

# Resource Summary Report

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## [imctools](#)

RRID:SCR\_017132

Type: Tool

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### Proper Citation

imctools (RRID:SCR\_017132)

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### Resource Information

**URL:** <https://github.com/BodenmillerGroup/imctools>

**Proper Citation:** imctools (RRID:SCR\_017132)

**Description:** Software Python package that implements preprocessing pipeline for imaging mass cytometry data. Can convert IMC raw files to tiff files that are used as inputs into CellProfiler, Ilastik, Fiji etc.

**Abbreviations:** IMCtools

**Synonyms:** imaging mass spectrometry tools

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

**Defining Citation:** [PMID:29605184](#)

**Keywords:** preprocessing, pipeline, imaging, mass, cytometry, data, convert, IMC, raw, file, TIFF

**Funding:** European Research Council ;  
NIDDK UC4 DK108132;  
PhosphonetPPM and MetastasiX SystemsX grant ;  
Roche Postdoctoral Fellowship ;  
SNSF Assistant Professorship grant ;  
Swiss National Science Foundation

**Availability:** Free, Available for download, Freely available

**Resource Name:** imctools

**Resource ID:** SCR\_017132

**Alternate URLs:** <https://bodenmillergroup.github.io/imctools/build/html/index.html>

**License:** copyright University of Zurich

**Record Creation Time:** 20220129T080333+0000

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## Ratings and Alerts

- Used for image processing operation by the Human Islet Research Network community. Contact(s): [Bernd Bodenmiller](#) - Human Islets Research Network <https://hirnetwork.org/>

No alerts have been found for imctools.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 25 mentions in open access literature.

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

Sun J, et al. (2026) Cinobufotalin reduces glioblastoma resistance to temozolomide by inhibiting the CCL5-mediated PI3K/Akt/mTOR signaling pathway. Translational cancer research, 15(3), 159.

Kuett L, et al. (2025) Distant Metastases of Breast Cancer Resemble Primary Tumors in Cancer Cell Composition but Differ in Immune Cell Phenotypes. Cancer research, 85(1), 15.

Mayer M, et al. (2025) Deciphering the cellular tumor microenvironment landscape in salivary gland carcinomas using multiplexed imaging mass cytometry. Journal of experimental & clinical cancer research : CR, 44(1), 288.

Egozi A, et al. (2023) Single-cell atlas of the human neonatal small intestine affected by necrotizing enterocolitis. PLoS biology, 21(5), e3002124.

Lien HE, et al. (2023) Single-cell profiling of low-stage endometrial cancers identifies low epithelial vimentin expression as a marker of recurrent disease. EBioMedicine, 92, 104595.

Weeratunga P, et al. (2023) Single cell spatial analysis reveals inflammatory foci of immature neutrophil and CD8 T cells in COVID-19 lungs. Nature communications, 14(1), 7216.

Bai Y, et al. (2023) Expanded vacuum-stable gels for multiplexed high-resolution spatial histopathology. Nature communications, 14(1), 4013.

- Scuiller Y, et al. (2023) YOUPI: Your powerful and intelligent tool for segmenting cells from imaging mass cytometry data. *Frontiers in immunology*, 14, 1072118.
- Strotton M, et al. (2023) Multielement Z-tag imaging by X-ray fluorescence microscopy for next-generation multiplex imaging. *Nature methods*, 20(9), 1310.
- Märtens K, et al. (2023) Rarity: discovering rare cell populations from single-cell imaging data. *Bioinformatics (Oxford, England)*, 39(12).
- Lu P, et al. (2023) IMC-Denoise: a content aware denoising pipeline to enhance Imaging Mass Cytometry. *Nature communications*, 14(1), 1601.
- van Eijs MJM, et al. (2023) Highly multiplexed spatial analysis identifies tissue-resident memory T cells as drivers of ulcerative and immune checkpoint inhibitor colitis. *iScience*, 26(10), 107891.
- Krijgsman D, et al. (2022) MATISSE: An analysis protocol for combining imaging mass cytometry with fluorescence microscopy to generate single-cell data. *STAR protocols*, 3(1), 101034.
- Xiao X, et al. (2022) Multiplexed imaging mass cytometry reveals distinct tumor-immune microenvironments linked to immunotherapy responses in melanoma. *Communications medicine*, 2, 131.
- Baars MJD, et al. (2021) MATISSE: a method for improved single cell segmentation in imaging mass cytometry. *BMC biology*, 19(1), 99.
- Zhang J, et al. (2021) Multiplex imaging reveals the architecture of the tumor immune microenvironment. *Cancer biology & medicine*, 18(4), 949.
- Stockton JD, et al. (2021) Complete response to neoadjuvant chemoradiotherapy in rectal cancer is associated with RAS/AKT mutations and high tumour mutational burden. *Radiation oncology (London, England)*, 16(1), 129.
- Deng Y, et al. (2021) *Helicobacter pylori* infection disturbs the tumor immune microenvironment and is associated with a discrepant prognosis in gastric de novo diffuse large B-cell lymphoma. *Journal for immunotherapy of cancer*, 9(10).
- Schlecht A, et al. (2021) Imaging mass cytometry for high-dimensional tissue profiling in the eye. *BMC ophthalmology*, 21(1), 338.
- Olaloye OO, et al. (2021) CD16+CD163+ monocytes traffic to sites of inflammation during necrotizing enterocolitis in premature infants. *The Journal of experimental medicine*, 218(9).