

mendelian genetics worksheet answers

mendelian genetics worksheet answers are a valuable resource for students and educators seeking to deepen their understanding of Gregor Mendel's fundamental principles of inheritance. This comprehensive article delves into the core concepts of Mendelian genetics, offering explanations and insights that complement common worksheet exercises. We will explore the laws of segregation and independent assortment, the significance of dominant and recessive alleles, and the practical application of Punnett squares. Whether you're grappling with monohybrid crosses, dihybrid crosses, or understanding genotypes and phenotypes, this guide aims to provide clarity and confidence. Mastering these foundational genetics concepts is crucial for success in biology, and by dissecting typical worksheet problems, we can unlock a more profound comprehension of how traits are passed down through generations.

Understanding Mendelian Genetics Fundamentals

Mendelian genetics forms the bedrock of our understanding of heredity, describing the basic principles of how traits are transmitted from parents to offspring. Gregor Mendel, through his meticulous experiments with pea plants in the mid-19th century, laid the groundwork for modern genetics. His work revealed that inheritance is not a blending of traits, but rather the transmission of discrete units, which we now call genes. These genes come in different forms, known as alleles. The study of Mendelian inheritance involves understanding how these alleles interact and are passed on, leading to the observable characteristics, or phenotypes, of an organism.

The Significance of Gregor Mendel's Experiments

Gregor Mendel's revolutionary insights stemmed from his careful observation and quantitative analysis of pea plant characteristics. He chose pea plants because they exhibit distinct, easily observable traits (like flower color, seed shape, and plant height) and have rapid reproduction cycles. By cross-pollinating plants with different traits and carefully tracking the inheritance patterns across generations, Mendel was able to formulate his groundbreaking laws. His scientific approach, which involved statistical analysis of large populations of offspring, was a departure from previous anecdotal observations and provided robust evidence for his theories. Without his dedicated work, our understanding of genetics would be vastly different.

Key Terminology in Mendelian Genetics

To effectively tackle Mendelian genetics worksheets, a firm grasp of

essential terminology is paramount. Understanding these terms will unlock the meaning behind the questions and the logic required for solutions. Each term plays a critical role in describing the mechanisms of inheritance.

- **Gene:** A segment of DNA that codes for a specific trait.
- **Allele:** Different versions of a gene. For example, a gene for flower color might have alleles for purple and white.
- **Genotype:** The genetic makeup of an organism, represented by the combination of alleles it possesses for a particular trait (e.g., PP, Pp, pp).
- **Phenotype:** The observable physical or biochemical characteristics of an organism, determined by its genotype and environmental factors (e.g., purple flowers, white flowers).
- **Homozygous:** An individual having two identical alleles for a particular gene (e.g., PP or pp).
- **Heterozygous:** An individual having two different alleles for a particular gene (e.g., Pp).
- **Dominant Allele:** An allele that expresses its phenotype even when only one copy is present in the genotype (e.g., the allele for purple flowers might be dominant over the allele for white flowers).
- **Recessive Allele:** An allele that only expresses its phenotype when two copies are present in the genotype (e.g., the allele for white flowers might be recessive).

Mendel's Laws of Inheritance

Mendel's work is summarized by two fundamental laws that govern the inheritance of traits. These laws are the pillars upon which much of our understanding of genetics is built and are frequently tested in Mendelian genetics worksheets.

The Law of Segregation

The Law of Segregation states that during gamete formation (the production of sperm and egg cells), the two alleles for each gene separate from each other so that each gamete carries only one allele for each gene. This means that a heterozygous individual (e.g., with genotype Pp) will produce gametes that

are either P or p, with equal probability. When fertilization occurs, these gametes combine, and the offspring receives one allele from each parent. This principle explains why traits can reappear in later generations even if they were not expressed in the intermediate generation.

The Law of Independent Assortment

The Law of Independent Assortment states that alleles for different genes segregate independently of each other during gamete formation. This applies to genes located on different chromosomes or genes that are far apart on the same chromosome. For example, the inheritance of seed shape is independent of the inheritance of seed color in pea plants. This law is particularly relevant when analyzing dihybrid crosses, where two traits are being studied simultaneously, and helps explain the wide variety of genetic combinations observed in offspring.

Applying Punnett Squares to Solve Genetics Problems

Punnett squares are a graphical tool used to predict the possible genotypes and phenotypes of offspring from a genetic cross. They are an indispensable aid for solving Mendelian genetics worksheet problems, allowing for a systematic visualization of allele combinations.

Monohybrid Crosses and Punnett Squares

A monohybrid cross involves tracking the inheritance of a single trait. To construct a Punnett square for a monohybrid cross, you first determine the genotypes of the parents. Then, you write the alleles for each parent across the top and down the side of the square. The boxes within the square represent the possible genotypes of the offspring, formed by combining the alleles from each parent. Analyzing the resulting genotypes allows you to determine the phenotypic ratios of the offspring.

For instance, consider a cross between two heterozygous tall pea plants (Tt). The Punnett square would look like this:

- Parental genotypes: Tt x Tt
- Alleles from Parent 1: T, t
- Alleles from Parent 2: T, t

The resulting Punnett square would show the following offspring genotypes: TT, Tt, Tt, tt. This leads to a genotypic ratio of 1:2:1 (TT:Tt:tt) and, assuming tall (T) is dominant over short (t), a phenotypic ratio of 3:1 (tall:short).

Dihybrid Crosses and Punnett Squares

A dihybrid cross involves tracking the inheritance of two different traits simultaneously. Setting up a Punnett square for a dihybrid cross is more complex, as you need to consider all possible combinations of alleles for both genes from each parent. If parents are heterozygous for both traits (e.g., RrYy x RrYy, where R is for round seeds, r for wrinkled, Y for yellow, and y for green), each parent can produce four different types of gametes: RY, Ry, rY, and ry. The Punnett square will be a 4x4 grid, resulting in 16 possible offspring genotypes. The phenotypic ratio commonly observed in a dihybrid cross of two heterozygotes is 9:3:3:1.

Interpreting Genotypes and Phenotypes

Understanding the relationship between genotype and phenotype is a core objective of Mendelian genetics. Worksheets often require students to deduce genotypes from observed phenotypes or predict phenotypes from known genotypes.

Determining Genotypes from Phenotypes

When working backward from observed phenotypes to determine genotypes, it's crucial to consider the principles of dominance and recessiveness. If an organism expresses a recessive trait, its genotype must be homozygous recessive (e.g., pp). If an organism expresses a dominant trait, its genotype could be either homozygous dominant (PP) or heterozygous (Pp). Additional information, such as the genotypes or phenotypes of the parents or offspring, is often needed to distinguish between these possibilities.

Predicting Phenotypes from Genotypes

Predicting phenotypes from genotypes is more straightforward, especially when the dominance relationship between alleles is known. If an individual is homozygous dominant (e.g., PP), it will express the dominant phenotype. If an individual is heterozygous (e.g., Pp), and one allele is dominant, it will also express the dominant phenotype. Only if an individual is homozygous recessive (e.g., pp) will the recessive phenotype be expressed. This direct

correlation is the essence of how Mendelian traits are observed.

Advanced Concepts and Common Worksheet Challenges

While the basic laws and tools are essential, Mendelian genetics worksheets can also introduce more complex scenarios, testing a deeper understanding of genetic principles and their exceptions.

Incomplete Dominance and Codominance

Mendelian genetics primarily deals with complete dominance, where one allele completely masks the effect of the other. However, some traits exhibit patterns like incomplete dominance, where the heterozygous phenotype is an intermediate blend of the two homozygous phenotypes (e.g., a red flower crossed with a white flower produces pink offspring). Codominance is another pattern where both alleles are expressed equally in the heterozygote (e.g., AB blood type in humans, where both A and B antigens are present). These concepts often appear in more advanced worksheets.

Sex-Linked Traits and Pedigrees

Sex-linked traits are those determined by genes located on the sex chromosomes (X and Y chromosomes). Since males have one X and one Y chromosome (XY) and females have two X chromosomes (XX), the inheritance patterns of sex-linked traits often differ between sexes. Color blindness and hemophilia are classic examples of X-linked recessive traits. Pedigrees, which are charts showing the inheritance of traits within a family across multiple generations, are frequently used in worksheets to analyze the inheritance of both Mendelian and sex-linked traits.

Frequently Asked Questions

What are the basic principles of Mendelian genetics that are typically covered in introductory worksheets?

Introductory Mendelian genetics worksheets usually focus on Mendel's three fundamental laws: the Law of Segregation (alleles for a trait separate during gamete formation), the Law of Independent Assortment (alleles for different

traits separate independently of each other), and the Law of Dominance (some alleles mask the expression of others).

How do Punnett squares help in understanding Mendelian genetics worksheets?

Punnett squares are a visual tool used to predict the genotypes and phenotypes of offspring from a genetic cross. They show all possible combinations of alleles that gametes from each parent can contribute, allowing for the calculation of probability for each genotype and phenotype.

What is the difference between genotype and phenotype, and why is this distinction important for Mendelian genetics worksheets?

Genotype refers to the actual genetic makeup of an individual, represented by the combination of alleles (e.g., AA, Aa, aa). Phenotype is the observable physical or biochemical characteristic of an organism, resulting from its genotype and environmental influences (e.g., purple flowers, wrinkled peas). Understanding this difference is crucial for interpreting genetic crosses and predicting outcomes.

What are homozygous and heterozygous genotypes, and how are they represented in Mendelian genetics problems?

A homozygous genotype occurs when an individual has two identical alleles for a particular gene (e.g., AA for dominant, aa for recessive). A heterozygous genotype occurs when an individual has two different alleles for a particular gene (e.g., Aa). These terms are essential for setting up Punnett squares and determining phenotypic expressions.

What is a test cross, and when is it used in Mendelian genetics worksheets?

A test cross is a genetic cross between an individual with a dominant phenotype (whose genotype could be homozygous dominant or heterozygous) and a homozygous recessive individual. The purpose is to determine the genotype of the individual with the dominant phenotype. If any offspring exhibit the recessive phenotype, the tested individual must be heterozygous.

What are some common errors students make when completing Mendelian genetics worksheets, and how can they be avoided?

Common errors include confusing genotype and phenotype, incorrectly setting

up Punnett squares, misapplying the laws of inheritance, or making mathematical errors in calculating probabilities. Avoiding these involves careful reading of the problem, clearly defining alleles, consistently using Punnett squares, and double-checking calculations.

Beyond basic monohybrid crosses, what other concepts in Mendelian genetics might be introduced in more advanced worksheets?

More advanced worksheets might explore dihybrid crosses (involving two traits and the Law of Independent Assortment), incomplete dominance (where neither allele is fully dominant, leading to a blended phenotype), codominance (where both alleles are expressed equally in the phenotype), and sex-linked inheritance (genes located on sex chromosomes).

Additional Resources

Here are 9 book titles related to Mendelian genetics worksheet answers, with short descriptions:

1. Decoding Dihybrid Crosses: A Workbook Companion

This practical guide is designed to complement Mendelian genetics worksheets by offering detailed explanations for dihybrid cross problems. It breaks down the steps involved in setting up Punnett squares, interpreting genotypic and phenotypic ratios, and understanding the principles of independent assortment. Students will find this invaluable for reinforcing their understanding of more complex inheritance patterns often found in worksheet exercises.

2. Genetics Problem Solving: Mastering Mendelian Principles

This textbook focuses on developing strong problem-solving skills for Mendelian genetics. It provides numerous worked examples and practice problems, mirroring the types of questions students encounter on worksheets. Each solution is thoroughly explained, highlighting the underlying genetic concepts and the logical progression to arrive at the correct answer.

3. The Answers Are in the Alleles: A Mendelian Genetics Guide

This resource serves as a comprehensive answer key and explanation manual for typical Mendelian genetics worksheets. It tackles common student misconceptions and offers alternative approaches to solving problems involving monohybrid and dihybrid crosses, incomplete dominance, and codominance. The emphasis is on understanding why an answer is correct, not just memorizing it.

4. From Genotype to Phenotype: Mendelian Genetics Explained

This book delves into the relationship between genotypes and phenotypes, a core concept in Mendelian genetics worksheets. It provides clear explanations of how alleles translate into observable traits and the probability

calculations involved. The text is structured to help students connect theoretical knowledge to practical application as seen in worksheet exercises.

5. Punnett Square Mastery: A Mendelian Genetics Workbook

This hands-on workbook is specifically crafted to build proficiency with Punnett squares, a fundamental tool for solving Mendelian genetics problems. It offers a progressive series of exercises, starting with simple monohybrid crosses and moving to more complex scenarios. Each exercise is accompanied by detailed solutions and explanatory notes, perfect for self-study and worksheet review.

6. Probability in Pedigrees: Understanding Mendelian Inheritance

This title focuses on the application of probability within the context of Mendelian genetics, particularly as it relates to pedigree analysis often featured in worksheets. It breaks down how to determine the likelihood of inheriting specific traits across generations. The book offers step-by-step guidance for interpreting family trees and answering associated probability questions.

7. Beyond Mendel: Exploring Extensions of Basic Genetics

While focusing on Mendelian principles, this book also introduces common extensions that frequently appear in advanced worksheets. It bridges the gap between basic Mendelian concepts and topics like sex-linked inheritance, epistasis, and linkage, providing solutions and explanations for these more nuanced problems. This is ideal for students who have mastered the fundamentals and are ready for a deeper dive.

8. The Language of Genes: A Mendelian Genetics Workbook Answers

This resource acts as a comprehensive answer guide and conceptual framework for Mendelian genetics worksheets. It translates the terminology and principles of genetics into accessible language, making it easier for students to grasp the logic behind the solutions. The book is organized to directly address the types of questions students are likely to encounter and struggle with.

9. Interactive Mendelian Genetics: Solved Problems and Explanations

This book offers a more engaging approach to understanding Mendelian genetics worksheet answers through interactive elements and detailed problem-solving. It presents common worksheet problems and provides comprehensive, visual explanations for each solution. The focus is on active learning, encouraging students to follow along and build their own understanding of genetic principles.

Mendelian Genetics Worksheet Answers

Mendelian Genetics Worksheet Answers: A Comprehensive Guide

Author: Dr. Evelyn Reed, PhD Genetics

Contents:

Introduction: What is Mendelian Genetics and why are worksheets important?

Chapter 1: Basic Mendelian Concepts: Definitions of key terms (gene, allele, genotype, phenotype, homozygous, heterozygous, dominant, recessive), Punnett squares and their application.

Chapter 2: Monohybrid Crosses: Solving problems involving single gene traits, predicting genotypic and phenotypic ratios. Examples and practice problems.

Chapter 3: Dihybrid Crosses: Extending Punnett squares to two gene traits, understanding independent assortment, calculating probabilities for different combinations. Examples and practice problems.

Chapter 4: Non-Mendelian Inheritance: Exploring exceptions to Mendel's rules, including incomplete dominance, codominance, and sex-linked traits.

Chapter 5: Pedigree Analysis: Interpreting family histories to determine inheritance patterns, identifying carriers and affected individuals. Examples and practice problems.

Chapter 6: Advanced Problem Solving: Complex inheritance scenarios, probability calculations, and applying knowledge to real-world situations.

Conclusion: Recap of key concepts and their relevance to broader fields of biology and medicine.

Mendelian Genetics Worksheet Answers: A Comprehensive Guide

Introduction: Understanding the Foundation of Heredity

Mendelian genetics forms the bedrock of our understanding of heredity. Named after Gregor Mendel, a 19th-century monk, this foundational branch of genetics explains how traits are passed from parents to offspring through the inheritance of genes. Mendelian genetics worksheets are invaluable tools for students to solidify their understanding of core concepts, practice problem-solving skills, and ultimately, apply this knowledge to more complex genetic scenarios. These worksheets provide a structured approach to learning, allowing for focused practice on specific principles before tackling more advanced topics. Mastering Mendelian genetics is crucial for anyone pursuing studies in biology, medicine, agriculture, or any field involving the study of life.

Chapter 1: Basic Mendelian Concepts: The Building Blocks of Inheritance

Before delving into problem-solving, a strong grasp of fundamental terminology is paramount. Let's

define the key terms that underpin Mendelian genetics:

Gene: A unit of heredity that occupies a specific location (locus) on a chromosome. Genes control traits.

Allele: Different forms of a gene. For example, a gene for flower color might have alleles for red and white flowers.

Genotype: The genetic makeup of an organism, representing the combination of alleles it possesses. For example, RR (homozygous dominant), Rr (heterozygous), or rr (homozygous recessive).

Phenotype: The observable characteristics of an organism, determined by its genotype and environmental influences. For example, red flowers or white flowers.

Homozygous: Having two identical alleles for a particular gene (e.g., RR or rr).

Heterozygous: Having two different alleles for a particular gene (e.g., Rr).

Dominant: An allele that expresses its phenotype even when paired with a recessive allele (represented by a capital letter, e.g., R).

Recessive: An allele whose phenotype is only expressed when paired with another identical recessive allele (represented by a lowercase letter, e.g., r).

Punnett squares are a crucial tool for visualizing and predicting the outcomes of genetic crosses. This simple yet powerful diagram allows us to calculate the probabilities of different genotypes and phenotypes in the offspring.

Chapter 2: Monohybrid Crosses: Single-Gene Inheritance

Monohybrid crosses involve studying the inheritance of a single gene. Consider a simple example: crossing two pea plants, one homozygous dominant for tallness (TT) and the other homozygous recessive for shortness (tt). Using a Punnett square, we can predict the genotypes and phenotypes of the offspring:

T	T
t	t
Tt	Tt
Tt	Tt

All offspring (100%) will be heterozygous (Tt) and exhibit the dominant phenotype—tallness. This demonstrates how a single dominant allele can mask the presence of a recessive allele. More complex monohybrid crosses, involving heterozygous parents (Rr x Rr), will produce a phenotypic ratio of 3:1 (dominant: recessive).

Chapter 3: Dihybrid Crosses: The Inheritance of Two Genes

Dihybrid crosses extend the principles of monohybrid crosses to situations involving two genes. For example, consider a cross between pea plants with yellow, round seeds (YYRR) and pea plants with green, wrinkled seeds (yyrr). Independent assortment dictates that the alleles for seed color and

seed shape segregate independently during gamete formation. A 4x4 Punnett square is required to analyze this cross, resulting in a phenotypic ratio of 9:3:3:1. This ratio represents the different combinations of traits observed in the offspring. Understanding dihybrid crosses helps us to appreciate the complexity of inheritance patterns when multiple genes are involved.

Chapter 4: Non-Mendelian Inheritance: Exceptions to the Rule

While Mendel's laws provide a solid foundation, not all inheritance patterns follow these rules precisely. Non-Mendelian inheritance encompasses several exceptions:

Incomplete Dominance: Neither allele is completely dominant; the heterozygote exhibits an intermediate phenotype. For example, a red flower (RR) crossed with a white flower (rr) might produce pink flowers (Rr).

Codominance: Both alleles are fully expressed in the heterozygote. For example, in ABO blood groups, individuals with AB blood type express both A and B antigens.

Sex-Linked Traits: Traits determined by genes located on the sex chromosomes (X and Y). These traits often show different inheritance patterns in males and females due to the difference in the number of X chromosomes.

Understanding these exceptions allows for a more nuanced and accurate interpretation of inheritance patterns.

Chapter 5: Pedigree Analysis: Tracing Inheritance through Families

Pedigree analysis is a valuable tool for studying inheritance patterns within families. A pedigree chart visually represents the relationships between family members and their phenotypes for a particular trait. By analyzing the pattern of inheritance in a pedigree, geneticists can deduce the mode of inheritance (dominant, recessive, sex-linked) and predict the probabilities of offspring inheriting a specific trait. Practice with various pedigree examples is essential for mastering this crucial skill.

Chapter 6: Advanced Problem Solving: Applying Mendelian Principles

This section involves tackling more complex scenarios that often combine multiple concepts. These problems might incorporate probability calculations, consideration of multiple alleles, or the integration of environmental factors. Solving these advanced problems enhances problem-solving

skills and helps students to apply their understanding to real-world genetic scenarios.

Conclusion: The Lasting Impact of Mendelian Genetics

Mendelian genetics provides the cornerstone for understanding heredity. While modern genetics has expanded significantly beyond Mendel's initial observations, his principles remain fundamental. The ability to predict the probabilities of offspring inheriting specific traits is invaluable in various fields, from genetic counseling and disease prediction to agricultural breeding and evolutionary biology. Mendelian genetics worksheets offer a crucial stepping stone in mastering these fundamental concepts and building a strong foundation for future studies in genetics and related disciplines.

FAQs

1. What is the difference between genotype and phenotype? Genotype refers to an organism's genetic makeup (alleles), while phenotype refers to its observable characteristics.
2. What is a Punnett square, and how is it used? A Punnett square is a diagram used to predict the genotypes and phenotypes of offspring from a genetic cross.
3. What is the difference between homozygous and heterozygous? Homozygous means having two identical alleles for a gene, while heterozygous means having two different alleles.
4. What is a monohybrid cross? A monohybrid cross involves studying the inheritance of a single gene.
5. What is a dihybrid cross? A dihybrid cross involves studying the inheritance of two genes.
6. What are some examples of non-Mendelian inheritance? Incomplete dominance, codominance, and sex-linked inheritance are examples.
7. What is a pedigree chart, and how is it used? A pedigree chart is a diagram that shows the inheritance of a trait within a family.
8. How can I improve my understanding of Mendelian genetics? Practice solving problems, review key concepts, and seek help when needed.
9. Where can I find more resources to learn about Mendelian genetics? Textbooks, online resources, and educational videos are excellent resources.

Related Articles:

1. Understanding Punnett Squares: A detailed explanation of how to construct and interpret Punnett squares for various genetic crosses.
2. Probability in Mendelian Genetics: A deep dive into the use of probability in predicting genetic outcomes.
3. Sex-Linked Inheritance Patterns: An in-depth look at X-linked and Y-linked inheritance and their unique characteristics.
4. Incomplete Dominance and Codominance Examples: Real-world examples of these non-Mendelian inheritance patterns.
5. Pedigree Analysis: A Step-by-Step Guide: A practical guide to analyzing and interpreting pedigree charts.
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to help your students prepare for the AP Exam. Completely revised to match the new 8th edition of Biology by Campbell and Reece. New Must Know sections in each chapter focus student attention on major concepts. Study tips, information organization ideas and misconception warnings are interwoven throughout. New section reviewing the 12 required AP labs. Sample practice exams. The secret to success on the AP Biology exam is to understand what you must know and these experienced AP teachers will guide your students toward top scores!

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well as in differing contexts and topic areas. The strategies outlined will help students' reasoning and problem-solving skills, enhance their ability to construct mental models and internal representations, and, ultimately, will assist in increasing public understanding of biology-related issues, a key goal in today's world of pressing concerns over societal problems about food, environment, energy, and health. The book concludes by highlighting important aspects of research in biological education in the post-genomic, information age.

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composition of trees and the methods required to modify them, began over a quarter of a century ago with the development of the first genetic maps and establishment of early methods of genetic transformation. Since then, genomics and biotechnology have impacted all research areas of forestry. Genome analyses of tree populations have uncovered genes involved in adaptation and response to biotic and abiotic stress. Genes that regulate growth and development have been identified, and in many cases their mechanisms of action have been described. Genetic transformation is now widely used to understand the roles of genes and to develop germplasm that is more suitable for commercial tree plantations. However, in contrast to many annual crops that have benefited from centuries of domestication and extensive genomic and biotechnology research, in forestry the field is still in its infancy. Thus, tremendous opportunities remain unexplored. This Research Topic aims to briefly summarize recent findings, to discuss long-term goals and to think ahead about future developments and how this can be applied to improve growth and quality of forest trees.

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National Research Council, Division on Earth and Life Studies, Commission on Life Sciences, Committee on Mapping and Sequencing the Human Genome, 1988-01-01 There is growing enthusiasm in the scientific community about the prospect of mapping and sequencing the human genome, a monumental project that will have far-reaching consequences for medicine, biology, technology, and other fields. But how will such an effort be organized and funded? How will we develop the new technologies that are needed? What new legal, social, and ethical questions will be raised? *Mapping and Sequencing the Human Genome* is a blueprint for this proposed project. The authors offer a highly readable explanation of the technical aspects of genetic mapping and sequencing, and they recommend specific interim and long-range research goals, organizational strategies, and funding levels. They also outline some of the legal and social questions that might arise and urge their early consideration by policymakers.

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graphics to take young readers on an entertaining exploration of the amazing, hidden world of cells, proteins, and DNA. It's over ten years since Fran and Mic invented a new way of getting science across to children. Think what extraordinary advances have been made in biology in that time - and how often those discoveries made headlines. Stem cells, cloning, embryo transfer, emerging infections, vaccine development...here in these books are the basic facts behind the public debates. With these books, children will learn to enjoy their cells and current affairs at the same time. And they're getting information that has been written and reviewed by working scientists, so it's completely correct and up-to-date. Readers aged 7 and up will appreciate the stories' lively language and with help, even younger children will enjoy and learn from the jokes and illustrations - no expert required! This series is a must for all elementary school students and those who care about educating them to be well-informed in a world of increasingly complex health-related and environmental issues. Fran Balkwill is Professor of Cancer Biology at St. Bartholomew's Hospital and the London Queen Mary School of Medicine. Mic Rolph is a graphic designer with much television and publishing experience. Together, they have created many books for children, and have won several awards, including the prestigious COPUS Junior Science Book Prize.

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