

LXXI REUNIÓN ANUAL DE LA SOCIEDAD ARGENTINA DE INMUNOLOGÍA

9 al 11 de noviembre de 2023 / San Luis



**LXXI REUNIÓN CIENTÍFICA ANUAL DE LA
SOCIEDAD ARGENTINA DE INMUNOLOGÍA (SAI)**

9 -11 de noviembre de 2023

Universidad Nacional de San Luis-San Luis

**LXXI ANNUAL MEETING OF THE
ARGENTINEAN SOCIETY OF IMMUNOLOGY (SAI)**

November 9 - 11, 2023

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EDITORES RESPONSABLES

Comisión Directiva SAI
María Silvia Di Genaro

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Camino del Macizo Central de San Luis
de Hebe Iriarte

Macizo de San Luis: cima de las Sierras de San Luis, a 2088m sobre el nivel del mar, con vistas inolvidables de cerros, quebradas, valles y pequeñas mesetas de altura. (Extraído de Ser Argentino.com).

Hebe Iriarte: Microbióloga, docente en el Área Microbiología e Inmunología de la Universidad Nacional de San Luis, personal técnica de apoyo de CONICET. Fotógrafa profesional, realiza fotografías de flora y fauna.



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85 (86) NOVEL, SENSITIVE, *IN VITRO* RADIOLABELING ASSAYS ALLOW THE MONITORING OF CYTOSOLIC VIMENTIN PROTEOFORMS BY SDS-PAGE IN NON-INFECTED, MYCOBACTERIAL-INFECTED AND TLR2-LIGATED THP-1 CELLS. POSSIBLE VIMENTIN ROLES IN MONOCYTE TO MACROPHAGE DIFFERENTIATION, INFECTION AND INFLAMMATION.

Cristian J. A. Asensio^{1,2,3}, Stefania Zampieri², Karim Zuppan², Corrado Guarnaccia², Alessandro Vindigni², Renato A. S. Oliveira², Jerry Saklatvala³, Francisco E. Baralle², Rodolfo C. García²

¹CONICET, Argentina. ²ICGEB, International Centre For Genetic Engineering and Biotechnology, Trieste, Italy. ³Kennedy Institute of Rheumatology, UK.

Discovering low-abundant macrophage proteins/proteoforms, altered in level/PTMs during intracellular bacterial infection and innate immune responses, needs sensitive proteome screening tools in electrophoretic gels, but outperforming dyes. AIMS: to search for, in human THP1 macrophage-like cells, cytosolic proteins reproducibly altered in a time-dependent and sustained manner, at days 1-4 post-infection with mycobacteria (live or killed). METHODS: The cytosolic fraction was obtained and used in novel, post-cell harvest, cell-free, *in vitro* radiolabeling (IVR) assays, allowing the covalent labeling of cytosolic proteomes with P-32. Labeled proteomes were separated in 1D/2D gels to detect bands/spots with altered labeling, normalizing them against total stained and total labeled proteomes. Proteins of interest were identified by MS and characterized. Bibliometric and bioinformatic studies were initiated to interpret findings in terms of PTMs, protein-protein interactions and possible roles of altered proteins and to plan how IVR might help future studies. RESULTS: in all 12 time-course infection experiments, cytosolic vimentin (VIM) was upregulated by infection in a time-dependent manner. In 3 monocytic- to-macrophage differentiation experiments (PMA-treated, non-infected), the VIM IVR increased during 4 days. We identified cytosolic kinases allowing detection of VIM with cleaved forms. Metabolic labeling in cell culture detected VIM profiles different to IVR. In WB, different antibodies and sera against other proteins often did bind non-specifically to VIM. So, to monitor minor cleavage/expression changes in VIM, IVR was more sensitive, quantitative and robust than WB. The literature indicated that VIM: **a)** is emerging as a multifunctional protein located in the perinuclear area, cytosol, endosomes, viral factories, cell surface, extracellular space and blood; **b)** has roles in auto-/xeno-/aggre-phagy, apoptosis, scaffolding of signaling complexes and in binding to DNA, RNA, phospholipids, O-GlcNAc, Rab7a, p62, HDAC6, MTOC, NFκB, NOD2, NLRP3, ERK; **c)** is a modulator of infectious, immune, autoimmune, inflammatory, cell stress, and fibrotic responses and is a target of toxins from many bacteria; **d)** has roles other than the cytoskeletal/mechanical by using different PTMs and by assembling as 1-, 2- and 4-mers, cages, and filamentous networks; **e)** Surface VIM binds many bacteria and viruses including SARS-Cov-2; **f)** Non-specific WB signals might depend on VIM-Fc and/or on citrullinated VIM-Ab interactions; **g)** VIM can be cleaved in cells. CONCLUSIONS: IVR helped detecting dynamic changes in cytosolic VIM levels, complementing WB. IVR would help to study VIM functional diversity, to correlate VIM alterations with those in binding partners, and to study VIM as biomarker or drug-target in cell infection and/or differentiation. Dissecting pro-infection and infection-restricting VIM roles will improve our knowledge of host-pathogen interaction complexity.