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PREFACE

The title of this book almost makes a preface superfluous. Genes is simply about genes, recognizing what amounts to a new field whose extraordinary progress has all but overwhelmed the traditional discipline of genetics. My aim is to cut through the enormous mass of information that has accumulated recently, to discern general principles and describe the state of the art in this exciting area. This text asks: what is a gene, how is it reproduced, how is it expressed, what controls its expression?

The underlying theme of this book is that the gene has at its disposal a vast repertoire of strategies for survival, different examples of which are displayed in various systems. Reflecting the perspective of current research, procaryotic and eucaryotic molecular biology are given equal weight. Both are now part of the same story. The starting point is the issue of how a gene is represented in protein; and from the protein, we work backward at the molecular level, as it were, to the DNA itself.

As a comprehensive introduction to the molecular biology of the gene, this book assumes no prior knowledge and is up to date with current research. I hope that this overview will make this rapidly advancing subject more accessible and readily allow readers to proceed to more advanced works. In view of the size of the relevant literature, it would only be confusing to rely on the citation of individual research articles. Each

chapter therefore concludes with a bibliography to suggest useful reviews and some research articles that lead more deeply into the subject.

I have tried to illustrate all important points diagramatically, and where appropriate the illustrations attempt to give some feeling for the scale and relationships of the elements involved. This preface would certainly be incomplete without acknowledgment of the considerable artistic endeavors of John Balbalis to realise this aim.

One of the pleasures of writing this book has been the ensuing discussions with my friends and colleagues who have commented on it. Many improvements have resulted from the generous efforts of Sankar Adhya, Sidney Altman, French Anderson, David Clayton, Nicholas Cozzarelli, Bernard Davis, Igor Dawid, Arg Efstratiadis, Nina Federoff, Alice Fulton, Joe Gall, Nicholas Gillham, Philip Hanawalt, Ira Herskowitz, Lee Hood, Joel Huberman, George Khoury, Nancy Kleckner, Marilyn Kozak, Charles Kurland, Art Landy, Jeffrey Miller, Masayasu Nomura, Charles Radding, Jeff Roberts, Rich Roberts, Gerry Rubin, Robert Schimke, David Schlessinger, David Shafritz, Phil Sharp, Allen Smith, Phang-C. Tai, Susumu Tonegawa, Harold Varmus, Alex Varshavsky, and Hal Weintraub. Finally, it hardly needs saying that the book would have been much less fun to write without the enthusiastic participation of my family. Benjamin Lewin

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GENES

BENJAMIN LEWIN

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Editor, Cell

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PART 1 THE NATURE OF GENETIC INFORMATION

THE STREETSHIE OF CHIPPMANERS ASTRONOMY

THE PASTIENT COMPONELIST P. O.

In calling the structure of the chromosome fibers a code-script we mean that the all-penetrating mind could tell from their structure whether the egg would develop, under suitable conditions, into a black cock or into a speckled hen, into a fly or a maize plant, a beetle, a mouse or a woman. . . But the term codescript is, of course, too narrow. The chromosome structures are at the same time instrumental in bringing about the development they foreshadow. They are law-code and executive power—or, to use another simile, they are architect's plan and builder's craft—in one.

GENERATION OF

ALLEGOA DIABELLA

Erwin Schrödinger, 1945

EV. NEW HOTTISEFSMART

CHAPTER 1 WHAT IS A GENE? A GENETIC VIEW

The concept of the gene has been the focus of some hundred years of work to establish the basis of heredity. It has been scrutinized from so many perspectives that there is no simple one-line answer to satisfy the question: what is a gene?

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A gene is a sequence of DNA that carries the information representing a protein. Until very recently this would have been an adequate (if incomplete) biochemical description. The sequence of DNA could be identified as a continuous stretch of nucleotides, related to the protein sequence by a colinear readout. But now it is clear that the sequence representing protein is not always continuous; it may be interrupted by sequences not concerned with specifying the protein. So genes may be in pieces that are put together during the process of gene expression.

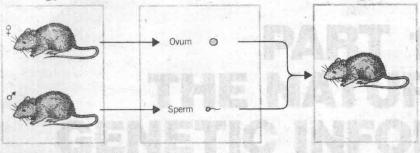
A gene can be identified as a cluster of mutations all of which prevent the production of the protein that it represents. Although this remains true, the properties of these clusters become more complex for interrupted genes. This issue has been moot, because most interrupted genes are found in situations in which detailed genetic analysis is not possible; but in principle, the genetic view of the gene now needs modification.

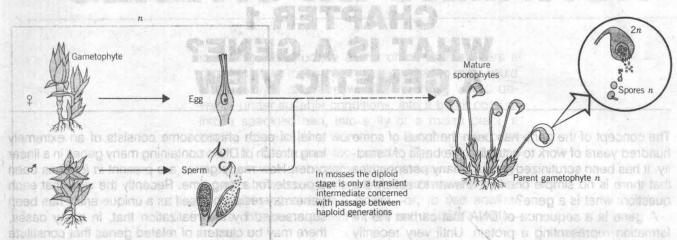
The large number of genes that make up the genome of any species are organized into a comparatively small number of chromosomes. The genetic ma-

terial of each chromosome consists of an extremely long stretch of DNA, containing many genes in a linear order. How many genes are present *in toto* has been a puzzle for a long time. Recently the view that each gene may reside by itself as a unique entity has been superseded by the realization that, in many cases, there may be clusters of related genes that constitute small families.

The genome generally has been viewed as rather stable, subject to changes in overall constitution and organization only on an extraordinarily slow evolutionary time scale. This contrasts with recent evidence that in some instances there may be rearrangements that occur regularly; and there may be components of the genome that are relatively mobile.

The concept of the gene has therefore undergone an evolution in which, although many of its traditional properties remain, exceptions have been found to show that none constitutes an absolute rule. Starting from the discovery of the gene as a fixed unit of inheritance, its properties were defined in terms of its residence at a definite position on the chromosome, this in turn leading to the view that the genetic material of the chromosome is a continuous length of DNA representing many genes. By tracing the development of these ideas, we can arrive at an operational description of the gene; although it is impractical to provide a capsule definition.





When eucaryotes perpetuate their genes through an alternation of diploid and haploid states, either type of state may provide the predominant "adult" type while the other is concerned solely with gamete and zygote formation. The mammals and the mosses are extreme examples

Red indicates diploid (2n) tissue; grey indicates haploid (n) tissue. that occur regularly; and there may be comboned to

A PARTICULATE FACTOR OF INHERITANCE

"The constant characters that appear in the group of plants may be obtained in all the associations that are possible according to the mathematical laws of combination. Those characters which are transmitted visibly, and therefore constitute the characters of the hybrid, are termed the dominant, and those that become latent in the process are termed recessive. The expression 'recessive' has been chosen because such characters withdraw or disappear entirely in the hybrids, but nevertheless reappear unchanged in their progeny in predictable proportions, without any essential alterations. Transitional forms were not observed in any experiment." All habit ad mad aften A

subudingo a sa baltimebi

This (abbreviated) quotation from Mendel's remarkable paper of 1865 introduces the fundamental concept of genetics: there is a unit of heredity consisting of some factor that is passed from parent to progeny. This particulate factor is, of course, what we now know as the gene. Mendel's work elucidated the general behavior of the gene in inheritance. Implicit in this is the concept that the organism is the gene's way of expressing and perpetuating itself, a point illustrated by the alternation of generations shown in Figure 1.1.

In striking contrast to the confusion of earlier thought

about heredity, Mendel's work established genetics as an experimental science through the introduction of several critical features. First, the initial set of plants was **true breeding** for the character under consideration: each gave rise to plants showing only that characteristic and none of the alternatives. Then the inheritance of only the one character was initially followed. With this established, the simultaneous behavior of two or more traits could be investigated. The experiments were placed on a quantitative basis by counting the numbers of each progeny type resulting from each mating.

The characteristics of the gene discernible from Mendel's work are summarized in what are known as his first and second laws. The first law deals with the properties of the individual gene. The organism has two copies of each gene: in modern terminology it is said to be **diploid**. Only one of the two copies is passed from parent to offspring through the **gametes** (sex cells). Thus when the gametes unite, the **zygote** (fertilized egg) gains one copy from each parent, restoring the situation in which every organism has one copy of paternal origin and one of maternal origin.

A gene may exist in alternative forms that result in the expression of a different characteristic (such as red versus white flower color). These forms are called **alleles.** The *law of independent segregation* states that these alleles do not affect each other when present in the same plant, but segregate unchanged by passing into different gametes when the next generation forms.

In a true-breeding organism, a **homozygote**, both alleles are the *same*. But a mating between two parents each of which is homozygous for a *different* allele generates a hybrid or **heterozygote**. If one allele is **dominant** and the other is **recessive**, the organism will have the appearance or **phenotype** only of the dominant type (so the heterozygote is indistinguishable from the true-breeding dominant parent).

But Mendel's first law recognizes that the genetic constitution or **genotype** of the hybrid comprises the presence of both alleles. This is revealed, as shown in **Figure 1.2**, when the hybrid is crossed with another hybrid to form the second generation. The critical point is that the alleles do not mix ("no transitional forms were observed"), but are physical entities whose interaction is at the level of expression.

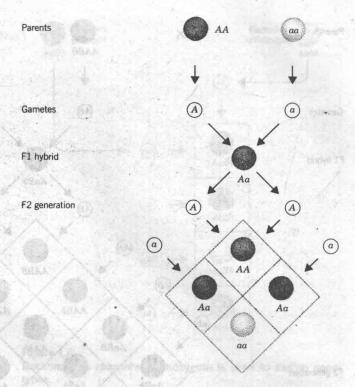


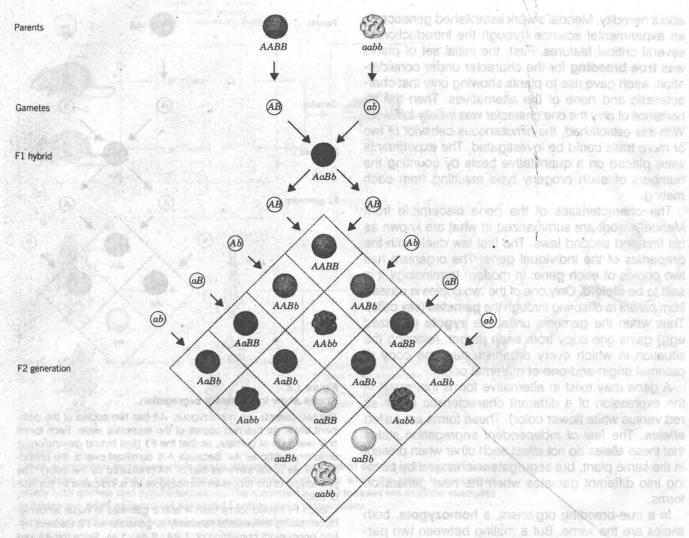
Figure 1.2
Alleles show independent segregation.

The two parents are homozygous: AA has two copies of the dominant allele; aa has two copies of the recessive allele. Each forms only one type of gamete, so that the **F1** (first hybrid generation) is uniformly hybrid as Aa. Because A is dominant over a, the phenotype of Aa is the same as that of AA (indicated by the color). The phenotype of the recessive homozygote aa is indicated by the lack of color.

Each F1 hybrid forms both A and a gametes in equal amounts. Upon mating these unite randomly to generate an **F2** (second hybrid generation) consisting of: 1 AA: 2 Aa: 1 aa. Since the AA and Aa have the same phenotype, this gives the classic 3: 1 ratio of dominant: recessive types.

In cases in which the heterozygote Aa has a phenotype intermediate between the parental AA and aa, the F1 would be distinct, and in the F2 the ratio of phenotypes would be 1 dominant: 2 intermediate: 1 recessive.

Although the characteristics studied by Mendel (fortunately) showed complete dominance, this is not necessarily always the case. Alleles may exhibit incomplete (partial) dominance or no dominance (sometimes known as codominance). In the latter case, the heterozygote is distinguished from the homozy-



officulty 1.2 when the hybri

Figure 1.3
Different genes assort independently.

One parent is homozygous for two dominant genes, A determining color, and B determining shape (shown by round structure). The other parent is homozygous for the recessive alleles a and b (characteristics shown by no color and wrinkled shape). The F1 is uniform with the dominant characteristics.

The F1 parents produce gametes in which there is independent segregation of alleles and independent assortment of genes, so that equal amounts are produced of each of the four possible types of gamete. These unite randomly to form 9 genotypic classes, which because of the dominance relationships appear as the four phenotypic classes: 9 colored-smooth: 3 colored-wrinkled: 3 non-colored-smooth: 1 noncolored-wrinkled.

Note that the same numbers are present of each reciprocal genotype; for example, the two parents (one each of *AABB* and *aaBb*) or the recombinant classes (one each of *AAbb* and *aaBB*). The 3:1 ratios are maintained for each individual segregating character.

The number of phenotypic classes will be greater if one or both of the characters is not dominant (so that heterozygotes appear different from either homozygote). It will be less if two genes affect a single characteristic. So if both A and B were needed for the production of color, a ratio of 9 colored: 7 noncolored would be seen in the F2.