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| 1. Record Nr. | UNINA9910716104803321 |
| Titolo | Ninth annual report of the Federal Farm Loan Board, year ended December 31, 1925. March 18, 1926. -- Referred to the Committee on Banking and Currency and ordered to be printed |
| Pubbl/distr/stampa | [Washington, D.C.] : , : [U.S. Government Printing Office], , 1926 |
| Descrizione fisica | 1 online resource (53 pages) : tables |
| Collana | House document / 69th Congress, 1st session. House ; ; no. 272 [United States congressional serial set] ; ; [serial no. 8579] |
| Soggetti | Agricultural credit Banks and banking Farms Revenue Factory and trade waste Foreclosure Land banks Loans Financial statements Legislative materials. |
| Lingua di pubblicazione | Inglese |
| Formato | Materiale a stampa |
| Livello bibliografico | Monografia |
| Note generali | Appendices, p. 17. Batch processed record: Metadata reviewed, not verified. Some fields updated by batch processes. FDLP item number not assigned. |

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| 2. Record Nr. | UNINA9910483993103321 |
| Titolo | Comparative Genomics : International Workshop, RECOMB-CG 2010, Ottawa, Canada, October 9-11, 2010, Proceedings // edited by Eric Tannier |
| Pubbl/distr/stampa | Berlin, Heidelberg : , : Springer Berlin Heidelberg : , : Imprint : Springer, , 2010 |
| ISBN | 3-642-16181-2 |
| Edizione | [1st ed. 2010.] |
| Descrizione fisica | 1 online resource (XII, 303 p.) |
| Collana | Lecture Notes in Bioinformatics, , 2366-6331 ; ; 6398 |
| Altri autori (Persone) | TannierEric |
| Disciplina | 572.8629 |
| Soggetti | Life sciences Computer programming Computer science Bioinformatics Life Sciences Programming Techniques Theory of Computation Computational and Systems Biology |
| Lingua di pubblicazione | Inglese |
| Formato | Materiale a stampa |
| Livello bibliografico | Monografia |
| Note generali | Bibliographic Level Mode of Issuance: Monograph |
| Sommario/riassunto | The complexity of genome evolution has given birth to exciting challenges for computational biologists. A various range of algorithmic, statistical, mathem- ical techniques to elucidate the histories of molecules are developed each year and many are presented at the RECOMB satellite workshop on Comparative Genomics. It is a place where scientists working on all aspects of comparative genomics can share ideas on the development of tools and their application to relevant questions. This volume contains the papers presented at RECOMB-CG 2010, held on October 9–11 in Ottawa. The ?eld is still ?ourishing as seen from the papers presented this year: many developments enrich the combinatorics of genome rearrangements, while gene order phylogenies are becoming more and more - curate, thanks to a mixing of combinatorial and statistical principles, |

associated with rapid and thoughtful heuristics. Several papers tend to refine the models of genome evolution, and more and more genomic events can be modeled, from single nucleotide substitutions in whole genome alignments to large structural mutations or horizontal gene transfers.
