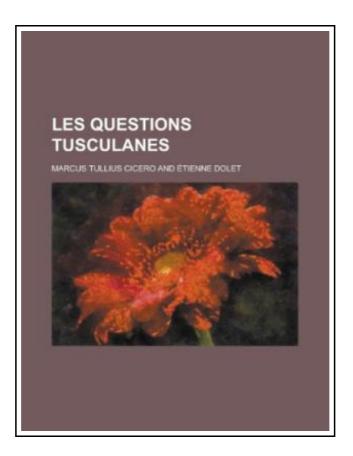
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Reviews

Extremely helpful to all of category of men and women. it had been writtern extremely completely and helpful. You are going to like the way the blogger compose this publication. (Johathan Haag)

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RareBooksClub. Paperback. Book Condition: New. This item is printed on demand. Paperback. 38 pages. Original publisher: Washington, D. C. : Executive Office of the President, National Science and Technology Council, 2004 OCLC Number: (OCoLC)233574204 Subject: Animal genome mapping. Excerpt: . . . value. The elucidation of these genomes will also benefit human medicine by identifying novel genes and their function that are critical for human health. It is important that the genetic and physical maps, and the reagents used to build them, are freely accessible and in the public domain. The bovine and swine genomics research communities need additional laboratory resources to allow more efficient use of current human and mouse mapping information to solve production inefficiency and health problems and contribute to increasing the global competitiveness of these livestock industries. Bovine BAC Map An international effort (International Bovine BAC Mapping Consortium; IBBMC) was initiated to develop a physical BAC-based map of the bovine genome. Clones from three publicly available bovine BAC libraries were fingerprinted and end-sequenced to develop the bovine BAC map. The three BAC libraries were generated from three common breeds, Holstein, Hereford, and Angus. A total of 295, 000 clones have been fingerprinted (15.8 X coverage) from these three libraries. Additional fingerprinting (100, 000 clones) was performed on a fourth BAC library by INRA (LInstitut National de la Recherche Agronomique), France and is being end-sequenced (12,000 clones to date). Overall, a total of 160, 000 clones have been sequenced with 140, 000 clones having both ends sequenced. End sequencing information from an additional 100, 000 clones by the INRA and international effort will be added and integrated in the near future. All sequence data are publicly available (www. ncbi. nih. gov Genbank). The...

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