

# Resource Summary Report

Generated by SPARC SAWG on Sep 14, 2026

## ArrayPipe

RRID:SCR\_010934

Type: Tool

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### Proper Citation

ArrayPipe (RRID:SCR\_010934)

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### Resource Information

**URL:** <http://koch.pathogenomics.ca/cgi-bin/pub/arraypipe.pl>

**Proper Citation:** ArrayPipe (RRID:SCR\_010934)

**Description:** A flexible tool for visualizing and analyzing your two-colour microarray slides.

**Abbreviations:** ArrayPipe

**Resource Type:** analysis service resource, data analysis service, production service resource, service resource

**Keywords:** bio.tools

**Resource Name:** ArrayPipe

**Resource ID:** SCR\_010934

**Alternate IDs:** OMICS\_00744, biotools:arraypipe

**Alternate URLs:** <https://bio.tools/arraypipe>

**Record Creation Time:** 20220129T080301+0000

**Record Last Update:** 20260912T200206+0000

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### Ratings and Alerts

No rating or validation information has been found for ArrayPipe.

No alerts have been found for ArrayPipe.

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### Data and Source Information

## Usage and Citation Metrics

We found 15 mentions in open access literature.

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

Yi SJ, et al. (2021) The KDM4B-CCAR1-MED1 axis is a critical regulator of osteoclast differentiation and bone homeostasis. *Bone research*, 9(1), 27.

Hernández-Cervantes A, et al. (2020) A conserved regulator controls asexual sporulation in the fungal pathogen *Candida albicans*. *Nature communications*, 11(1), 6224.

Fujita M, et al. (2019) A TonB-dependent receptor constitutes the outer membrane transport system for a lignin-derived aromatic compound. *Communications biology*, 2, 432.

Znaidi S, et al. (2018) Systematic gene overexpression in *Candida albicans* identifies a regulator of early adaptation to the mammalian gut. *Cellular microbiology*, 20(11), e12890.

Priyadarshini S, et al. (2018) Role of murine macrophage in temporal regulation of cortisol- and serotonin-induced adipogenesis in pre-adipocytes when grown together. *Biology open*, 7(8).

Pradhan B, et al. (2017) Comparative efficacy analysis of anti-microbial peptides, LL-37 and indolicidin upon conjugation with CNT, in human monocytes. *Journal of nanobiotechnology*, 15(1), 44.

Bibollet-Bahena O, et al. (2017) A dual-strategy expression screen for candidate connectivity labels in the developing thalamus. *PloS one*, 12(5), e0177977.

Sundqvist M, et al. (2017) Elevated Mitochondrial Reactive Oxygen Species and Cellular Redox Imbalance in Human NADPH-Oxidase-Deficient Phagocytes. *Frontiers in immunology*, 8, 1828.

Tremblay N, et al. (2016) Spliceosome SNRNP200 Promotes Viral RNA Sensing and IRF3 Activation of Antiviral Response. *PLoS pathogens*, 12(7), e1005772.

Kim JM, et al. (2015) Cooperation between SMYD3 and PC4 drives a distinct transcriptional program in cancer cells. *Nucleic acids research*, 43(18), 8868.

Fitzgerald S, et al. (2015) Re-engineering cellular physiology by rewiring high-level global regulatory genes. *Scientific reports*, 5, 17653.

Cabral V, et al. (2014) Targeted changes of the cell wall proteome influence *Candida albicans* ability to form single- and multi-strain biofilms. *PLoS pathogens*, 10(12), e1004542.

Li AH, et al. (2010) Contrasting transcriptional responses of a virulent and an attenuated strain of *Mycobacterium tuberculosis* infecting macrophages. *PloS one*, 5(6), e11066.

Gong JH, et al. (2010) G-protein-coupled receptor independent, immunomodulatory properties of chemokine CXCL9. *Cellular immunology*, 261(2), 105.

Brown KL, et al. (2008) ROS-deficient monocytes have aberrant gene expression that correlates with inflammatory disorders of chronic granulomatous disease. *Clinical immunology (Orlando, Fla.)*, 129(1), 90.