

Resource Summary Report

Generated by ASWG on Aug 4, 2026

U.S. Department of Defense

RRID:SCR_012712

Type: Tool

Proper Citation

U.S. Department of Defense (RRID:SCR_012712)

Resource Information

URL: <http://www.defense.gov/>

Proper Citation: U.S. Department of Defense (RRID:SCR_012712)

Description: Executive branch department of the federal government charged with coordinating and supervising all agencies and functions of the government directly related to national security and the United States Armed Forces.

Abbreviations: DOD

Synonyms: Department of Defense, United States Department of Defense, US Department of Defense

Resource Type: institution

Keywords: Government granting agency

Resource Name: U.S. Department of Defense

Resource ID: SCR_012712

Alternate IDs: ISNI: 0000 0004 0478 6223, grid.420391.d, nlx_143670, Crossref funder ID: 100000005, Wikidata: Q11209

Alternate URLs: <https://ror.org/0447fe631>

Record Creation Time: 20220129T080311+0000

Record Last Update: 20260801T190438+0000

Ratings and Alerts

No rating or validation information has been found for U.S. Department of Defense.

No alerts have been found for U.S. Department of Defense.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 41 mentions in open access literature.

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

Park J, et al. (2025) Astrocytic modulation of population encoding in mouse visual cortex via GABA transporter 3 revealed by multiplexed CRISPR/Cas9 gene editing. eLife, 14.

Avraham-Davidi I, et al. (2025) Spatially defined multicellular functional units in colorectal cancer revealed from single cell and spatial transcriptomics. eLife, 14.

Galassi C, et al. (2025) An immunological mechanism of resistance to CDK4/6 inhibitors in HR+ breast cancer. Oncoimmunology, 14(1), 2520269.

Dunn MR, et al. (2024) A latent class assessment of healthcare access factors and disparities in breast cancer care timeliness. PLoS medicine, 21(12), e1004500.

Cortes DRE, et al. (2024) Modeling normal mouse uterine contraction and placental perfusion with non-invasive longitudinal dynamic contrast enhancement MRI. PloS one, 19(7), e0303957.

Pellizzari S, et al. (2023) Neuron populations use variable combinations of short-term feedback mechanisms to stabilize firing rate. PLoS biology, 21(1), e3001971.

Gorenberg EL, et al. (2022) Identification of substrates of palmitoyl protein thioesterase 1 highlights roles of depalmitoylation in disulfide bond formation and synaptic function. PLoS biology, 20(3), e3001590.

Illingworth CJR, et al. (2020) A de novo approach to inferring within-host fitness effects during untreated HIV-1 infection. PLoS pathogens, 16(6), e1008171.

Johnston WJ, et al. (2020) Nonlinear mixed selectivity supports reliable neural computation. PLoS computational biology, 16(2), e1007544.

Fisher J, et al. (2020) Neuroinflammation associated with scrub typhus and spotted fever group rickettsioses. PLoS neglected tropical diseases, 14(10), e0008675.

Jones MU, et al. (2020) Seizing opportunities for intervention: Changing HIV-related knowledge among men who have sex with men and transgender women attending trusted community centers in Nigeria. PloS one, 15(3), e0229533.

Robbins SJ, et al. (2020) Oral sex practices among men who have sex with men and transgender women at risk for and living with HIV in Nigeria. PloS one, 15(9), e0238745.

Peniche AG, et al. (2020) Efficacy of histamine H1 receptor antagonists azelastine and fexofenadine against cutaneous *Leishmania major* infection. *PLoS neglected tropical diseases*, 14(8), e0008482.

Li J, et al. (2020) Robust genome-wide ancestry inference for heterogeneous datasets: illustrated using the 1,000 genome project with 3D facial images. *Scientific reports*, 10(1), 11850.

Bovin MJ, et al. (2019) Using the World Health Organization Disability Assessment Schedule 2.0 to assess disability in veterans with posttraumatic stress disorder. *PloS one*, 14(8), e0220806.

Sero D, et al. (2019) Facial recognition from DNA using face-to-DNA classifiers. *Nature communications*, 10(1), 2557.

Hefti MM, et al. (2018) High-resolution temporal and regional mapping of MAPT expression and splicing in human brain development. *PloS one*, 13(4), e0195771.

Jenkins GJ, et al. (2018) Automatic characterization of stride parameters in canines with a single wearable inertial sensor. *PloS one*, 13(6), e0198893.

Song Y, et al. (2017) RB inactivation in keratin 18 positive thymic epithelial cells promotes non-cell autonomous T cell hyperproliferation in genetically engineered mice. *PloS one*, 12(2), e0171510.

Xu B, et al. (2017) The ESCRT-III pathway facilitates cardiomyocyte release of cBIN1-containing microparticles. *PLoS biology*, 15(8), e2002354.