

14.3 Studying the Human Genome

Lesson Objectives

- ☑ Summarize the methods of DNA analysis.
- ☑ State the goals of the Human Genome Project and explain what we have learned so far.

Lesson Summary

Manipulating DNA Since the 1970s, techniques have been developed that allow scientists to cut, separate, and replicate DNA base-by-base. Using these tools, scientists can read the base sequences in DNA from any cell.

- ▶ **Restriction enzymes** cut DNA into smaller pieces, called restriction fragments, which are several hundred bases in length. Each restriction enzyme cuts DNA at a different sequence of bases.
- ▶ **Gel electrophoresis** separates different-sized DNA fragments by placing them at one end of a porous gel, then applying an electrical voltage. The electrical charge moves the DNA.
- ▶ Using dye-labeled nucleotides, scientists can stop replication at any point along a single DNA strand. The fragments can then be separated by size using gel electrophoresis and “read,” base-by-base.

The Human Genome Project was a 13-year international effort to sequence all 3 billion base pairs in human DNA and identify all human genes. The project was completed in 2003.

- ▶ The researchers identified markers in widely separated strands of DNA.
- ▶ They used “shotgun sequencing,” which uses a computer to match DNA base sequences.
- ▶ To identify genes, they found promoters, exons, and other sites on the DNA molecule.
- ▶ To locate and identify as many haplotypes (collections of linked single-base differences) in the human population as possible, the International HapMap Project began in 2002.
- ▶ The Human Genome Project identified genes associated with many diseases and disorders. From the project came the new science of **bioinformatics**, the creation and use of databases and other computing tools to manage data. Bioinformatics launched **genomics**, the study of whole genomes.
- ▶ The human genome project pinpointed genes and associated particular sequences in those genes with numerous diseases and disorders. It also found that the DNA of all humans matches base-for-base at most sites, but can vary at 3 million sites.
- ▶ The 1000 Genomes Project, launched in 2008, will catalogue the variation among 1000 people.

Chapter 14 The Human Genome Section 1 Answer Key

Sylvia S. Mader



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Chapter Resource 11 Gene Technology Biology Holt Rinehart & Winston, Holt, Rinehart and Winston Staff, 2004

Human Genetics Ricki Lewis, 2008 Human Genetics Eighth Edition is a non science majors human genetics text that clearly explains what genes are how they function how they interact with the environment and how our understanding of genetics has changed since completion of the human genome project It is a clear modern and exciting book for citizens who will be responsible for evaluating new medical options new foods and new technologies in the age of genomics *Williams Obstetrics, 24th Edition, Study Guide* Robyn Horsager, Scott W. Roberts, Vanessa L. Rogers, Patricia C. Santiago-Muñoz, Kevin C. Worley, Barbara L. Hoffman, 2014-09-22 Assess your knowledge and retention of Williams Obstetrics with the only study guide keyed to that trusted text There is no better way to understand and remember the information found in Williams Obstetrics Twenty Fourth Edition than this unique study guide You will find everything you need for the most thorough assessment of your knowledge possible It is all here in one comprehensive book hundreds of full color images multiple choice questions keyed to the world s premier obstetrics textbook and clinical case questions to test your decision making skills and ability to apply the content to real clinical situations Here s why this is the best obstetrics review available More than 2 100 evidence based multiple choice questions carefully selected to highlight key points from each chapter in Williams Obstetrics Twenty Fourth Edition The answer key guides you to the pages in Williams Obstetrics Twenty Fourth Edition that contains the answers and further discussion the perfect way to strengthen your weak areas More than 450 color images are included as question material Organization follows the chronology of pregnancy from Maternal and Fetal Anatomy and Physiology to Labor and Delivery with additional sections on the Fetus and Newborn Puerperium Obstetric Complications and Medical and Surgical Complications Clinical case questions give your knowledge practical real world application The most detailed comprehensive and rigorously referenced text on obstetrics a true must have for anyone in the field **Pharmacotherapy A Pathophysiologic Approach 9/E** Joseph T. DiPiro, Robert L. Talbert, Gary C. Yee, Barbara G. Wells, L. Michael Posey, 2014-03-22 25 ADDITIONAL CHAPTERS AVAILABLE ONLINE THE LANDMARK TEXT ON HOW TO ACHIEVE OPTIMAL PATIENT OUTCOMES THROUGH EVIDENCE BASED MEDICATION THERAPY Pharmacotherapy A Pathophysiologic Approach is written to help you advance the quality of patient care through evidence based medication therapy derived from sound pharmacotherapeutic principles The scope of this trusted classic goes beyond drug indications and dosages to include the initial selection proper administration and monitoring of drugs You will find everything you need to provide safe effective drug therapy across the full range of therapeutic categories This edition is enriched by more than 300 expert contributors and every chapter has been revised and updated to reflect the latest evidence based information and recommendations Important features such as Key Concepts at the beginning of each chapter Clinical Presentation tables that summarize disease signs and symptoms and Clinical Controversies boxes that examine the complicated issues faced by

students and clinicians in providing drug therapy make this text an essential learning tool for students patient focused pharmacists and other health care providers NEW TO THIS EDITION A section on personalized pharmacotherapy appears in most sections All diagnostic flow diagrams treatment algorithms dosing guideline recommendations and monitoring approaches have been updated in full color to clearly distinguish treatment pathways New drug monitoring tables have been added Most of the disease oriented chapters have incorporated evidence based treatment guidelines that often include ratings of the level of evidence to support the key therapeutic approaches Twenty five online only chapters are available at www.pharmacotherapyonline.com

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Neuroproteomics Andrew K. Ottens, Kevin K.W. Wang, 2009-08-05 With the development of polypeptide ionization processes for mass spectrometry and the rapid processing of large scale datasets with bioinformatics researchers now have the tools necessary to propel neuroproteomics research to new levels In *Neuroproteomics Methods and Protocols* experts in the neurological and analytical sciences present experimental details for applying proteomics to the study of the central nervous system CNS and its dysfunction through trauma and disease Divided into four convenient sections this collection covers CNS animal models used for neuroproteomics research methods for separating and analyzing discrete subcomponents of the neuroproteome large scale approaches for CNS proteome characterization and quantification as well as methods that evaluate biofluids and translate neuroproteomic results into clinical platforms As a volume in the highly successful *Methods in Molecular Biology*™ series chapters include brief introductions to their respective topics lists of the necessary materials and reagents step by step readily reproducible laboratory protocols and notes on troubleshooting and avoiding known pitfalls Authoritative and cutting edge *Neuroproteomics Methods and Protocols* includes the clear how to information needed to critically assess what can be accomplished in the field and how to best perform and evaluate neuroproteomic experiments in one s own research

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