Journal of Integrated

a methodological journal

OMICS

Editors-in-Chief

Carlos Lodeiro-Espiño

Florentino Fdez-Riverola

Jens Coorssen

Jose-Luís Capelo-Martínez

JIOMICS

Journal of Integrated OMICS

Focus and Scope

Journal of Integrated OMICS, JIOMICS, provides a forum for the publication of original research papers, preliminary communications, technical notes and critical reviews in all branches of pure and applied "-omics", such as genomics, proteomics, lipidomics, metabolomics or metallomics. The manuscripts must address methodological development. Contributions are evaluated based on established guidelines, including the fundamental nature of the study, scientific novelty, and substantial improvement or advantage over existing technology or method. Original research papers on fundamental studies, and novel sensor and instrumentation development, are especially encouraged. It is expected that improvements will also be demonstrated within the context of (or with regard to) a specific biological question; ability to promote the analysis of molecular mechanisms is of particular interest. Novel or improved applications in areas such as clinical, medicinal and biological chemistry, environmental analysis, pharmacology and materials science and engineering are welcome.

Editors-in-Chief

Carlos Lodeiro-Espiño, University NOVA of Lisbon, Portugal Florentino Fdez-Riverola, University of Vigo, Spain Jens R. Coorssen, Brock University, Ontario, Canada Jose-Luís Capelo-Martínez, University NOVA of Lisbon, Portugal

Regional editors

ASIA

Nelson Cruz Saudi Arabia

Europe

Gilberto Igrejas

University of Trás-os-Montes and Alto Douro, Life Sciences and Environmental School, Centre of Genetics and Biotechnology Department of Genetics and Biotechnology, 5001-801 Vila Real, Portugal

Randen Patterson

Center for Computational Proteomics, The Pennsylvania State University,

Jens R. Coorssen

Brock University, Ontario, Canada

South America

Carlos H. I. Ramos

Marco Aurélio Zezzi Arruda

University of Campinas - Unicamp

ChemistryInstitute - UNICAMP, Brazil

Associated editors

AFRICA

Saffaj Taouqif

Centre Universitaire Régional d'Interface, Université Sidi Mohamed Ben Abdallah, route d'Imouzzar-Fès, Morocco

ASIA

Amita Pal

Division of Plant Biology, Bose Institute, Kolkata, India **Ashish Gupta**

Centre of Biomedical Magnetic Resonance, SGPGIMS Campus, India Canhua Huang

The State Key Laboratory of Biotherapy, West China Hospital, Sichuan University, PR China

Ching-Yu Lin

Institute of Environmental Health, College of Public Health, National Taiwan University, Taipei, Taiwan

Chantragan Srisomsap

Chulabhorn Research Institute, Bangkok, Thailand

Debmalya Barh

Institute of Integrative Omics and Applied Biotechnology (IIOAB), India

Eiii Kinoshita

Department of Functional Molecular Science, Graduate School of Biomedical Sciences, Hiroshima University, Japan

Fan Cher

Institute of Genetics and Developmental Biology, Chinese Academy of Sciences (CAS), China

Ganesh Chandra Sahoo

BioMedical Informatics Center of Rajendra Memorial Research Institute of Medical Science (RMRIMS), Patna, India

Guangchuang Yu

Institute of Life & Health Engineering, Jinan University, Guangzhou, China **Hai-Lei Zheng**

School of Life Sciences, Xiamen University, China

Hsin-Yi Wu

Institute of Chemistry, Academia Sinica, Taiwan

Ibrokhim Abdurakhmonov

Institute of Genetics and Plant experimental Biology Academy of Sciences of Uzbekistan, Uzbekistan

Jianghao Sun

Food Composition and Method Development Lab, U.S. Dept. of Agriculture, Agricultural Research Services. Beltsville, USA

Juan Emilio Palomares-Rius

Forestry and Forest Products Research Institute, Tsukuba, Japan

Jung Min Kim

Liver and Immunology Research Center, Daejeon Oriental Hospital of Daejeon University, Republic of Korea

Kobra Pourabdollah

Razi Chemistry Research Center (RCRC), Shahreza Branch, Islamic Azad University, Shahreza, Iran

Krishnakumar Menon

Amrita Center for Nanosciences and Molecular Medicine, Amrita Institute of Medical Sciences, Kochi, Kerala, India

Mohammed Rahman

Center of Excellence for Advanced Materials Research (CEAMR), King Abdulaziz University, Jeddah, Saudi Arabia

Ningwei Zhao

Life Science & Clinical Medicine Dept.; Shimadzu (China) Co., Ltd

Poh-Kuan Chong

National University of Singapore, Singapore

Sanjay Gupta

Advanced Centre for Treatment, Research and Education in Cancer (ACTREC), Tata Memorial Centre, Kharghar, Navi Mumbai, India

Sanjeeva Srivastava

Indian Institute of Technology (IIT) Bombay, India

Suresh Kumar

Department of Applied Chemistry, S. V. National Institute of Technology, Gujarat, India

Toshihide Nishimura

Department of Surgery I, Tokyo Medical University, Tokyo, Japan

Vishvanath Tiwari

Department of Biochemistry, Central University of Rajasthan, India

Xuanxian Peng

School of Life Sciences, Sun Yat-sen University, Guangzhou, China

Youxiong Que

National Research & Development Center for Sugarcane, China Agriculture Research System(CARS), Fujian Agriculture & Forestry University, Republic of China

Yu Wang

Department of Pharmacology and Pharmacy, the University of Hong Kong,

Zhiqiang Gao

Department of Chemistry, National University of Singapore

AUSTRALIA AND NEW ZEALAND

Emad Kiriakous

Queensland University of Technology (QUT), Brisbane, Australia

Joëlle Coumans-Moens

School of Science and Technology, School of Medicine, University of New England, Australia

Maurizio Ronci

Mawson Institute, University of South Australia, Mawson Lakes, Australia

Michelle Colgrave

CSIRO Livestock Industries, St Lucia, Australia

Peter Hoffmann

Institute for Photonics & Advanced Sensing (IPAS), School of Chemistry and Physics, University of Adelaide, Australia

Valerie Wasinger

Bioanalytical Mass Spectrometry Facility, Mark Wainwright Analytical Centre, University of NSW, Australia

Wuiun Ma

Centre for Comparative Genomics, Murdoch University, Australia

EUROPE

AhmetKoc, PhD

Izmir Institute of Technology, Department of Molecular Biology & Genetics, Urla, İzmir, Turkey

Alejandro Gella

Department of Basic Sciences, Neuroscience Laboratory, Faculty of Medicine and Health Sciences, Universitat Internacional de Catalunya, Sant Cugat del Vallès-08195, Barcelona, Spain

Angelo D'Alessandro

Università degli Studi della Tuscia, Department of Ecological and Biological Sciences, Viterbo, Italy

Antonio Gnoni

Department of Medical Basic Sciences, University of Bari "Aldo Moro", Bari, Italy

Ana Varela Coelho

Instituto de Tecnologia Química e Biológica (ITQB) Universidade Nova de Lisboa (UNL), Portugal

Anna Maria Timperio

Dipartimento Scienze Ambientali Università della Tuscia Viterbo, Italy

Andrea Scaloni

Proteomics and Mass Spectrometry Laboratory, ISPAAM, National Research Council, via Argine 1085, 80147 Napoli, Italy

Angel P. Diz

Department of Biochemistry, Genetics and Immunology, Faculty of Biology, University of Vigo, Spain

Angela Chambery

Department of Life Science, Second University of Naples, Italy

Anna-Irini Koukkou

University of Ioannina, Department of Chemistry, Biochemistry Laboratory, Greece

António Sebastião Rodrigues

Departamento de Genética, Faculdade de Ciências Médicas, Universidade Nova de Lisboa,Portugal

Arzu Umar

Department of Medical Oncology, Laboratory of Breast Cancer Genomics and Proteomics, Erasmus Medical Center Rotterdam Josephine Nefkens Institute, Rotterdam, The Netherlands

Bart Devreese

Laborartory for Protein Biochemistry and Biomolecular Engineering, Department for Biochemistry and Microbiology, Ghent University, Belgium **Bernard Corfe**

Department of Oncology, University of Sheffield, Royal Hallamshire Hospital, United Kingdom

Bruno Manadas

Center for Neuroscience and Cell Biology, University of Coimbra, Portugal Carla Pinheiro

Plant Sciences Division, Instituto de Tecnologia Química e Biológica (ITOB), Universidade Nova de Lisboa, Portugal

Claudia Desiderio

Consiglio Nazionale delle Ricerche, Istituto di Chimica del Riconoscimento Molecolare (UOS Roma), Italy

Claudio De Pasquale

SAgA Department, University of Palermo, Italy

Celso Vladimiro Cunha

Medical Microbiology Department, Institute of Hygiene and Tropical Medicine, New University of Lisbon, Portugal

Christian Lindermayr

Institute of Biochemical Plant Pathology, Helmholtz Zentrum München, German Research Center for Environmental Health, Neuherberg, Germany

Christiane Fæste

Section for Chemistry and Toxicology Norwegian Veterinary Institute, Oslo, Norway

Christophe Cordella

UMR1145 INRA, Laboratoire de Chimie Analytique, Paris, France

Cosima Damiana Calvano

Universita' degli Studi di Bari, Dipartimento di Chimica, Bari, Italy

Daniela Cecconi

Dip. diBiotecnologie, LaboratoriodiProteomica e Spettrometriadi Massa, Universitàdi Verona, Verona, Italy

Deborah Penque

Departamento de Genética, Instituto Nacional de Saúde Dr Ricardo Jorge (INSA, I.P.), Lisboa, Portugal

Dilek Battal

Mersin University, Faculty of Pharmacy, Department of Toxicology, Turkey

Domenico Garozzo

CNR ICTP, Catania, Italy

Ed Dudley

Institute of Mass Spectrometry, College of Medicine Swansea University, Singleton Park, Swansea, Wales, UK

Elia Ranzato

Dipartimento di Scienze e Innovazione Tecnologica, DiSIT, University of Piemonte Orientale, Alessandria, Italy

Elisa Bona

Università del Piemonte Oientale, DISIT, Alessandria, Italy

Elke Hammer

Interfaculty Institute for Genetics and Functional Genomics, Ernst-Moritz-Arndt Universität, Germany

Enrica Pessione

University of Torino, Life Sciences and Systems Biology Department, Torino, Italy

Federica Pellati

Department of Life Sciences, University of Modena and Reggio Emilia, Italy François Fenaille

CEA, IBiTecS, Service de Pharmacologie et DImmunoanalyse (SPI), France

Fulvio Magni

Department of Health Science, Monza, Italy

Georgios Theodoridis

Department of Chemistry, Aristotle University, Greece

Gianfranco Romanazzi

Department of Environmental and Crop Sciences, Marche Polytechnic University, Italy

Giorgio Valentini

Università degli Studi di Milano, Dept. of Computer Science, Italy

Helen Gika

Chemical Engineering Department, Aristotle University of Thessaloniki, Greece

Hugo Miguel Baptista Carreira dos Santos

REQUIMTE-FCT Universidade NOVA de Lisboa, Portugal

Iñaki Álvarez

Institut de Biotecnologia i Biomedicina Vicent Villar Palasí, Universitat Autònoma de Barcelona, Barcelona

Jane Thomas-Oates

Centre of Excellence in Mass Spectrometry and Department of Chemistry, University of York, Heslington, UK

Jens Allmer

Molecular Biology and Genetics, Izmir Institute of Technology, Urla, Izmir, Turkey

Jesús Jorrín Novo

Agricultural and Plant Biochemistry, Proteomics Research Group, Department of Biochemistry and Molecular Biology, Córdoba, Spain

Johan Palmfeldt

Research Unit for Molecular Medicine, Aarhus University Hospital, Skejby, Aarhus, Denmark

Jose Câmara

University of Madeira, Funchal, Portugal

Juraj Gregan

Max F. Perutz Laboratories, University of Vienna, Austria

Karin Stensiö

Department of Photochemistry and Molecular Science, Ångström laboratory, Uppsala University, Sweden

Kay Ohlendieck

Department of Biology, National University of Ireland, Maynooth, Co. Kildare, Ireland

Konstantinos Kouremenos

Department of Chemistry, Umea University, Sweden

Luisa Brito

Laboratório de Microbiologia, Instituto Superior de Agronomia, Tapada da Ajuda, Lisbon, Portugal

Marco Lemos

GIRM & ESTM - Polytechnic Institute of Leiria, Peniche, Portugal

María Álava

Departamento de Bioquimica y Biologia Molecular y Celular, Facultad de Ciencias, Universidad de Zaragoza, Spain

María de la Fuente

Legume group, Genetic Resources, Mision Biologica de Galicia-CSIC, Pontevedra, Spain

Maria Gabriela Rivas

REQUIMTE/CQFB, Departamento de Química, Faculdade de Ciências e Tecnologia, Universidade Nova de Lisboa, Portugal

Marie Arul

Muséum National Histoire Naturelle, Département RDDM, Plateforme de spectrométrie de masse et de protéomique, Paris, France

Marie-Pierre Bousquet

Institut de Pharmacologieet de Biologie Structurale, UPS/CNRS, Tolouse, France

Mario Diniz

Dept. Química-REQUIMTE, Faculdade de Ciências e Tecnologia, Universidade Nova de Lisboa, Portugal

Martina Marchetti-Deschmann

Institute of Chemical Technologies and Analytics, Vienna University of Technology, Vienna, Austria

Maxence Wisztorski

University Lille 1, Laboratoire de Spectrométrie de Masse Biologique, Fondamentale & Appliquée, Villeneuve d'ascq, France

Michel Jaquinod

Exploring the Dynamics of Proteomes/Laboratoire Biologie à Grande Echelle, Institut de Recherches en Technologies et Sciences pour le Vivant, Grenoble, France

Mónica Botelho

Pantelis Bagos

Department of Computer Science and Biomedical Informatics, University of Central Greece, Greece

Patrice Francois

Genomic Research Laboratory, Service of Infectious Diseases, Department of Internal Medicine, Geneva

Patrícia Alexandra Curado Quintas Dinis Poeta

University of Trás-os-Montes and Alto Douro (UTAD), School of Agrary and Veterinary Sciences, Veterinary, Science Department, Portugal

Pedro Rodrigues

Centro de Ciências do Mar do Algarve, CCMAR, Faro, Portugal

Per Bruheim

Department of Biotechnology, Norwegian University of Science and Technology, Trondheim, Norway

Philipp Hess

Institut Universitaire Mer et Littoral(CNRS - Université de Nantes - Ifremer), Nantes, France

Pieter de Lange

DipartimentodiScienzedellaVita, SecondaUniversità degli Studi di Napoli, Caserta, Italy

Ralph Fingerhut

University Children's Hospital, Swiss Newborn Screening Laboratory, Children's Research Center, Zürich, Switzerland

Rubén Armañanzas

Computational Intelligence Group, Departamento de Inteligencia Artificial, Universidad Politécnica de Madrid, Spain

Ruth Birner-Gruenberger

Medical University Graz, Austria

Sebastian Galuska

Institute of Biochemistry, Faculty of Medicine, Justus-Liebig-University of Giessen, Germany

Serge Cosnier

Department of Molecular Chemistry, Grenoble university/CNRS, Grenoble, France

Serhat Döker

Cankiri Karatekin University, Chemistry Department, Cankiri, Turkey

Simona Martinotti

Dipartimento di Scienze e Innovazione Tecnologica, DiSIT, University of Piemonte Orientale, Alessandria, Italy

Spiros D. Garbis

Biomedical Research Foundation of the Academy of Athens, Center for Basic Research - Division of Biotechnology, Greece

Steeve Thany

Laboratoire Récepteurs et Canaux Ioniques Membranaires, UFR Science, Université d'Angers, France

Stefania Orrù

University if Naples Parthenope, Naples, Italy

Tâmara García Barrera

Departamento de Química y Ciencia de los Materiales, Facultad de Ciencias Experimentales, Universidad de Huelva, Spain

Vera Muccilli

 $Dipartimento di Scienze Chimiche, Universit\`a di Catania, Catania, Italy$

Yuri van der Burgt

Leiden University Medical Center, Department of Parasitology, The Netherlands

SOUTH AMERICA

Andréa P.B. Gollucke

Hexalab/Catholic University of Santos, Brazil

Arlindo Moura

Department of Animal Science - College of Agricultural Sciences - Federal University of Ceara, Fortaleza, Brasil

Bruno Lomonte

Instituto Clodomiro Picado, Universidad de Costa Rica

Edson Guimarães Lo Turco

São Paulo Federal University, Brasil

Fabio Ribeiro Cerqueira

Department of Informatics and NuBio (Research Group for Bioinformatics), University of Vicosa, Brazil

Fernando Barbosa

Faculty of Pharmaceutical Sciences of Ribeirão Preto University of São Paulo, Brazil

Mário Hiroyuki Hirata

Laboratório de Biologia Molecular Aplicado ao Diagnóstico, Departamento de Análises Clínicas e Toxicológicas, Faculdade de Ciências Farmacêuticas, Universidade de São Paulo, Brazil

Jan Schripsema

Grupo Metabolômica, Laboratório de Ciências Quimicas, Universidade Estadual do Norte Fluminense, Campos dos Goytacazes, Brazil

Jorg Kobarg

Centro Nacional de Pesquisa em Energia e Materiais, Laboratório Nacional de Biociências, Brazil

Rossana Arroyo

Department of Infectomic and Molecular Biology, Center of Research and Advanced Studies of the National, Polytechnical Institute (CINVESTAV-IPN), Mexico City, Mexico

Rubem Menna Barreto

Laboratorio de Biología Celular, Instituto Oswaldo Cruz, Fundaçao Oswaldo Cruz, Rio de Janeiro, Brazil

Vasco Azevedo

BiologicalSciencesInstitute, Federal University of Minas Gerais, Brazil

NORTH AMERICA

Amosy M'Koma

Vanderbilt University School of Medicine, Department of General Surgery, Colon and Rectal Surgery, Nashville, USA

Anthony Gramolini

Department of Physiology, Faculty of Medicine, University of Toronto, Canada

Anas Abdel Rahman

Department of Chemistry, Memorial University of Newfoundland and Labrador St. John's, Canada

Christina Ferreira

Purdue University - Aston Laboratories of Mass Spectrometry, Hall for Discovery and Learning Research, West Lafayette, US

Eustache Paramithiotis

Caprion Proteomics Inc., Montreal, Canada

Jagjit Yadav

Microbial Pathogenesis and Toxicogenomics, Laboratory, Environmental Genetics and Molecular, Toxicology Division, Department of Environmental

Health, University of Cincinnati College of Medicine, Ohio, USA

Jiaxu Li

Department of Biochemistry and Molecular Biology, Mississippi State University, $\ensuremath{\mathsf{USA}}$

Laszlo Prokai

Department of Molecular Biology & Immunology, University of North Texas Health Science Center, Fort Worth, USA

Madhulika Gupta

Children's Health Research Institute, University of Western Ontario London, ON, Canada

Michael H.A. Roehrl

Department of Pathology and Laboratory Medicine, Boston Medical Center Boston, USA

Olgica Trenchevska

Molecular Biomarkers, Biodesign Institute at Arizona State University, USA

Robert Powers

University of Nebraska-Lincoln, Department of Chemistry, USA

Susan Hester

United Stated Environmental Protection Agency, Durnam, USA

Thomas Kislinger

Department of Medical Biophysics, University of Toronto, Canada

William A LaFramboise

Department of Pathology, University of Pittsburgh School of Medicine Shadyside Hospital, Pittsburgh, USA

Xuequn Chen

Department of Molecular & Integrative Physiology, University of Michigan, Ann Arbor, USA

Ying Qu

Microdialysis Experts Consultant Service, San Diego, USA

JOURNAL OF INTEGRATED OMICS

A methodological Journal

ORIGINAL ARTICLES

Transformational Goal for Science Education: The 1 Student–1 Apparatus (1S1A) Model

1

ORIGINAL ARTICLES



JOURNAL OF INTEGRATED OMICS

A METHODOLOGICAL JOURNAL HTTP://www.jiomics.com



ORIGINAL ARTICLE | DOI: 10.5584/jiomics.v15i3.250

Transformational Goal for Science Education: The 1Student-1Apparatus (1S1A) Model

Inês F. Domingos ^{1,2}, André Q. Figueiredo^{1,2}, Hugo Miguel Santos^{1,2} Carlos Lodeiro^{1,2}, Orfeu Flores³, Gonçalo Doria³, Rodrigo Patricio^{3,4}, Ivo Lopes^{3,4}, Laura Mercolini⁵, Tereza Cartaxo⁶, Jerome Zoidakis^{7,8}, José Luis Capelo^{1,2,*}

¹LAQV-REQUIMTE, Department of Chemistry, NOVA School of Science and Technology, NOVA University of Lisbon, Lisbon, 2829-516 Caparica, Portugal;
²PROTEOMASS Scientific Society, Caparica, Portugal. ³STAB VIDA Lda, Caparica, Portugal. ⁴Department of Computer Science, NOVA, School of Science and Technology, NOVA University of Lisbon, 2829-516 Caparica, Portugal. ⁵Alma Mater Studiorum Università di Bologna, Bologna, Italy. ⁶University of Pernambuco, Institute of Biological Sciences – Recife (PE), Brazil. ⁷Department of Biology, National and Kapodistrian University of Athens, Greece. ⁸Department of Biotechnology, Biomedical Research Foundation, Academy of Athens, Greece.

Available Online: November 2025

ABSTRACT

Limited access to hands-on laboratory equipment remains a significant barrier to effective science education. To address this challenge, we evaluated the analytical performance of the Doctor Vida* Education platform - an ultra-compact, low-cost, multifunctional analytical device designed under the 'One Student–One Apparatus' (1S1A) model. Using the Bradford method, we quantified total protein in urine and serum samples and compared results against those obtained from a commercial CLARIOstar* microplate reader. Calibration curves constructed from eight independent replicates revealed comparable slopes and intercepts between the two systems, with Doctor Vida* Education device demonstrating high linearity and repeatability. Despite the CLARIOstar* achieving lower limits of detection and quantification, the Doctor Vida* Education device showed superior reproducibility, with consistently lower relative standard deviations across operators and experimental conditions. Statistical analysis of urine and serum measurements confirmed strong agreement between methods, with no significant differences in most samples and improved precision observed with Doctor Vida* Education device in serum analysis. With a unit cost below 1000 €, the Doctor Vida* Education device platform proves to be a reliable, robust, and accessible solution for individualized, competence-based learning in analytical sciences.

Keywords: 1S1A, Science Education, Protein Quantification, Urine, Serum, Bradford Method.

1. Introduction

A persistent challenge in science education is the limited access students have to practical, hands-on laboratory experiences. This gap between theory and experimentation often leads to disengagement and a superficial understanding of scientific concepts. In many educational systems, science is taught in abstract terms with limited infrastructure for experimentation, particularly in secondary and undergraduate settings [1], [2]. Addressing this issue requires innovative tools that are not only effective but also accessible and scalable across diverse educational contexts, including under-resourced schools and remote learning environments [3]. A key principle for improving science education is the 'one student–one apparatus' (1S1A) model, which ensures that each learner has direct, personal access to experimental tools. Educational research consistently shows that individualized

practical engagement enhances understanding, retention, and critical thinking skills [4], [5]. When students share equipment in overcrowded labs, their opportunities to practice and explore independently are limited, reducing the effectiveness of the learning experience [6]. In contrast, placing scientific instruments directly in each student's hands fosters ownership of the learning process and allows for repeated, self-paced experimentation [7].

The Doctor Vida® Education device - developed by STAB VIDA Lda, Caparica, Portugal - is a groundbreaking instrument designed with the 1S1A model in mind. Compact, portable, and battery operated, Doctor Vida® is built on energy-efficient LED technology and designed for ease of use in both formal and informal learning spaces. Measuring just a few centimeters in size and weighing less than 1 kg, the device is engineered for mobility, allowing students to perform real-time experiments whether in the classroom, in the field, or at home. Despite its modest size and eco-friendly

^{*}Corresponding author: José Luis Capelo, jlcm@fct.unl.pt

construction, Doctor Vida® Education supports a wide range of tasks, including fluorescence and spectrophotometry, PCR-based DNA amplification, colorimetric assays, and time-resolved data acquisition. Its built-in microcontroller and optical sensors enable high sensitivity and reliability, comparable to benchtop equipment used in conventional laboratories. Crucially, each unit costs less than 1000 €, making it a highly affordable solution for schools and institutions with limited resources. This low cost democratizes access to experimental science, enabling broader adoption in underserved educational settings. Taking advantage of modern digital and optical technologies in a compact format, the Doctor Vida® device represents a paradigm shift toward personalized, active, and inclusive science education. It empowers students to learn by doing and promotes scientific literacy by bridging the gap between theoretical knowledge and practical skills. This approach aligns with recent global educational frameworks emphasizing competence-based learning and equity in STEM access [8]. In this work, we present the analytical performance of Doctor Vida® Education as a compact device for the quantification of total

protein in urine and serum samples via the Bradford method.

2. Materials and Methods

2.1. Reagents and Protein Standards

Protein concentration was determined using the Bradford colorimetric assay with Bradford reagent (Sigma-Aldrich, B69168) [9]. Ultrapure water (18.2 m Ω ·cm $^{-1}$ at 25 °C) obtained from a Milli-Q* purification system (Merck Millipore) was used for all reagent preparations and dilutions. Bovine serum albumin (BSA) analytical standard (200 mg/mL stock; Sigma-Aldrich, P5369) was diluted in ultrapure water to obtain a 100 µg/mL working solution. This working standard was used to prepare a six-point calibration curve. All calibration solutions were prepared fresh on the day of analysis. Urine samples were taken from healthy volunteers from our lab, who signed an informed consent. Serum samples were acquired previously from Haematology service of Hospital Garcia de Orta (HGO) in Almada, Portugal, the same samples used in a previous study [10].

2.2. Calibration Curve Preparation

Tube 1 (T1) was prepared by combining 200 μ L of the BSA working standard (100 μ g/mL) with 600 μ L of ultrapure water, yielding a concentration of 25.0 μ g/mL. The remaining calibration standards (T2–T5) were prepared as two-fold serial dilutions by transferring 400 μ L from the preceding tube into 400 μ L of ultrapure water, resulting in concentrations of 12.5, 6.25, 3.13, and 1.56 μ g/mL, respectively. Tube 6 (T6), containing only ultrapure water, served as the 0 μ g/mL blank.

2.3. Urine and Serum Samples Preparation

Urine samples were taken from healthy volunteers from our lab, who signed an informed consent. Urine samples were prepared as two-fold serial dilutions in ultrapure water at final dilution factors of 1:2, 1:4, and 1:8. Serum samples were diluted in two steps: first to 1:50 by diluting 0.1 mL of serum to a final volume of 5 mL with ultrapure water, followed by a subsequent dilution to 1:100 by diluting 0.1 mL of the initial dilution to a final volume of 10 mL. This resulted in a final dilution factor of 1:5000 for serum samples. All diluted samples were prepared immediately before Bradford analysis.

2.4. Bradford Assay using the CLARIOstar® Microplate Reader

For microplate measurements, 150 μ L of each BSA standard or sample was combined with 150 μ L of Bradford reagent in flatbottom 96-well microplates (final volume: 300 μ L per well). After gentle mixing, the plate was incubated for 10 min at room temperature to allow full colour development. Absorbance at 595 nm was recorded using the CLARIOstar microplate reader (BMG LABTECH). All conditions were analysed in technical triplicate, and blanks containing Bradford reagent and ultrapure water were included for baseline correction.

2.5. Bradford Assay using the Doctor Vida® Spectrophotometric System

For cuvette-based measurements, 50 μ L of each BSA standard or sample was mixed with 50 μ L of Bradford reagent in Doctor Vida* disposable optical tubes (final volume: 100 μ L). After a 10-minute incubation at room temperature, absorbance at 595 nm was measured using the Doctor Vida* Education Spectrophotometric System (STAB VIDA Lda., Caparica, Portugal). All measurements were performed in technical triplicate with ultrapure water blanks for background subtraction.

2.6. Doctor Vida® Education device

Figure 1 shows an exploded schematic overview of the Doctor Vida® Education device. All analytical measurements were performed using the Doctor Vida® Education device, a compact, battery-powered, and multifunctional instrument designed to support hands-on science education and low-cost field-based experimentation. The device integrates several analytical capabilities within a single portable platform, allowing for realtime, user-directed operation in diverse learning or research environments. The core structure of the device comprises a multilayer modular architecture, beginning with an external top cover featuring an integrated rotary encoder with push-button functionality, enabling manual control of settings and measurements. A colour LCD display is mounted on the side of the encoder for real-time output visualization. Power is supplied via a rechargeable lithium-ion battery, providing several hours of autonomous operation. A USB-C port supports both battery charging and wired communication. The device's onboard lowpower microcontroller unit (MCU) includes embedded Wi-Fi and Bluetooth connectivity, allowing wireless data transfer and potential integration with mobile or cloud-based applications.

Doctor Vida® education (Exploded view) Сар Top cover Rotary encoder & button **Analysis chamber** (Top section view) Temperature White LED controlled LCD Battery block Blue LED USB-C port Green LED Multi-spectral filter CMOS sensor Buzzer Low-power MCU Bottom cover (inc. Wi-Fi & Bluetooth Copyright © 2025 STAB VIDA Lda. All rights reserved.

Figure 1. Doctor Vida* Education device depicting the internal structure and analysis chamber composition.

At the heart of the system lies the analysis chamber, which includes:

- A Complementary metal oxide semiconductor (CMOS) optical sensor for light detection.
- A multi-spectral optical filter enabling wavelength-specific measurements.
- A temperature-controlled block, suitable for isothermal reactions or thermal cycling (e.g., PCR).
- White, blue, and green LEDs for excitation in fluorescence and absorbance-based assays.
- This optical block allows the device to perform a range of standard laboratory techniques, including UV-VIS spectrophotometry, fluorimetry, and colorimetric assays. Additionally, its thermal control module supports nucleic acid amplification procedures.
- The bottom section houses the core processing and sensor electronics, securely mounted to a robust base plate that also contains a buzzer for audio feedback, enhancing user interaction during experimental procedures.

The total unit weighs under 1 kg and is small enough to be handheld or placed on a benchtop. All measurements in this study were conducted using individual units of the Doctor Vida* device *per* student, in alignment with the 1S1A pedagogical model.

3. Results and Discussion

3.1. Comparison of Analytical Performance

The analytical performance of both the Doctor Vida $^{\circ}$ Education and CLARIOstar devices was assessed based on the linear calibration curves generated from eight independent replicates for each instrument. The slope (m) and intercept (b) of the calibration lines were determined along with their respective standard deviations (mean \pm SD). For the Doctor Vida $^{\circ}$ Education device, the slope was $(1.13\pm0.14)\times10^{-2}$, while the intercept was $(7.7\pm5.6)\times10^{-3}$. In comparison, the CLARIOstar device showed a slope of $(1.00\pm0.16)\times10^{-2}$, and an intercept of $(6.2\pm5.8)\times10^{-3}$. These results demonstrate that both systems yield calibration curves with similar slopes and intercepts. The relatively low standard deviations in both parameters confirm the good repeatability and reliability of the calibration process across devices. Furthermore, the close

agreement between the slopes indicates comparable analytical sensitivity, supporting the potential use of Doctor Vida® Education for educational and screening applications where simplified instrumentation is required.

The comparison of limit detection (LOD) and limit of quantification (LOQ) obtained with the Doctor Vida® Education and CLARIOstar® systems reveals differences in analytical performance, as can be read in **Table 1**. The CLARIOstar® exhibited lower LOD and LOQ values, indicating higher instrumental

sensitivity. However, the Doctor Vida* Education showed lower variability across days, users, and calibration curves, reflected in its smaller RSD values for both LOD and LOQ, demonstrating a similar precision and robustness between devices. These results suggest that, while the CLARIOstar* is better suited for detecting very low analyte concentrations, the Doctor Vida* provides consistent and reproducible measurements as well, which is particularly relevant for applications requiring analytical stability and reduced dependence on operational conditions.

Table 1. Comparison of LOD and LOQ values for Doctor Vida® Education and CLARIOstar®.

Operator	Batch	Curve	Doctor Vida [®] LOD¹ (μg/mL)	Doctor Vida [®] LOQ ² (μg/mL)	CLARIOstar [®] LOD¹ (µg/mL)	CLARIOstar° LOQ² (μg/mL)
1	1	R1	3.6	12.2	0.9	3.1
	1	R2	5.1	16.9	1.0	3.2
	2	R1	4.4	14.8	0.9	3.0
	2	R2	4.9	16.3	0.8	2.8
	3	R1	3.9	12.9	1.3	4.3
	3	R2	4.1	13.8	1.1	3.7
2	4	R1	3.8	12.6	1.3	4.3
	4	R2	-	_	1.1	3.7
	5	R1	4.5	14.9	1.3	4.3
	3	R2	-	-	1.1	3.7
	Mean		4.3	14.3	1.1	3.6
SD			0.5	1.7	0.2	0.6
	RSD (%)		12.4	12.1	16.8	15.7

 1 Calculated as three times the standard deviation of the 10 blanks divided by the slope of the calibration curve (LOD = 3σ /S). 2 Calculated as 10 times the standard deviation of the 10 blanks divided by the slope of the calibration curve (LOD = 10σ /S).

3.2. Proof of concept

To assess the analytical performance of the proposed Doctor Vida® Education platform relative to an established reference system (CLARIOstar*), a series of comparative experiments were conducted using both urine and serum matrices. Statistical tests were applied to determine whether the new platform delivers results that are both accurate and precise when compared to the commercial system. This sections present the results of this initial validation, including F-tests to assess precision and t-tests to evaluate agreement in quantitative results. Each urine sample was analyzed in triplicate using both the CLARIOstar® (established method) and the Doctor Vida® Education (proposed method) as shown in **Table 2**. To evaluate the comparability between methods, an F-test was initially conducted to assess the homogeneity of variances for each sample. All calculated F-values were below the critical threshold ($F_{\text{critic}} = 19.0$; df = 2, 2), indicating that the assumption of equal variances was satisfied. Consequently, a twosample Student's t-tests was applied (P = 0.05, df = 4). No statistically significant differences were observed between the two methods for samples 1, 2, and 3 (t_{calc} < 2.776). In contrast, significant differences were detected in samples 4 and 5 (t = 5.01

and 4.47, respectively), which may reflect enhanced sensitivity or sample-specific variability influencing the performance of the new method. Serum samples were also analyzed in triplicate using both methods, as shown in Table 3. The average concentrations obtained by Doctor Vida® Education and CLARIOstar® were broadly consistent across all samples. F-test results revealed markedly higher variances in the CLARIOstar® measurements, with F-values exceeding the critical threshold (F > 19) in all cases. Due to these results, an additional Welch's t-test was applied to confirm whether the average concentrations obtained have any statistically difference between them. According to Welch's t-test (P = 0.05, $df \sim 5$), no statistically significant differences were detected between methods in any of the samples. This indicates that Doctor Vida® Education consistently achieved better precision, as evidenced by lower relative standard deviations (%RSD < 2%). These results further support the reliability and precision of the Doctor Vida® Education platform for serum analysis. Nonetheless, additional studies involving a wider range of concentrations and sample types are warranted to fully establish its analytical robustness. Overall, the results demonstrate a high degree of agreement between methods, supporting the potential of Doctor Vida® Education as an affordable platform for laboratory experiments at the learning laboratory.

Table 2. Comparison of total protein concentrations in urine samples determined by Doctor Vida Education* and CLARIOstar* devices.

Sample Number	Doctor Vida [®] x ± SD(μg/mL) ^a	RSD (%)	CLARIOstar° x ± SD(μg/mL) ^a	RSD (%)	F^1	$t_{statistic}^2$
1	43.4 ± 0.7	1.6	42.7 ± 1.5	3.5	4.6	0.76
2	50.9 ± 0.7	1.3	54.2 ± 3.1	5.8	18.0	1.69
3	45.2 ± 2.6	5.7	39.6 ± 1.6	4.0	3.5	2.60
4	41.7 ± 0.7	1.7	36.2 ± 1.3	3.5	3.5	5.01*
5	42.6 ± 0.9	2.2	38.4 ± 1.1	2.9	1.5	4.47*

an = 3.1F-test for the comparison of standard deviations. Paired t -test, P=0.05, df=4, t_{critical}=2.78. *Statistically significant difference.

Table 3. Comparison of total protein concentrations in serum samples determined by Doctor Vida® Education and CLARIOstar®.

Sample Number	Doctor Vida® x±SD(mg/mL)a	RSD (%)	CLARIOstar° x±SD(mg/mL) ^a	RSD (%)	F^1	t _{experimental} (Welch's t-test) ²
6	64.1 ± 0.6	0.96	65.9 ± 4.7	7.19	69.4	0.97
7	64.2 ± 0.1	0.16	61.9 ± 3.1	5.08	900.0	1.77
8	63.9 ± 1.0	1.50	67.1 ± 4.6	6.92	25.0	1.63
9	76.0 ± 0.9	1.13	76.0 ± 5.7	7.51	44.4	0.003
10	62.6 ± 0.5	0.83	65.8 ± 3.9	5.92	64.0	2.05

a n= 3. ¹F-test for the comparison of standard deviations. ²| $t_{experimental}$ | value of Welch's t-test; df ~ 5 ; $t_{critical}$ =2.57 for P=0.05.

4. Conclusion

The results confirm the reliability and analytical suitability of the Doctor Vida® Education platform as a practical alternative to the commercial CLARIOstar® system for basic quantitative analysis. Although the CLARIOstar® exhibited lower detection and quantification limits - indicating higher instrumental sensitivity the Doctor Vida® Education demonstrated excellent robustness across operators, water baths, and calibration curves, as evidenced by its consistently lower RSD values. Statistical comparisons of urine and serum samples showed strong agreement between the two systems, with no significant differences in most cases and superior precision from Doctor Vida® Education device in serum analysis. Importantly, beyond its analytical capabilities, the Doctor Vida® Education stands out for its compact design, ease of use, and affordable cost, priced under 1000 € per unit. These features make it particularly well-suited for educational settings, where individual students can operate their own real analytical device. This promotes hands-on learning, fosters experimental independence, and enhances understanding of key concepts in quantitative analysis. In summary, the Doctor Vida® Education platform offers a reliable, accessible, and scalable solution for modern teaching laboratories, supporting the broader goal of democratizing access to analytical science tools.

Acknowledgements

PROTEOMASS Scientific Society is acknowledged by the funding provided to the Laboratory for Biological Mass Spectrometry Isabel Moura (#PM001/2024, #PM001/2019 and #PM003/2016). This work received support from Fundação para a Ciência e a Tecnologia and Ministério da Ciência, Tecnologia e Ensino Superior (FCT/MCTES) through the projects LA/P/0008/2020 DOI 10.54499/LA/P/0008/2020, UIDP/50006/2020 DOI 10.54499/UIDP/50006/2020 and UIDB/50006/2020 DOI 10.54499/UIDB/50006/2020. HMS acknowledges the Associate Laboratory for Green Chemistry – LAQV (LA/P/0008/2020) DOI 10.54499/LA/P/0008/2020 funded by FCT/MCTES for his research contract. AQF thanks FCT/MCTES for his PhD grant reference 2023.00528.BD. IFD thanks FCT/MCTES for his PhD grant reference 2024.00745.BD.

Funding

PROTEOMASS Scientific Society, #PM001/2019, #PM003/2016. Fundação para a Ciência e a Tecnologia/ Ministério da Ciência, Tecnologia e Ensino Superior (FCT/MCTES), LA/P/0008/2020 DOI 10.54499/LA/P/0008/2020, UIDP/50006/2020 DOI 10.54499/UIDP/50006/2020 and UIDB/50006/2020 DOI 10.54499/UIDB/50006/2020.

References

- [1] A. S. Pamungkas, S. Muslim, and K. Khaerudin, "Innovation of Mobile Science Laboratories as a Solution to Access to Practical Learning in Schools: A Case Study at an Education Quality Assurance Center in Banten, Indonesia," International Journal of STEM Education for Sustainability, vol. 5, no. 2, pp. 184–198, Aug. 2025, doi: 10.53889/ijses.v5i2.700.
- [2] K. Kroes, D. Lefler, A. Schmitt, and C. Supalo, "Development of Accessible Laboratory Experiments for Students with Visual Impairments," Journal of Science Education for Students with Disabilities, vol. 19, no. 1, pp. 61–67, Dec. 2016, doi: 10.14448/jsesd.09.0006.
- [3] Irene Uzezi Berezi, "Virtual Learning Environment: Redefining Higher Educational Delivery for Efficiency and Accessibility," International Journal of Educational Management, Rivers State University, vol. 1, no. 1, Jan. 2025.
- [4] T. M. Bhuttah, Q. Xusheng, M. N. Abid, and S. Sharma, "Enhancing student critical thinking and learning outcomes through innovative pedagogical approaches in higher education: the mediating role of inclusive leadership," Sci Rep, vol. 14, no. 1, p. 24362, Oct. 2024, doi: 10.1038/s41598-024-75379-0.
- [5] D. W. Johnson and R. T. Johnson, "Cooperative Learning: The

- Foundation for Active Learning," in Active Learning Beyond the Future, IntechOpen, 2019. doi: 10.5772/intechopen.81086.
- [6] J. Garcia, A. Y. Uluan, I. J. Barat, J. N. Lubay, I. Macagba, and H. Mahinay, "Lived Experiences of Science Major Students in the Absence of Laboratory Activities," American Journal of Education and Technology, vol. 1, no. 2, pp. 75–82, Sep. 2022, doi: 10.54536/ajet.v1i2.513.
- [7] K. T. Kotsis, "Significance of Experiments in Inquiry-based Science Teaching," European Journal of Education and Pedagogy, vol. 5, no. 2, pp. 86–92, Apr. 2024, doi: 10.24018/ejedu.2024.5.2.815.
- [8] C. Jackson et al., "Equity-Oriented Conceptual Framework for K-12 STEM literacy," Int J STEM Educ, vol. 8, no. 1, p. 38, Jun. 2021, doi: 10.1186/s40594-021-00294-z.
- [9] M. M. Bradford, "A rapid and sensitive method for the quantitation of microgram quantities of protein utilizing the principle of protein-dye binding," Anal Biochem, vol. 72, no. 1–2, pp. 248–254, May 1976, doi: 10.1016/0003-2697(76)90527-3.
- [10] I. F. Domingos et al., "Dithiothreitol-based protein equalisation in the context of multiple myeloma: Enhancing proteomic analysis and therapeutic insights," Talanta, vol. 279, p. 126589, Nov. 2024, doi: 10.1016/j.talanta.2024.126589.