Sabrina F.P. ROSA - Translation Professional

Ph.D. in Biological Sciences - <u>it.linkedin.com/in/sabrinarosa</u> Email: <u>sabrinarosa@me.com</u> Phone: +39.320.062.9031

PROFESSIONAL SUMMARY

Scientist with 10+ years of interdisciplinary and multicultural experience proposing expertise in: scientific writing, proofreading and editing of technical and scientific manuscripts, translation of technical and scientific documents.

SKILLS

- Languages: English, French, Italian
- Project Management
- Scientific & Technical Writing
- Editing & Proofreading

- Communication & Cultural Sensitivity
- Translation
- Organization (trainings, events)
- Molecular Biology & Biotechnology

PROFESSIONAL EXPERIENCE

Alpine mountain huts of Aosta Valley (Italy) Tourism officer / Small business manager / Seasonal worker

2016 - 2018

- Wrote winning project proposal for the management of an alpine hut and applied it
- Developed a survey to understand mountain tourism expectations

European Commission Joint Research Centre, Ispra (Italy) Scientific & Technical Project Officer

2013 - 2016

- Wrote standard operating procedures (SOP), guidelines, manuals, peer-reviewed manuscripts and book chapter
- Developed and optimized multi-target genetically modified organisms (GMO) detection tools to improve and harmonize GMO testing within the EU
- Organized and followed up EU-wide inter-laboratory projects to test the developed tools
- Planned and supervised the tools technology transfer to other laboratories
- Organized and supported training activities in the field of GMO analysis
- Conducted surveys and conferred with scientists from EU Member States to target research and technology development to actual needs

NYU Medical Center (Dept of Environmental Medicine), New York, NY, USA 2011 – 2012 Postdoctoral Fellow

- Wrote scientific grant and obtained own funding
- Developed and validated new molecular and cellular markers to assess polychlorinated biphenyls immunotoxicity

Mount Sinai School of Medicine (Dept of Liver Diseases), New York, NY, USA 2010 - 2011 Postdoctoral Fellow

Drafted manuscripts

- Presented oral and written data at international conferences and at monthly meetings with collaborators
- Developed zebrafish transgenic lines using molecular genetic engineering techniques with the aim to perform compound testing for a group of rare inherited diseases (congenital disorders of glycosylation (CDG))

Yale University (Dept of Ecology & Evolutionary Biology), New Haven, CT, USA 2004 - 2005 Research Assistant

- Provided assistance with the preparation of projects-related reports, manuscripts and presentations
- Managed the animal resources and genetic stocks of the laboratory for researchers

ADDITIONAL EXPERIENCE

- Doctoral research: Prepared and published manuscripts and thesis *
- M.Sc. research: Prepared and published manuscript (publication now used as reference for the ongoing captive breeding programs) **
- Translation and interpretation work for the Yale Law School Immigration Legal Services clinics (New Haven, USA)
- Probono work:
 - translation of documents on the occasion of the "International Year of Microcredit" (2005) organized by the United Nations
 - translation of touristic booklets
- *, ** See publication list

EDUCATION

Yale University (New Haven, CT, USA) & Free University of Brussels (Belgium) Ph.D. in Biological Sciences (Molecular Biology and Genetics)	2005 – 2010
Free University of Brussels (Belgium) M.Sc. in Molecular Biology (Population/Conservation Genetics)	2002 - 2004
Free University of Brussels (Belgium) B.Sc. ("Licence") in Sciences (Molecular Biology)	1998 – 2002

LIST OF PUBLICATIONS

Rosa S., Gatto F., Angers-Loustau A. Petrillo, M., Querci M., Kreysa J.: Development and Applicability of a Ready-to-Use Multi-Target Analytical System for GMO Screening (2016). Food Chemistry, 201:110-119

Gatto F., Bassani, N., **Rosa S.**, Lievens A. Brustio R., Kreysa J. & Querci M.: Semi-quantification of GM maize using ready-to-use RTi-PCR plates (2016). Food Analytical Methods. doi:10.1007/s12161-016-0609-0

Bonfini L., Angers-Loustau A., Petrillo M., Ciabatti I.M., Gatto F., Rosa S., Lievens, A., Kreysa J.: The European Union Reference Methods Database and Decision Supporting Tool for the Analysis of Genetically Modified Organisms: GMOMETHOD and JRC GMO-Matrix (2015). In book: Genetically Modified Organisms in Food, Edition: First Edition, Publisher: Elsevier, Editors: Ronald Ross Watson,

Angers-Loustau A., Petrillo M., Bonfini L., Gatto F., **Rosa S.**, Patak A., Kreysa J.: JRC GMO-Matrix: a web application to support Genetically Modified Organisms detection strategies (2014). BMC Bioinformatics, 15 (1):659

Cline A., Gao N., Flanagan-Steet H., Sharma V., **Rosa S**, Sonon R., Azadi P., Sadler, K.C., Freeze H.H., Lehrman, M.A., and Steet R.: A zebrafish model of PMM2-CDG reveals altered neurogenesis and a substrate-accumulation mechanism for N-linked glycosylation deficiency (2012). Mol.Biol.Cell (in press, doi:10.1091/mbc.E12-05-0411)

Chu J., Mir A., Gao N., **Rosa S.**, Monson C., Sharma V., Steet R., Freeze H., Lehrman M. and Sadler K.: A zebrafish model of congenital disorders of glycosylation with phospomannose isomerase deficiency reveals a developmental window for corrective mannose supplementation (2013). Disease Models and Mechanisms (in press, doi:10.1242/dmm.010116)

Rosa S.: Positional cloning of the allorecognition gene *alr1* in the cnidarian Hydractinia *symbiolongicarpus* (2010). Doctoral thesis. Accessible at: http://difusion.ulb.ac.be/vufind/Record/ULB-DIPOT:oai:dipot.ulb.ac.be:2013/210157/Holdings *

Rosa S., Powell A.E., Rosengarten R.D., Nicotra, M.L., Moreno M., Grimwood J., Lakkis F.G., Dellaporta S.L., and Buss L.W. (2010): *Hydractinia* allodeterminant *alr1* resides in an immunoglobulin superfamily-like gene complex. *Current Biology*, 20, 1122-1127 *

Steinfartz S., Glaberman S., Lanterbecq D., Russello M., **Rosa S.**, Hanley T.C., Marquez C., Snell H.L., Snell H.M., Gentile G., dell'Olmo G., Powell A.M., and Caccone A. (2009): Progressive colonization and restricted gene flow shape island-dependent population structure in Galápagos marine iguanas (*Amblyrhynchus cristatus*). *BMC Evolutionary Biology*, 9, 297

Rosa S., Monteyne D., and Milinkovitch M.C. (2009): Development of 10 highly polymorphic microsatellite markers in the vulnerable Galápagos land iguanas (genus *Conolophus*). *Molecular Ecology Resources*, 9, 376-379 **

°Tzika A.C., °Rosa S., Snell H.L., Snell H.M., Marquez C., Tapia W., Rassmann K., Gentile G., and Milinkovich M.C. (2008): Population genetics of Galápagos land iguana (genus *Conolophus*) remnant populations. *Molecular Ecology*, 17, 4943-4952 (°co-first authors) **

Poudyal M., Rosa S., Powell A.E., Moreno M., Dellaporta S.L., Buss L.W., and Lakkis F.G. (2007): Embryonic chimerism does not induce tolerance in an invertebrate model organism. *PNAS* 104 (11), 4559-4564

Rosa S., Milinkovitch M.C., Van Waerebeek K., Berck J., Oporto J., Alfaro-Sigueto J., VanBressem M.F., Goodall N., and Cassens I. (2005): Population structure of nuclear and mitochondrial DNA variation among South American Burmeister's porpoises (*Phocoena spinipinnis*). *Conservation Genetics*, **6**, 431-443