

Resource Summary Report

Generated by [nidm-terms](#) on Aug 10, 2026

Eton Bioscience

RRID:SCR_003533

Type: Tool

Proper Citation

Eton Bioscience (RRID:SCR_003533)

Resource Information

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

Proper Citation: Eton Bioscience (RRID:SCR_003533)

Description: An Antibody supplier

Synonyms: Eton Bioscience Inc.

Resource Type: organization portal, portal, data or information resource, department portal

Resource Name: Eton Bioscience

Resource ID: SCR_003533

Alternate IDs: nlx_152359

Record Creation Time: 20220129T080219+0000

Record Last Update: 20260810T040345+0000

Ratings and Alerts

No rating or validation information has been found for Eton Bioscience.

No alerts have been found for Eton Bioscience.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 87 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](https://nidm-terms.org) .

Segovia-Mota JP, et al. (2026) Enhanced Detection of *Coccidioides* spp. Fungi from Environmental Samples Using Droplet Digital PCR. *Emerging infectious diseases*, 32(4), 500.

Bhattacharjee D, et al. (2026) Genomic insights into the functional and metabolic versatility of gut microbiome *Anaerostipes* species. *Microbial genomics*, 12(2).

Pedroza Romo MJ, et al. (2026) Optimal 1TEL-target protein linker character is target protein-dependent. *Acta crystallographica. Section D, Structural biology*, 82(Pt 5), 516.

Osborne C, et al. (2026) A candidate gene marker at the red kidney color locus (Rk) enables the development of slow-darkening pink beans. *TAG. Theoretical and applied genetics. Theoretische und angewandte Genetik*, 139(6).

Bhattacharjee D, et al. (2025) Diversity and prevalence of *Anaerostipes* in the human gut microbiota. *bioRxiv : the preprint server for biology*.

Lee GY, et al. (2024) Single missense mutations in Vi capsule synthesis genes confer hypervirulence to *Salmonella* Typhi. *Nature communications*, 15(1), 5258.

McClean PE, et al. (2024) T and Z, partial seed coat patterning genes in common bean, provide insight into the structure and protein interactions of a plant MBW complex. *G3 (Bethesda, Md.)*, 14(10).

Lee GY, et al. (2024) Single missense mutations in Vi capsule synthesis genes confer hypervirulence to *Salmonella* Typhi. *bioRxiv : the preprint server for biology*.

Dao UM, et al. (2023) Stripes and loss of color in ball pythons (*Python regius*) are associated with variants affecting endothelin signaling. *G3 (Bethesda, Md.)*, 13(7).

Jordan JT, et al. (2023) Prospective phase II trial of the dual mTORC1/2 inhibitor vistusertib for progressive or symptomatic meningiomas in persons with neurofibromatosis 2. *Neuro-oncology advances*, 5(1), vdad041.

Bahar T, et al. (2023) Unveiling *Lathyrus aphaca* L. as a Newly Identified Host for Begomovirus Infection: A Comprehensive Study. *Genes*, 14(6).

Arad N, et al. (2023) RNA interference-mediated knockdown of genes involved in sugar transport and metabolism disrupts psyllid *Bactericera cockerelli* (Order: Hemiptera) gut physiology and results in high mortality. *Frontiers in insect science*, 3, 1283334.

Gajjar PL, et al. (2023) Increasing the bulk of the 1TEL-target linker and retaining the 10??His tag in a 1TEL-CMG2-vWa construct improves crystal order and diffraction limits. *Acta crystallographica. Section D, Structural biology*, 79(Pt 10), 925.

Chen K, et al. (2023) Detection of *Orientia* spp. Bacteria in Field-Collected Free-Living *Eutrombicula Chigger* Mites, United States. *Emerging infectious diseases*, 29(8), 1676.

Brown AR, et al. (2022) A community-science approach identifies genetic variants associated with three color morphs in ball pythons (*Python regius*). PloS one, 17(10), e0276376.

Bucholz JR, et al. (2022) RAD-tag and mitochondrial DNA sequencing reveal the genetic structure of a widespread and regionally imperiled freshwater mussel, *Obovaria olivaria* (Bivalvia: Unionidae). Ecology and evolution, 12(1), e8560.

McClellan PE, et al. (2022) The Common Bean V Gene Encodes Flavonoid 3'5' Hydroxylase: A Major Mutational Target for Flavonoid Diversity in Angiosperms. Frontiers in plant science, 13, 869582.

Leyn SA, et al. (2021) Experimental evolution in morbidostat reveals converging genomic trajectories on the path to triclosan resistance. Microbial genomics, 7(5).

Kumar A, et al. (2021) Generation and validation of CRISPR-engineered human natural killer cell lines for research and therapeutic applications. STAR protocols, 2(4), 100874.

Zhao X, et al. (2021) Glutamate metabolism directs energetic trade-offs to shape host-pathogen susceptibility in *Drosophila*. Cell metabolism, 33(12), 2428.