

Resource Summary Report

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Adaptive Biotechnologies

RRID:SCR_014709

Type: Tool

Proper Citation

Adaptive Biotechnologies (RRID:SCR_014709)

Resource Information

URL: <http://www.adaptivebiotech.com>

Proper Citation: Adaptive Biotechnologies (RRID:SCR_014709)

Description: Commercial organization which combines high-throughput sequencing and bioinformatics to profile T-cell and B-cell receptors.

Resource Type: resource

Keywords: high throughput, sequencing, bioinformatics, t cell, b cell, biotechnology

Availability: Commercially available

Resource Name: Adaptive Biotechnologies

Resource ID: SCR_014709

Record Creation Time: 20220129T080321+0000

Record Last Update: 20260808T190016+0000

Ratings and Alerts

No rating or validation information has been found for Adaptive Biotechnologies.

No alerts have been found for Adaptive Biotechnologies.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 24 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Zhai Y, et al. (2026) In vitro recapitulation of intramuscular mRNA vaccination with naive and recall antigens using a human lymphoid follicle chip platform. *iScience*, 29(7), 116416.

Tran B, et al. (2026) Safety, Pharmacokinetics, Pharmacodynamics, and Preliminary Efficacy from a First-in-Human Study of Volrustomig, a Novel PD-1/CTLA-4 Bispecific Antibody. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 32(14), 2850.

Robinson SI, et al. (2025) Oncolytic measles virotherapy encoding the neutrophil-activating protein is effective in synovial sarcoma. *Molecular therapy. Oncology*, 33(4), 201062.

Beck S, et al. (2025) Immunogenomic profiles of HIV-associated classic Hodgkin lymphoma from Malawi. *Blood global hematology*, 1(3).

Brummel K, et al. (2025) Spatiotemporal Immune Landscape and Long-term Immune Memory in POLE-Mutant Endometrial Cancer at the Single-Cell Level. *Cancer immunology research*, 13(12), 1911.

Gonzalez-Kozlova E, et al. (2024) Tumor-Immune Signatures of Treatment Resistance to Brentuximab Vedotin with Ipilimumab and/or Nivolumab in Hodgkin Lymphoma. *Cancer research communications*, 4(7), 1726.

Kato H, et al. (2023) Immune repertoire profiling for disease pathobiology. *Pathology international*, 73(1), 1.

Nawrocki ST, et al. (2023) Comprehensive Single-Cell Immune Profiling Defines the Patient Multiple Myeloma Microenvironment Following Oncolytic Virus Therapy in a Phase Ib Trial. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 29(24), 5087.

Chow A, et al. (2021) Tim-4+ cavity-resident macrophages impair anti-tumor CD8+ T cell immunity. *Cancer cell*, 39(7), 973.

Springer I, et al. (2021) Contribution of T Cell Receptor Alpha and Beta CDR3, MHC Typing, V and J Genes to Peptide Binding Prediction. *Frontiers in immunology*, 12, 664514.

Lan Y, et al. (2021) Simultaneous targeting of TGF- β /PD-L1 synergizes with radiotherapy by reprogramming the tumor microenvironment to overcome immune evasion. *Cancer cell*, 39(10), 1388.

El Meskini R, et al. (2021) Distinct Biomarker Profiles and TCR Sequence Diversity Characterize the Response to PD-L1 Blockade in a Mouse Melanoma Model. *Molecular cancer research : MCR*, 19(8), 1422.

Gurusamy D, et al. (2020) Multi-phenotype CRISPR-Cas9 Screen Identifies p38 Kinase as a Target for Adoptive Immunotherapies. *Cancer cell*, 37(6), 818.

Welsh RA, et al. (2020) Lack of the MHC class II chaperone H2-O causes susceptibility to autoimmune diseases. *PLoS biology*, 18(2), e3000590.

Mandric I, et al. (2020) Profiling immunoglobulin repertoires across multiple human tissues using RNA sequencing. *Nature communications*, 11(1), 3126.

Springer I, et al. (2020) Prediction of Specific TCR-Peptide Binding From Large Dictionaries of TCR-Peptide Pairs. *Frontiers in immunology*, 11, 1803.

Shi B, et al. (2020) The Usage of Human IGHJ Genes Follows a Particular Non-random Selection: The Recombination Signal Sequence May Affect the Usage of Human IGHJ Genes. *Frontiers in genetics*, 11, 524413.

Glass DR, et al. (2020) An Integrated Multi-omic Single-Cell Atlas of Human B Cell Identity. *Immunity*, 53(1), 217.

Corgnac S, et al. (2020) CD103+CD8+ TRM Cells Accumulate in Tumors of Anti-PD-1-Responder Lung Cancer Patients and Are Tumor-Reactive Lymphocytes Enriched with Tc17. *Cell reports. Medicine*, 1(7), 100127.

Wolf K, et al. (2018) Identifying and Tracking Low-Frequency Virus-Specific TCR Clonotypes Using High-Throughput Sequencing. *Cell reports*, 25(9), 2369.