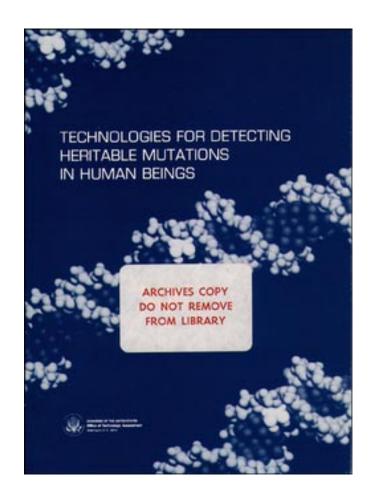
Technologies for Detecting Heritable Mutations in Human Beings

September 1986

NTIS order #PB87-140158



Recommended Citation:

U.S. Congress, Office of Technology Assessment, Technologies *for Detecting* Heritable *Mutations in* Human Beings, OTA-H-298 (Washington, DC: U.S. Government Printing Office, September 1986).

Library of Congress Catalog Card Number 86-600523

For sale by the Superintendent of Documents U.S. Government Printing Office, Washington, DC 20402

Foreword

Ensuring the health of future generations of children is of obvious importance to American society. Heritable mutations, permanent changes in the genetic material that can be passed on to succeeding generations, are the cause of a large but currently unquantifiable share of embryonic and fetal loss, disease, disability, and early death in the United States today. The methods now available to study heritable mutations, however, offer relatively little information about the kinds of mutations that can occur, their frequency, or their causes. Recent advances in molecular genetics have opened the door to new and innovative technologies that may offer a great deal more information about DNA. It may soon be possible to characterize mutations with exposures to specific mutagenic influences. While some of the new technologies are still *on* the drawing board, they are developing quickly and several of them may become available for wide-scale use in the next 5 to 10 years.

The Senate Committee on Veterans' Affairs, the House Committee on Science and Technology, and the House Committee on Energy and Commerce requested that OTA assess the available information about current and proposed means for detecting heritable mutations and on the likelihood and potential impact of such technological advances. These committees have wrestled with the problems of determining whether past exposures to potential mutagens have affected the health of Americans, and of framing reasonable public health laws, given current knowledge and technologies. This report summarizes OTA's findings as they relate to these issues.

An advisory panel, chaired by Arno G. Motulsky, provided guidance and assistance during the assessment. The OTA Health Program Advisory Committee, OTA staff, and scientific and policy experts from the private sector, academia, and the Federal Government provided information during the assessment and reviewed drafts of the report. We thank all who assisted us. As with all OTA reports, the content of the assessment is the sole responsibility of OTA and does not necessarily constitute the consensus or endorsement of the advisory panel or the Technology Assessment Board. Key OTA staff involved in the assessment were Michael Gough, Julie Ostrowsky, and Hellen Gelband,

John H. Libbou

JOHN H. GIBBONS Director

Advisory Panel for Technologies for Detecting Heritable Mutations in Human Beings

Arno G. Motulsky, Panel *Chair* Center for Inherited Diseases University of Washington School of Medicine

Richard J. Albertini Department of Medicine University of Vermont College of Medicine

Michael S. Baram School of Medicine and Public Health Boston University

Charles R. Cantor Department of Human Genetics and Development Columbia University College of Physicians and Surgeons

Dale Hattis Center for Policy Alternatives Massachusetts Institute of Technology

Ernest B. Hook Bureau of Maternal and Child Health New York State Department of Health

Alfred G. Knudson, Jr. Institute for Cancer Research Fox Chase Cancer Center

Nan M. Laird Department of Biological Statistics Harvard University School of Public Health

Mortimer L. Mendelssohn Lawrence Livermore National Laboratory Jeffrey H. Miller Department of Biology University of California at Los Angeles

James V. Neel Department of Human Genetics University of Michigan School of Medicine

Norton Nelson Department of Environmental Medicine New York University School of Medicine

Mark L. Pearson E.I. du Pent de Nemours & Co.

Richard K. Riegelman Department of Medicine George Washington University School of Medicine

Liane B. Russell Oak Ridge National Laboratory

Richard B. Setlow Brookhaven National Laboratory

William J. Schull The University of Texas Health Science Center

William G. Thilly Department of Nutrition and Food Science Massachusetts Institute of Technology

Richard M. Myers, Special Consultant University of California at San Francisco

NOTE: OTA appreciates and is grateful for the valuable assistance and thoughtful critiques provided by the advisory panel members. The panel does not, however, necessarily approve, disapprove, or endorse this report. OTA assumes full responsibility for the report and the accuracy of its contents.

OTA Project Staff—Technologies for Detecting Heritable Mutations in Human Beings

Roger C. Herdman, Assistant Director, OTA Health and Life Sciences Division

Clyde J. Behney, Health Program Manager

Michael Gough, *Project Director* Julia T. Ostrowsky, *Analyst*

Hellen Gelband, Analyst

Other Contributing Staff Cheryl M. Corsaro, Analyst' Virginia Cwalina, Administrative Assistant Carol Ann Guntow, Secretary/Word Processor Specialist Diann G. Hohenthaner, Word Processor/P. C. Specialist Eric Passaglia, Senior Distribution Specialist

Contractors

Elbert Branscomb, Lawrence Livermore National Laboratory Neal Cariello, Massachusetts Institute of Technology Leonard Lerman, Genetics Institute Harvey Mohrenweiser, University of Michigan Medical School Richard M. Myers, University of California at San Francisco Janice A. Nicklas, University of Vermont School of Medicine Maynard Olson, Washington University Medical School Cassandra Smith, Columbia University College of Physicians and Surgeons L.H.T. Van der Ploeg, Columbia University College of Physicians and Surgeons Diane K. Wagener, Washington, DC

^{&#}x27;On detail from the National Institutes of Health, April-June 1985.