

Resource Summary Report

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DoG picker

RRID:SCR_016655

Type: Tool

Proper Citation

DoG picker (RRID:SCR_016655)

Resource Information

URL: <https://omictools.com/dog-picker-tool>

Proper Citation: DoG picker (RRID:SCR_016655)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 18,2023. Software tool for general particle picking in the single-particle processing of unknown macromolecules. Reference free particle picker with ability to sort particles based on size or it can be used to bootstrap the creation of templates or training datasets for other particle pickers. Used to facilitate particle selection in single particle electron microscopy., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: DoG picker

Synonyms: Difference of Gaussians (DoG) picker, Difference of Gaussians Picker, Difference of Gaussians picker

Resource Type: data processing software, image processing software, software application, software resource

Defining Citation: [PMID:19374019](#)

Keywords: general, single, particle, picking, macromolecule, size, selection, electron, microscopy, image, transform

Funding: NCRR RR17573;
NCRR RR23093

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: DoG picker

Resource ID: SCR_016655

Alternate URLs: http://emg.nysbc.org/redmine/projects/appion/wiki/Appion_Home

Record Creation Time: 20220129T080331+0000

Record Last Update: 20260912T200019+0000

Ratings and Alerts

No rating or validation information has been found for DoG picker.

No alerts have been found for DoG picker.

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 [SPARC SAWG](#) .

Huang AY, et al. (2026) Somatic cancer variants enriched in Alzheimer's disease microglia-like cells drive inflammatory and proliferative states. *Cell*, 189(12), 3719.

Li SH, et al. (2026) Childhood immunological imprinting of cross-subtype antibodies targeting the hemagglutinin head domain of influenza viruses. *Cell host & microbe*, 34(5), 873.

Zhang P, et al. (2024) Design of soluble HIV-1 envelope trimers free of covalent gp120-gp41 bonds with prevalent native-like conformation. *Cell reports*, 43(8), 114518.

Mostofian B, et al. (2022) Continuum dynamics and statistical correction of compositional heterogeneity in multivalent IDP oligomers resolved by single-particle EM. *Journal of molecular biology*, 434(9), 167520.

Clausen TM, et al. (2020) SARS-CoV-2 Infection Depends on Cellular Heparan Sulfate and ACE2. *Cell*, 183(4), 1043.

Zhou T, et al. (2020) Structure-Based Design with Tag-Based Purification and In-Process Biotinylation Enable Streamlined Development of SARS-CoV-2 Spike Molecular Probes. *Cell reports*, 33(4), 108322.

Hashimoto H, et al. (2020) Structural Basis of Protein Arginine Methyltransferase Activation by a Catalytically Dead Homolog (Prozyme). *Journal of molecular biology*, 432(2), 410.

Gorman J, et al. (2020) Structure of Super-Potent Antibody CAP256-VRC26.25 in Complex with HIV-1 Envelope Reveals a Combined Mode of Trimer-Apex Recognition. *Cell reports*, 31(1), 107488.

Ringe RP, et al. (2020) Neutralizing Antibody Induction by HIV-1 Envelope Glycoprotein SOSIP Trimers on Iron Oxide Nanoparticles May Be Impaired by Mannose Binding Lectin. *Journal of virology*, 94(6).

Yoder N, et al. (2020) The His-Gly motif of acid-sensing ion channels resides in a reentrant 'loop' implicated in gating and ion selectivity. *eLife*, 9.

Bepler T, et al. (2019) Positive-unlabeled convolutional neural networks for particle picking in cryo-electron micrographs. *Nature methods*, 16(11), 1153.

Wang N, et al. (2019) Structural Definition of a Neutralization-Sensitive Epitope on the MERS-CoV S1-NTD. *Cell reports*, 28(13), 3395.

Turner HL, et al. (2019) Potent anti-influenza H7 human monoclonal antibody induces separation of hemagglutinin receptor-binding head domains. *PLoS biology*, 17(2), e3000139.

Yuan M, et al. (2019) Conformational Plasticity in the HIV-1 Fusion Peptide Facilitates Recognition by Broadly Neutralizing Antibodies. *Cell host & microbe*, 25(6), 873.

Dubrovskaya V, et al. (2019) Vaccination with Glycan-Modified HIV NFL Envelope Trimer-Liposomes Elicits Broadly Neutralizing Antibodies to Multiple Sites of Vulnerability. *Immunity*, 51(5), 915.

Zhu X, et al. (2019) Structural Basis of Protection against H7N9 Influenza Virus by Human Anti-N9 Neuraminidase Antibodies. *Cell host & microbe*, 26(6), 729.

He L, et al. (2018) HIV-1 vaccine design through minimizing envelope metastability. *Science advances*, 4(11), eaau6769.

Noreng S, et al. (2018) Structure of the human epithelial sodium channel by cryo-electron microscopy. *eLife*, 7.

Dingens AS, et al. (2018) Complete functional mapping of infection- and vaccine-elicited antibodies against the fusion peptide of HIV. *PLoS pathogens*, 14(7), e1007159.

Borst AJ, et al. (2018) Germline VRC01 antibody recognition of a modified clade C HIV-1 envelope trimer and a glycosylated HIV-1 gp120 core. *eLife*, 7.