

Abstracts from the

IV JOINT MEETING OF THE BIOLOGY SOCIETIES OF ARGENTINA

(Cuarta Reunión Conjunta de Sociedades de Biología de la República Argentina)

XXXVII Annual Scientific Meeting of the Tucumán Biology Association XXIII Annual Scientific Meeting of the Córdoba Biology Society XXXVIII Annual Scientific Meeting of the Cuyo Biology Society Argentine Biology Society Rosario Biology Society Chilean Society of Reproduction and Development

September 9–15, 2020 Online edition

The abstracts have been revised and evaluated by a Scientific Committee prior to publication

SCIENTIFIC COMMITTEE

ASOCIACIÓN DE BIOLOGÍA DE TUCUMÁN

Dra. María Teresa Ajmat Dra. Patricia L. Albornoz

Dr. Mario Fortuna

Dra. Lucrecia Iruzubieta Villagra

Mag. Analía Salvatore
Dr. Federico Bonilla
Dra. Liliana I. Zelarayán
Dra. María Eugenia Pérez
Dra. Elisa Ofelia Vintiñi

SOCIEDAD DE BIOLOGÍA DE CÓRDOBA

Dra. Graciela Borioli Dra. Paola Boeris Dra. Cecilia Conde Dra. Marta Dardanelli Dra. Elena Fernández Dr. Leonardo Fruttero

Dra. Susana Genti-Raimondi Dr. Alejandro Guidobaldi

Dr. Edgardo Jofré Dra. Melina Musri

Dra. Graciela Panzetta-Dutari

Dr. Germán Robert Dra. Luciana Torre Dra. Cristina Torres

SOCIEDAD DE BIOLOGÍA DE CUYO

MENDOZA

Dra. Nora Arenas Dra. Silvia Belmonte Dra. Alejandra Camargo Dr. Diego Cargnelutti Dra. María Teresa Damiani Dra. María Inés Echeverría Dr. Carlos Gamarra-Luques

Vet. Paula Ginevro Dr. Diego Grilli Dr. Eduardo Koch Dra. Myriam Laconi Dr. Luis López

Dra. Alejandra Mampel Dr. Walter Manucha Dr. Ricardo Masuelli Dra. Marcela Michaut Dra. Adriana Telechea Dr. Roberto Yunes

SAN LUIS

Dra. Silvina Álvarez Dra. Cristina Barcia

Dra. María Eugenia Ciminari Dr. Juan Gabriel Chediack

Dr. Fabricio Cid Dra. Gladys Ciuffo

Lic. Óscar Córdoba Mascali Dra. María Esther Escudero

Dra. Susana Ferrari Dra. Lucia Fuentes

Esp. Mónica Laurentina Gatica Dra. Nidia Noemí Gomez

Dra. Marta Moglia Esp. Facundo Morales

Dra. Edith Pérez

Dra. María Verónica Pérez Chaca Dra. Hilda Elizabeth Pedranzani

Dra. Graciela Wendel Dra. Alba Edith Vega Dra. Liliana Villegas

SAN JUAN

Dra. Gabriela Feresín

SOCIEDAD DE BIOLOGÍA DE ROSARIO

Dra. Ariana Diaz

Méd. Vet. Melina Gay

Dra. Graciela Klekailo

Dra. Milagros López Hiriart

Dra. Stella Mattaloni

Dra. Nidia Montechiarini

Dra. Alejandra Peruzzo

Dr. Claudio Luis Pidone

Dra. Marta Posadas

Dra. Mariana Raviola

Dra. María Elena Tosello

Dra. Silvina Villar

SOCIEDAD ARGENTINA DE BIOLOGÍA

Dra. Fernanda Parborell

Dra. Débora Cohen

Dra. Griselda Irusta

Dra. Isabel María Lacau

Dra. Silvina Pérez Martínez

Dra. Evelin Elia

Dra. Clara I. Marín Briggiler

Dr. Leandro Miranda

Dr. Pablo Cetica

BIOCELL, Vol. 45, Suppl. 3, 2021 ISSN 0327-9545 ISSN 1667-5746 (online version)

A31

IDENTIFICATION OF POTENTIAL PROTEINS INVOLVED IN ANGIOGENESIS ASSOCIATED WITH CERVICAL CANCER USING PROTEOMICS AND BIOINFORMATICS APPROACHES

 $Valero\ V^I$, $Homann\ L^I$, $Carriere\ P^I$, $Novoa\ MB^I$, $Gentili\ C^I$, $Calvo\ N^I$ IINBIOSUR , $Dpto.\ de\ Biología$, $Bioquímica\ y\ Farmacia$, UNS-CONICET. E-mail: ncalvo@criba.edu.ar

Angiogenesis is the growth of blood vessels from the existing vasculature and is essential in the progression of cervical cancer (CC), the fourth most common tumor in women worldwide. This process studies in search of potential biomarkers and therapeutic targets since the endothelial cells that form the abnormal tumor vasculature are characterized by changes at the protein level when are regulated by tumor and microenvironmental factors. The objective of this work was to identify potential proteins involved in the response of endothelial cells to soluble factors released by tumor cells derived from CC, using proteomics and bioinformatics approaches. We previously observed that treatment with conditioned media from CC HeLa cells (TCMs) for 24 h increases the number of endothelial HMEC-1 cells. In this work, the proteome response of HMEC-1 cells was studied under these experimental conditions, performing a Label-Free quantitative (LFQ) mass spectrometry (MS) at the CEQUIBIEM Proteomics Center. Proteins were identified and quantified with the Proteome Discoverer software and the Uniprot database. Also, a more in-depth statistical study was performed using the Perseus software. Proteomic analysis revealed 26 proteins with increased expression levels in endothelial cells treated with TCM $(P \le 0.05)$. Then, to evaluate the biological characteristics of these proteins, they were classified using the PANTHER analysis tool, according to their molecular function and biological processes. As a result of this study, catalytic activity was the most represented molecular function (11/26), followed by binding (4/26). Respect to biological processes, proteins were mainly classified into cellular processes (12/26) and energy metabolism (11/26). This analysis suggests that factors released by tumor cells mainly increase the expression of proteins involved in metabolic processes in endothelial cells. Within these proteins, the probable ATP-dependent RNA helicase DDX47 showed the greatest magnitude of change (>2). DDX47 is related to rRNA processing and ribosome biogenesis, which are processes associated with cell proliferation and cancer progression. Furthermore, ribosomal activity is also a critical regulator of metabolism. These results highlight the use of Label-Free spectrometry and bioinformatics approaches in an initial phase of discovery of potential proteins involved in cancer and suggest the potential role of DDX47 in angiogenesis associated with CC.

A32

THE NATURAL FLAVONOID APIGENIN IS ACTIVE AGAINST Trypanosoma cruzi EPIMASTIGOTES

Cano R^1 , Cifuente D^2 , Sosa M^3 , Lozano $E^{1,4}$

¹Consejo Nacional de Investigaciones Científicas y Técnicas, Instituto de Medicina y Biología Experimental de Cuyo, Mendoza-Argentina. ²Consejo Nacional de Investigaciones Científicas y Técnicas, Instituto de Investigación en Tecnología Química—UNSL- San Luis-Argentina. ³Consejo Nacional de Investigaciones Científicas y Técnicas, Instituto de Histología y Embriología de Mendoza, Mendoza-Argentina. ⁴Universidad Nacional de Cuyo, Facultad de Ciencias Médicas, Instituto de Fisiología, Mendoza-Argentina. E-mail: elozano@mendoza-conicet.gob.ar

Trypanosoma cruzi is the causal etiologic agent of Chagas disease. In cultures, this parasite is mainly found in the epimastigote form and a low percentage in the infective form trypomastigote. The current chemotherapy against T. cruzi is insufficient because the available drugs, Nifurtimox and Benznidazole, have limited activity, and show toxic side effects in patients. Therefore, the "screening" of purified molecules from natural sources, mainly plant leaves has become an important tool for the fight against Chagas disease. Many natural compounds, extracted from native plants of Argentina, have been shown to be effective against the parasite. Among them, flavonoids are an important family of molecules that have been widely studied. In this work we analyze the effect of the natural flavonoid Apigenin (AGN) isolated from Larrea divaricata, on the growth of T. cruzi epimastigotes (strain Dm28c). AGN showed an antiproliferative effect on epimastigotes, even at low concentrations. This effect was irreversible even in the short term of exposure to the compound. AGN does not significantly affect the mitochondrial activity of the parasites, at all the concentrations tested (1, 5, and 10 μ g/mL) but alteration in ROS levels were observed when 5 and 10 μ g/mL of AGN were used. When we analyzed the ultrastructure of the parasites, we observed an increase in cytoplasmic vacuolization and the presence of structures that appear to be like "membrane blisters". From these results it is necessary to identify the molecular targets of the parasites for the action of this compound and to determine if AGN can affect the life cycle of T cruzi.

A33

BEHAVIOR OF RENAL HEK293 CELLS ON HYDROGELS BASED ON POLY-*N*-ISOPROPYL ACRYLAMIDE (PNIPAM)

 $\underline{Capella\ V^{l}}$, Liaudat AC^{2} , Rivero R^{l} , Barbero CA^{l} , Bosch P^{2} , Rivarola CR^{l} , Rodriguez N^{2} $\overline{}^{l}$ Institute of Research in Energy Technologies and Advanced Materials (IITEMA) – Chemistry Department. 2 Institute of Environmental Biotechnology and Health (INBIAS) – Molecular Biology Department and Faculty of Exact, Physical-Chemical and Natural Sciences, National University of Río Cuarto – National Council of Scientific and Technical Research (CONICET) E-mail: v-capella@exa.unrc.edu.ar

Synthetic or natural hydrogels have mechanical properties and physicochemical characteristics that simulate the properties of the extracellular matrix (EMC) of body tissues, providing an environment that mimics the native cellular milieus and allows cell growth and adhesion. Hydrogels based on poly-*N*-isopropylacrylamide (PNIPAM) and their co-polymers have attracted a great attention in the biomedical field due to their biocompatibility, smooth texture, and absence of cytotoxicity against several cell lines, characteristics that would allow tissue development. Based on this background, the aim of this study focused on evaluating the biocompatibility of PNIPAM and co-polymers surfaces with the HEK293 cell line, constituted by human embryonic kidney cells, under*in vitro* conditions. Cytotoxicity, genotoxicity, proliferation, adhesion, and cellular mitotic/fragmentation relation in contact with surfaces of PNIPAM and co-polymers with neutral or ionic characteristics in different proportions were carried out. The