Resource Summary Report

Generated by NIF on Aug 27, 2026

Tufts University; Massachusetts; USA

RRID:SCR_000994
Type: Tool

Proper Citation

Tufts University; Massachusetts; USA (RRID:SCR_000994)

Resource Information

URL: http://www.tufts.edu/

Proper Citation: Tufts University; Massachusetts; USA (RRID:SCR_000994)

Description: Private research university in Medford and Somerville, Massachusetts, United States, with additional facilities in Boston and Grafton, as well as Talloires, France. Provides undergraduate, graduate, and professional degree programs.

Synonyms: Tufts University, Tufts

Resource Type: university

Keywords: tufts, university, undergraduate, graduate, professional

Resource Name: Tufts University; Massachusetts; USA

Resource ID: SCR_000994

Alternate IDs: Crossref funder ID:100008090, ISNI:0000 0004 1936 7531, Wikidata:Q49120, nlx_38314, grid.429997.8

Alternate URLs: https://ror.org/05wvpxv85

Record Creation Time: 20220129T080204+0000

Record Last Update: 20260815T182150+0000

Ratings and Alerts

No rating or validation information has been found for Tufts University; Massachusetts; USA.

No alerts have been found for Tufts University; Massachusetts; USA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 9 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF .

Yin Q, et al. (2018) Long-range haplotype analysis of the malaria parasite receptor gene ACKR1 in an East-African population. Human genome variation, 5, 26.

Al-Naemi AH, et al. (2018) Is the rs1801282 (G/C) Polymorphism of PPAR - Gamma Gene Associated with T2DM in Iraqi People? Open access Macedonian journal of medical sciences, 6(3), 447.

Al-Bustan SA, et al. (2017) Increased Risk of the APOB rs11279109 Polymorphism for CHD among the Kuwaiti Population. Disease markers, 2017, 6963437.

Zurawek M, et al. (2017) MAVS is not a Likely Susceptibility Locus for Addison's Disease and Type 1 Diabetes. Archivum immunologiae et therapiae experimentalis, 65(3), 271.

Dawidowska M, et al. (2016) Association of germline genetic variants in RFC, IL15 and VDR genes with minimal residual disease in pediatric B-cell precursor ALL. Scientific reports, 6, 29427.

Ozbek IC, et al. (2016) Timing of coronary artery bypass surgery in patients with non-ST-segment elevation myocardial infarction and postoperative outcomes. Archives of medical science : AMS, 12(4), 766.

El Khoury L, et al. (2016) MMP3 and TIMP2 gene variants as predisposing factors for Achilles tendon pathologies: Attempted replication study in a British case-control cohort. Meta gene, 9, 52.

Sheahan KL, et al. (2015) Identification of mammalian proteins that collaborate with type III secretion system function: involvement of a chemokine receptor in supporting translocon activity. mBio, 6(1), e02023.

Sen A, et al. (2011) Impact of interleukin-6 promoter polymorphism and serum interleukin-6 level on the acute inflammation and neovascularization stages of patients with Eales' disease. Molecular vision, 17, 2552.