is situated on human chromosomes 10, 19 and Y. *J Gen Virol* 80 (4), 835–839, [10.1099/0022-1317-80-4-835](https://doi.org/10.1099/0022-1317-80-4-835)
104. Khil PP, **Lebedev YB**, Sverdlov ED (1998). A new putative gene preferentially expressed in the human brain located on chromosome 19q12 near the human endogenous virus HERV-K LTR. *Russ. J. Bioorganic Chem.* 24 (1), 62–63.
105. Khil PP, **Lebedev YB**, Sverdlov ED (1998). The human endogenous virus HERV-K long terminal repeat in an intron of the ZNF91 gene. *Russ. J. Bioorganic Chem.* 24 (2), 112–116.
106. Khil PP, **Lebedev YB**, Sverdlov ED (1998). A New Putative Gene Preferentially Expressed in the Human Brain Located on Chromosome 19q12 near the Human Endogenous Virus HERV-K LTR. *Bioorg Khim* 24 (1), 73–74.
107. Klinov DV, Lagutina IV, Prokhorov VV, Neretina T, Khil PP, **Lebedev YB**, Cherny DI, Demin VV, Sverdlov ED (1998). High resolution mapping DNAs by R-loop atomic force microscopy. *Nucleic Acids Res* 26 (20), 4603–4610, [10.1093/nar/26.20.4603](https://doi.org/10.1093/nar/26.20.4603)
108. Lavrentieva I, Khil P, Vinogradova T, Akhmedov A, Lapuk A, Shakhova O, **Lebedev Y**, Monastyrskaya G, Sverdlov ED (1998). Subfamilies and nearest-neighbour dendrogram for the LTRs of human endogenous retroviruses HERV-K mapped on human chromosome 19: Physical neighbourhood does not correlate with identity level. *Hum Genet* 102 (1), 107–116, [10.1007/s004390050662](https://doi.org/10.1007/s004390050662)
109. Akopov SB, Nikolaev LG, Khil PP, **Lebedev YB**, Sverdlov ED (1998). Long terminal repeats of human endogenous retrovirus K family (HERV-K) specifically bind host cell nuclear proteins. *FEBS Lett* 421 (3), 229–233, [10.1016/S0014-5793\(97\)01569-X](https://doi.org/10.1016/S0014-5793(97)01569-X)
110. Khil PP, **Lebedev YB**, Sverdlov ED (1998). The Human Endogenous Virus HERV-K Long Terminal Repeat in an Intron of the ZNF91 Gene. *Bioorg Khim* 24 (2), 130–131.
111. Khil PP, Kostina MB, Azhikina TL, Kolesnik TB, **Lebedev YB**, Sverdlov ED (1997). Structural Features of Four Long Terminal Repeats of Human Endogenous Retroviruses and Their Integration Sites. *Bioorg Khim* 23 (5), 440.
112. Vinogradova T, Volik S, **Lebedev Y**, Shevchenko Y, Lavrentyeva I, Khil P, Grzeschik KH, Ashworth LK, Sverdlov E (1997). Positioning of 72 potentially full size LTRs of human endogenous retroviruses HERV-K on the human chromosome 19 map. Occurrences of the LTRs in human gene sites. *Gene* 199 (12), 255–264, [10.1016/S0378-1119\(97\)00376-4](https://doi.org/10.1016/S0378-1119(97)00376-4)
113. Khil PP, Kostina MB, Azhikina TL, Kolesnik TB, **Lebedev YB**, Sverdlov ED (1997). Structural features of four long terminal repeats of human endogenous retroviruses and their integration sites. *Russ. J. Bioorganic Chem.* 23 (5), 406–411.
114. **Lebedev Y**, Akopyants N, Azhikina T, Shevchenko Y, Potapov V, Stecenko D, Berg D, Sverdlov E (1996). Oligonucleotides containing 2-aminoadenine and 5-methylcytosine are more effective as primers for PCR amplification than their nonmodified counterparts. *Genet Anal Tech Appl* 13 (1), 15–21, [10.1016/1050-3862\(96\)00139-8](https://doi.org/10.1016/1050-3862(96)00139-8)
115. **Lebedev YB**, Volik SV, Obradovic D, Ermolaeva OD, Ashworth LK, Lennon GG, Sverdlov ED (1995). Physical mapping of sequences homologous to an endogenous retrovirus LTR on human chromosome 19.

Mol Gen Genet 247 (6), 742–748, [10.1007/BF00290406](https://doi.org/10.1007/BF00290406)

116. Volik S, **Lebedev Y**, Nikolaev L, Shevchenko Y, Vinogradova T, Kopantzev E, Kolesnik T, Monastyrskaya G, Kunz U, Grzeschik KH, Ashworth LK, Lennon G, Sverdlov E (1995). Mapping of transcribed sequences on human chromosome 19. *Mitochondrial DNA* 6 (1), 13–26, [10.3109/10425179509074694](https://doi.org/10.3109/10425179509074694)
117. Borodin A, Kopantzev E, Wagner L, Volik S, Ermolaeva O, **Lebedev Y**, Monastyrskaya G, Kunz J, Grzeschik KH, Sverdlov E (1995). An arrayed library enriched in hncDNA corresponding to transcribed sequences of human chromosome 19: preparation and analysis. *Genet Anal Tech Appl* 12 (1), 23–31, [10.1016/1050-3862\(95\)00106-9](https://doi.org/10.1016/1050-3862(95)00106-9)