

Resource Summary Report

Generated by [dkNET](#) on Aug 8, 2026

TraIT

RRID:SCR_010493

Type: Tool

Proper Citation

TraIT (RRID:SCR_010493)

Resource Information

URL: <http://www.ctmm.nl/en/programmas/infrastructuren/traitprojecttranslationeleresearch>

Proper Citation: TraIT (RRID:SCR_010493)

Description: An ambitious project to develop an IT infrastructure for translational research that aims to facilitate the collection, storage, analysis, archiving, sharing and securing of data.

Synonyms: Translational Research IT, TraIT project

Resource Type: software resource

Resource Name: TraIT

Resource ID: SCR_010493

Alternate IDs: nlx_158039

Record Creation Time: 20220129T080259+0000

Record Last Update: 20260801T190414+0000

Ratings and Alerts

No rating or validation information has been found for TraIT.

No alerts have been found for TraIT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 23 mentions in open access literature.

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

Gee M, et al. (2026) Yield impact of source-sink-regulated senescence in hybrid maize and genetic architecture in exPVP inbreds. TAG. Theoretical and applied genetics. Theoretische und angewandte Genetik, 139(5).

Jacques A, et al. (2026) Optimising the Use of Cryopreserved Genetic Resources for the Selection and Conservation of Animal Populations. Journal of animal breeding and genetics = Zeitschrift fur Tierzuchtung und Zuchtungsbiologie, 143(1), 50.

Mwangangi W, et al. (2026) Yield Performance, Combining Ability and Stability of Early- to Medium-Maturing Doubled-Haploid Maize Lines in Eastern Africa. Plant breeding = Zeitschrift fur Pflanzenzuchtung, 145(3), 671.

Horn SS, et al. (2025) Genetic and metabolic characterization of individual differences in liver fat accumulation in Atlantic salmon. Frontiers in genetics, 16, 1512769.

Baucom RS, et al. (2025) Off-target drift of the herbicide dicamba disrupts plant-pollinator interactions via novel pathways. The New phytologist, 247(2), 850.

Aspromonte MC, et al. (2025) CAGI6 ID panel challenge: assessment of phenotype and variant predictions in 415 children with neurodevelopmental disorders (NDDs). Human genetics, 144(2-3), 227.

Rochelle TL, et al. (2025) The influence of gender and body satisfaction on attentional bias among young adults in Hong Kong: An eye-tracking study. Journal of health psychology, 30(10), 2696.

Proß T, et al. (2025) Within-individual leaf trait response to local light availability and biodiversity in a subtropical forest experiment. Ecology, 106(7), e70160.

Zhao Y, et al. (2025) Identification of Candidate Lung Function-Related Plasma Proteins to Pinpoint Drug Targets for Common Pulmonary Diseases: A Comprehensive Multi-Omics Integration Analysis. Current issues in molecular biology, 47(3).

Ngetich C, et al. (2024) The potential for use of haematological and anti-IgE humoral responses as phenotypic markers for tick resistance in cattle. Current research in parasitology & vector-borne diseases, 5, 100159.

Vexler L, et al. (2024) QTL discovery for agronomic and quality traits in diploid potato clones using PotatoMASH amplicon sequencing. G3 (Bethesda, Md.), 14(10).

Rubino C, et al. (2024) Gamified Practice Improves Paretic Arm Motor Behavior in Individuals With Stroke. Neurorehabilitation and neural repair, 38(11-12), 832.

Desai R, et al. (2023) Examining the Lancet Commission risk factors for dementia using Mendelian randomisation. BMJ mental health, 26(1).

Fontana D, et al. (2023) Evolutionary signatures of human cancers revealed via genomic analysis of over 35,000 patients. Nature communications, 14(1), 5982.

Valdés-Hernández J, et al. (2023) Global analysis of the association between pig muscle fatty acid composition and gene expression using RNA-Seq. *Scientific reports*, 13(1), 535.

Lin A, et al. (2021) Transcriptomic profiling of whole blood in 22q11.2 reciprocal copy number variants reveals that cell proportion highly impacts gene expression. *Brain, behavior, & immunity - health*, 18, 100386.

Kontio JAJ, et al. (2021) Model guided trait-specific co-expression network estimation as a new perspective for identifying molecular interactions and pathways. *PLoS computational biology*, 17(5), e1008960.

Antunes R, et al. (2020) Exploring Lifestyle Habits, Physical Activity, Anxiety and Basic Psychological Needs in a Sample of Portuguese Adults during COVID-19. *International journal of environmental research and public health*, 17(12).

Ramazzotti D, et al. (2019) Learning mutational graphs of individual tumour evolution from single-cell and multi-region sequencing data. *BMC bioinformatics*, 20(1), 210.

Jalalzadeh H, et al. (2019) Design and protocol of a comprehensive multicentre biobank for abdominal aortic aneurysms. *BMJ open*, 9(8), e028858.