

1. Record Nr.	UNINA9911079588903321
Titolo	In Senate of the United States, March 1, 1826. Mr. Benton, from the select committee, to whom was referred the proposition to amend the Constitution of the United States with respect to the appointment of senators and representatives to offices under the federal government, reported .
Pubbl/distr/stampa	[Washington, D.C.] : , : [publisher not identified], , 1826
Descrizione fisica	1 online resource (6 pages)
Collana	Senate document / 19th Congress, 1st session. Senate ; ; no. 52 [United States congressional serial set] ; ; [serial no. 126]
Altri autori (Persone)	BentonThomas Hart <1782-1858> (Jacksonian (MO))
Soggetti	Constitutional amendments Political ethics Presidents Government executives - Selection and appointment - United States Separation of powers Legislators - United States Legislative materials.
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Title from opening lines of text. Batch processed record: Metadata reviewed, not verified. Some fields updated by batch processes. FDLP item number not assigned.

2. Record Nr.	UNINA9911123356303321
Autore	Arnold Claus
Titolo	The Reception and Application of the Encyclical Pascendi : The Reports of the Diocesan Bishops and the Superiors of the Religious Orders Until 1914
Pubbl/distr/stampa	Venice, Italy : , : Edizioni Ca' Foscari - Digital Publishing, , 2017 ©2017
ISBN	88-6969-130-6
Edizione	[1st ed.]
Descrizione fisica	1 online resource (328 pages)
Collana	Studi Di Storia Series ; ; v.3
Disciplina	282.0904
Soggetti	Modernism (Christian theology) - History Encyclicals, Papal
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Nota di contenuto	Pages:1 to 25 -- Pages:26 to 50 -- Pages:51 to 75 -- Pages:76 to 100 -- Pages:101 to 125 -- Pages:126 to 150 -- Pages:151 to 175 -- Pages:176 to 200 -- Pages:201 to 225 -- Pages:226 to 250 -- Pages: 251 to 275 -- Pages:276 to 300 -- Pages:301 to 325 -- Pages:326 to 328.
Sommario/riassunto	This volume examines the reception and application of Pope Pius X's 1907 encyclical 'Pascendi dominici gregis,' which condemned modernism as a heresy. The book compiles and analyzes reports from diocesan bishops and religious order superiors worldwide, submitted between 1908 and 1914, detailing their implementation of the encyclical's measures. It provides insights into the global response to the modernist crisis within Roman Catholicism, exploring themes such as doctrinal enforcement, disciplinary actions, and the broader implications for the Church during the early 20th century. The research is based on archival materials from institutions such as the Vatican Secret Archives and the Congregation for the Doctrine of the Faith.

3. Record Nr.	UNINA9911138213203321
Titolo	Proceedings of the 3rd Asia-Pacific Bioinformatics Conference : Institute for Infocomm Research (Singapore), 17-21 January 2005 / / editors, Yi-Ping Phoebe Chen, Limsoon Wong
Pubbl/distr/stampa	London, : Imperial College Press Hackensack, NJ, : distributed by World Scientific, c2005
ISBN	9786611866822 9781281866820 1281866822 9781860947322 1860947328
Edizione	[1st ed.]
Descrizione fisica	1 online resource (401 p.)
Collana	Series on advances in bioinformatics and computational biology ; ; v. 1
Altri autori (Persone)	ChenYi-Ping Phoebe WongLimsoon <1965->
Disciplina	572.0285
Soggetti	Bioinformatics Biology - Data processing
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Description based upon print version of record.
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Preface; APBC 2005 Organization; CONTENTS; Contributed Papers; Keynote Papers; Invited Paper; Author Index
Sommario/riassunto	High-throughput sequencing and functional genomics technologies have given us a draft human genome sequence and have enabled large-scale genotyping and gene expression profiling of human populations. Databases containing large number of sequences, polymorphisms, and gene expression profiles of normal and diseased tissues in different clinical states are rapidly being generated for human and model organisms. Bioinformatics is thus rapidly growing in importance in the annotation of genomic sequences, in the understanding of the interplay between genes and proteins, in the analysis the genetic va