

Resource Summary Report

Generated by T1D on Aug 4, 2026

U.S. Army Medical Research and Materiel Command

RRID:SCR_011586

Type: Tool

Proper Citation

U.S. Army Medical Research and Materiel Command (RRID:SCR_011586)

Resource Information

URL: <http://mrmc.amedd.army.mil/>

Proper Citation: U.S. Army Medical Research and Materiel Command (RRID:SCR_011586)

Description: United States Army's medical materiel developer, with responsibility for medical research, development, and acquisition and medical logistics management.

Abbreviations: MRMC, USAMRMC

Synonyms: Medical Research and Materiel Command, Medical Research and Materiel Command - U.S. Army Medical Department, Medical Research and Materiel Command U.S. Army Medical Department, U.S. Army Medical Department Medical Research and Materiel Command, US Army Medical Research and Materiel Command

Resource Type: institution

Keywords: Government granting agency

Resource Name: U.S. Army Medical Research and Materiel Command

Resource ID: SCR_011586

Alternate IDs: nlx_157932, Crossref funder ID: 100000182, Wikidata: Q7889498, ISNI: 0000 0001 0036 4726, grid.420210.5

Alternate URLs: <https://ror.org/03cd02q50>

Record Creation Time: 20220129T080305+0000

Record Last Update: 20260801T190428+0000

Ratings and Alerts

No rating or validation information has been found for U.S. Army Medical Research and Materiel Command.

No alerts have been found for U.S. Army Medical Research and Materiel Command.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 15 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Rifkin RE, et al. (2019) Use of a pressure-sensing walkway system for biometric assessment of gait characteristics in goats. PloS one, 14(10), e0223771.

Weightman MM, et al. (2017) Further Development of the Assessment of Military Multitasking Performance: Iterative Reliability Testing. PloS one, 12(1), e0169104.

Fukuda MM, et al. (2017) A randomized, double-blind, active-control trial to evaluate the efficacy and safety of a three day course of tafenoquine monotherapy for the treatment of Plasmodium vivax malaria. PloS one, 12(11), e0187376.

Adebiyi A, et al. (2017) Assessment of feedback modalities for wearable visual aids in blind mobility. PloS one, 12(2), e0170531.

Grin'kina NM, et al. (2016) Righting Reflex Predicts Long-Term Histological and Behavioral Outcomes in a Closed Head Model of Traumatic Brain Injury. PloS one, 11(9), e0161053.

Banerjee A, et al. (2016) YY1 Is Required for Germinal Center B Cell Development. PloS one, 11(5), e0155311.

Minlikeeva AN, et al. (2016) Single-Nucleotide Polymorphisms and Markers of Oxidative Stress in Healthy Women. PloS one, 11(6), e0156450.

Tsianos GA, et al. (2016) Validated Predictions of Metabolic Energy Consumption for Submaximal Effort Movement. PLoS computational biology, 12(6), e1004911.

AbdulHameed MD, et al. (2016) Mining kidney toxicogenomic data by using gene co-expression modules. BMC genomics, 17(1), 790.

Vital-Lopez FG, et al. (2015) Biofilm Formation Mechanisms of Pseudomonas aeruginosa Predicted via Genome-Scale Kinetic Models of Bacterial Metabolism. PLoS computational biology, 11(10), e1004452.

Moore SE, et al. (2015) Statistical Machines for Trauma Hospital Outcomes Research: Application to the PProspective, Observational, Multi-Center Major Trauma Transfusion

(PROMMTT) Study. PloS one, 10(8), e0136438.

R??bago CA, et al. (2015) Reliability and Minimum Detectable Change of Temporal-Spatial, Kinematic, and Dynamic Stability Measures during Perturbed Gait. PloS one, 10(11), e0142083.

Nagaraja S, et al. (2015) Computational Identification of Mechanistic Factors That Determine the Timing and Intensity of the Inflammatory Response. PLoS computational biology, 11(12), e1004460.

Heier CR, et al. (2014) Non-invasive MRI and spectroscopy of mdx mice reveal temporal changes in dystrophic muscle imaging and in energy deficits. PloS one, 9(11), e112477.

AbdulHameed MD, et al. (2014) Systems level analysis and identification of pathways and networks associated with liver fibrosis. PloS one, 9(11), e112193.