

Resource Summary Report

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ArrayPipe

RRID:SCR_010934

Type: Tool

Proper Citation

ArrayPipe (RRID:SCR_010934)

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

Ratings and Alerts

No rating or validation information has been found for ArrayPipe.

No alerts have been found for ArrayPipe.

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 [NIF](#) .

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.