

Resource Summary Report

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EndoMap

RRID:SCR_026690

Type: Tool

Proper Citation

EndoMap (RRID:SCR_026690)

Resource Information

URL: <https://endomap.hms.harvard.edu/>

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Description: Structural interactome viewer. Interactive database of endosomal protein-protein interactions identified by cross-linking mass spectrometry and modeled by AlphaFold multimer. Structural protein interactome of human early endosomes.

Synonyms: EndoMAP.v1

Resource Type: data or information resource, database

Defining Citation: [DOI:10.1101/2025.02.07.636106](https://doi.org/10.1101/2025.02.07.636106)

Keywords: Structural protein interactome, structural interactome viewer, endosomal protein-protein interactions, human early endosomes,

Funding: Aligning Science Across Parkinson ;
NINDS R01NS110395;
NIGMS RO1 GM132129;
Michael J Fox Foundation

Availability: Free, Freely available

Resource Name: EndoMap

Resource ID: SCR_026690

Record Creation Time: 20250401T060330+0000

Record Last Update: 20260729T044644+0000

Ratings and Alerts

No rating or validation information has been found for EndoMap.

No alerts have been found for EndoMap.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Gonzalez-Lozano MA, et al. (2025) EndoMAP.v1 charts the structural landscape of human early endosome complexes. Nature, 643(8070), 252.

Holzer LA, et al. (2019) The accuracy of digital templating in uncemented total hip arthroplasty. Archives of orthopaedic and trauma surgery, 139(2), 263.