1. Record Nr. UNISA996386668703316

Autore	Buchanan George <1506-1582.>
Titolo	De prosodia libellus [[electronic resource] /] / authore G. Buchanano, Scoto
Pubbl/distr/stampa	Edinburgi, : Excudebat Thomas Brown, anno Dom. 1678
Descrizione fisica	61 [i.e. 16], [4] p. : ill. (woodcut)
Soggetti	Latin language - Metrics and rhythmics Borders (Type evidence)17th century.ScotlandEdinburgh Printers' devices (Printing)17th century.ScotlandEdinburgh
Lingua di pubblicazione	Latino
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	With a title page woodcut. Page 16 misnumbered 61. Copy has print show-through, affecting text; leaf A10 torn and stained. Reproduction of the original in the Aberdeen University Library.
Sommario/riassunto	eebo-0009

2. Record Nr. UNINA9910261140103321

Autore	Katrine L. Whiteson
Titolo	Virus Discovery by Metagenomics: The (Im)possibilities
Pubbl/distr/stampa	Frontiers Media SA, 2017
Descrizione fisica	1 online resource (216 p.)
Collana	Frontiers Research Topics
Soggetti	Microbiology (non-medical)
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Sommario/riassunto	Since the late 1800s, the discovery of new viruses was a gradual process. Viruses were described one by one using a suite of techniques such as (electron) microscopy and viral culture. Investigators were usually interested in a disease state within an organism, and expeditions in viral ecology were rare. The advent of metagenomics using high-throughput sequencing has revolutionized not only the rate of virus discovery, but also the nature of the discoveries. For example, the viral ecology and etiology of many human diseases are being characterized, non-pathogenic viral commensals are ubiquitous, and the description of environmental viromes is making progress. This Frontiers in Virology Research Topic showcases how metagenomic and bioinformatic approaches have been combined to discover, classify and characterize novel viruses.