1. Record Nr. UNINA9910131968403321

Titolo	Suburbium 2 [[electronic resource]] : il suburbio di Roma dalla fine dell'eta monarchica alla nascita del sistema delle ville, 5.-2. secolo a.C. / / a cura di Vincent Jolivet ... [et al.]
Pubbl/distr/stampa	Roma, : Ecole française de Rome, 2009
Descrizione fisica	755 p., [2] fold. maps : ill
Collana	Collection de l'Ecole fran-caise de Rome ; ; 419
Altri autori (Persone)	JolivetVincent
Disciplina	937
Soggetti	Rome Antiquities Congresses
Lingua di pubblicazione	Italiano
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Papers presented to 2 meetings, Rome, 2004, 2005. Incl. 1 text in Eng. and 1 text in Fre. Rome (Lazio).
Nota di bibliografia	Includes bibliographical references.
Nota di contenuto	(http://www.ilibri.casalini.it/toc/09280707.pdf)

2. Record Nr. UNINA9910229834903321

Titolo	The digest : National Italian American Bar Association law journal
Pubbl/distr/stampa	Washington, D.C. : , : National Italian American Bar Association, , [1991] -
Descrizione fisica	1 online resource (volumes)
Soggetti	Law - United States Law - Italy Law Bar journals. Periodicals. Italy United States
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Periodico

3. Record Nr. UNINA9910220052303321

Autore	Alessandro Lagana
Titolo	Bioinformatics of Non-Coding RNAs with Applications to Biomedicine: Recent Advances and Open Challenges
Pubbl/distr/stampa	Frontiers Media SA, 2017
Descrizione fisica	1 online resource (95 p.)
Collana	Frontiers Research Topics
Soggetti	Biotechnology
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Sommario/riassunto	The recent discovery of small and long non-coding RNAs (ncRNAs) has represented a major breakthrough in the life sciences. These molecules add a new layer of complexity to biological processes and pathways by revealing a sophisticated and dynamic interconnected system whose structure is just beginning to be uncovered. Genetic and epigenetic aberrations affecting ncRNA gene sequences and their expression have been linked to a variety of pathological conditions, including cancer, cardiovascular and neurological diseases. Latest advances in the development of high throughput analysis techniques may help to shed light on the complex regulatory mechanisms in which ncRNA molecules are involved. Bioinformatics tools constitute a unique and essential resource for non-coding RNA studies, providing a powerful technology to organize, integrate and analyze the huge amount of data produced daily by wet biology experiments in order to discover patterns, identify relationships among heterogeneous biological elements and formulate functional hypotheses. This Research Topic reviews current knowledge, introduces novel methods, and discusses open challenges of this exciting and innovative field in connection with the most important biomedical applications. It consists of four reviews and six original research and methods articles, spanning the full scope of the Research Topic.