

Riccardo Bellazzi, is Full Professor of Bioengineering and Biomedical Informatics at the Department of Electrical, Computer and Biomedical Engineering of the University of Pavia. He is the director of the Laboratory of Medical Informatics "Mario Stefanelli" and the chair of the PhD program of Bioengineering and Bioinformatics, University of Pavia. Moreover, he leads Laboratory of biomedical informatics at the hospital "Salvatore Maugeri" in Pavia.

Last year he also became the Chair of the Interdepartmental Centre for Health Technologies (CHT) of the University of Pavia.

The scientific interests of Prof. Bellazzi are highly interdisciplinary and are aimed at applications of informatics to medicine and life sciences, comprising data mining, temporal data analysis, decision support, clinical research informatics.

Prof. Bellazzi has a wide and internationally recognized research activity.

In 2000 he founded the working group on "Intelligent Data Analysis and Data Mining" of the International Association of Medical Informatics (IMIA), of which he was chairman from 2000 to 2008. In 2009 he became a Fellow of the American College of Medical Informatics for his international achievement. He was Vice-President of IMIA in the period 2011-2014.

He has been track chairman of the conference of the American Association of Medical Informatics (AMIA) in 2015. He was Scientific Program Chairman of the World conference of Medical Informatics (Medinfo 2010) and Program Chairman of the 2007 Artificial Intelligence in Medicine (AIME 2007).

He is a member of the editorial board of the journals "Methods of Information in Medicine", "Journal of the American Medical Informatics Association", "International Journal of Biomedical Informatics", "Journal of Diabetes Science and Technology" and Associate Editor of the "Journal of Biomedical Informatics".

Finally, he is co-founder of the academic spin-off Biomeris, which implements software to support clinical research and Engenome, which is specialized on the analysis of Next Generation Sequencing data.