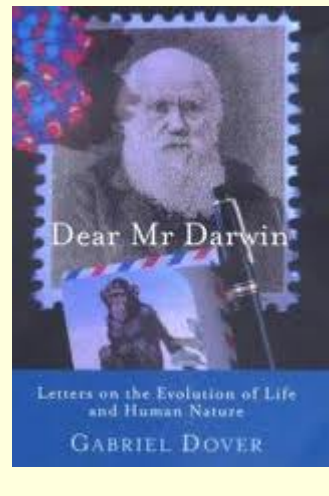


Molecular Drive: the Third Force in evolution

Gabriel Dover: Dear Mr Darwin

by Gert Korthof
updated: 4 Jun 2015. First published: 24 Mar 2001



Geneticist Gabriel Dover claims that there is a third force in evolution: 'Molecular Drive' beside natural selection and neutral drift. Molecular drive is operationally distinct from natural selection and neutral drift. According to Dover it explains biological phenomena, such as the 700 copies of a ribosomal RNA gene and the origin of the 173 legs of the centipede, which natural selection and neutral drift alone cannot explain.

Were Darwin and Mendel both wrong?

Molecular Drive is, according to Dover, an important factor in evolution, because it shapes the genomes and forms of organisms. Therefore Neo-Darwinism is incomplete without Molecular Drive. It is no wonder that the spread of novel genes was ascribed to natural selection, because it was the only known process that could promote the spread of novel genes. Dover doesn't reject the existence of natural selection but points out cases where natural selection clearly fails as a mechanism. Molecular drive is a non-Darwinian mechanism because it is independent of selection. We certainly need forces in evolution, since natural selection itself is not a force. It is the passive outcome of other processes. It is not an active process, notwithstanding its name. Conclusion: depending on how the theories are formulated, one should say either that orthodox-neo-Darwinism and orthodox-Mendelianism are 'refuted' and replaced or that both neo-Darwinism and Mendelianism are supplemented with additional mechanisms. Dover is clearly non-orthodox in the sense that he rejects the 'gene as the ultimate selfish unit of selection' view of Richard Dawkins, John Maynard Smith, Robert Trivers, William Hamilton, Edward O. Wilson and George C. Williams.

The dominance misunderstanding

Most students mistakenly believe that alleles are intrinsically either dominant or recessive, as did Mendel. But dominance is a relationship between alleles—no allele is dominant to another if its homozygous phenotype is also seen in the heterozygote.

Students also mistakenly think that dominant/recessive relationships are the norm. This is largely because almost all the alleles they see in their genetics course are presented in dominant/recessive pairs, with alternatives presented only as variants of or exceptions to dominance ('codominance' and 'incomplete dominance').

From: Rosemary J. Redfield *"Why Do We Have to Learn This Stuff?" – A New Genetics for 21st Century Students*, PLoS Biol 10(7). July 3, 2012.

Paramutation violates Mendel's first law

A paramutation is an interaction between two alleles at a single locus, whereby one allele induces a heritable epigenetical change in the other allele. This change is meiotically inheritable, and therefore paramutation violates Mendel's first law. (wiki) [added: 4 Oct 2012]

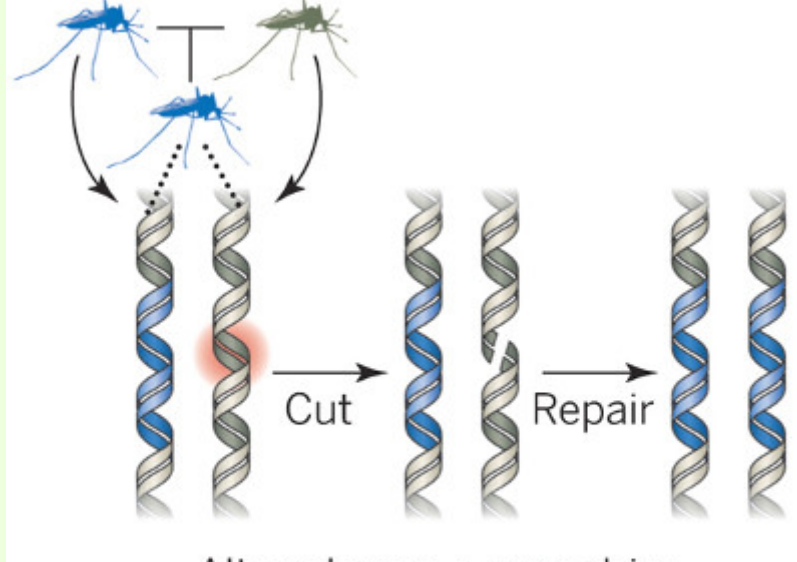
There is now evidence for RNA-directed non-Mendelian epigenetic inheritance ('paramutation') in plants and animals. [added: 15 Jun 2014]
John S Matick (2009) Has evolution learnt how to learn? EMOB Rep. Jul 2009; 10(7): 665.
V. L. Chandler (2007) Paramutation: from maize to mice. Cell 128: 643-646.

Segregation distortion

In principle, the two parental alleles have equal probabilities of being present in the mature gamete. However, there are several mechanisms that lead to an unequal transmission of parental alleles from parents to offspring. One example is a gene, called a segregation distorter, that "cheats" during meiosis or gametogenesis and thus is present in more than half of the functional gametes.

Sources: wiki and Evolution website.

Gene drives

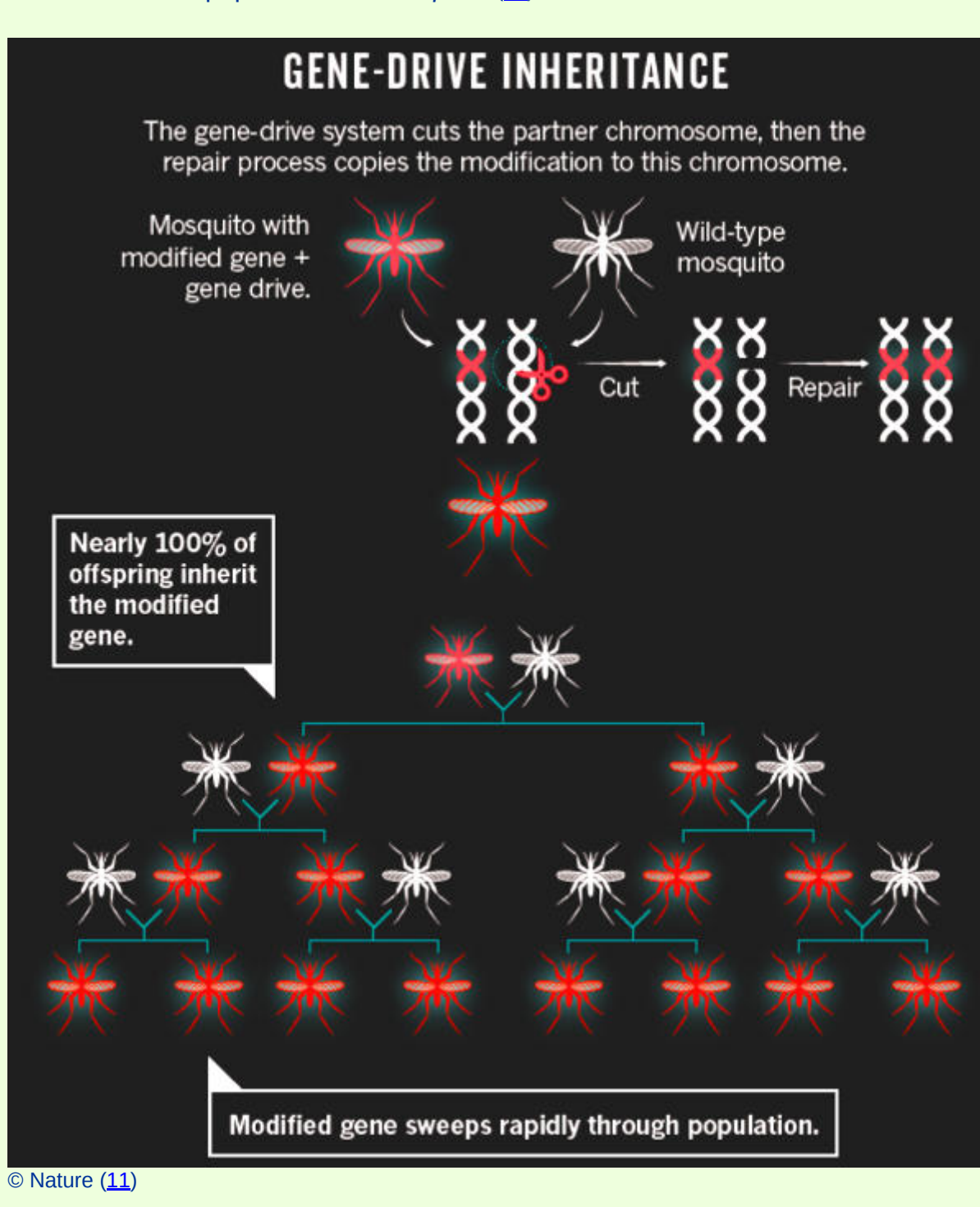


Altered gene + gene drive
1 copy → 2 copies
100% chance of passing it on
© Science

Genes in sexually reproducing organisms normally have, on average, a 50% chance of being inherited, but some genes have a higher chance of being inherited. Gene drives can increase this chance to nearly 100% by cutting homologous chromosomes [by endonucleases]. These genes can increase in relative frequency in a population even if they reduce the odds that each organism will reproduce.

Source: Kenneth A. Oye et al (2014) Regulating gene drives, Science 8 August 2014. See also Gene drive (wiki).

In 2015 researchers engineered gene drive in a lab population of *Drosophila* (10)



© Nature (11)

Mitochondrial inheritance is non-Mendelian

"Because each cell contains many mitochondria, and each mitochondrion has up to 10 copies of its genome, some cells and tissues can acquire more faulty genes than others. This randomness means that apparently healthy women can be carriers of mitochondrial disease, not discovering the problem until they give birth to children who are very ill."

Source: Gretchen Vogel (2004) FDA Considers Trials of 'Three-Parent Embryos', Science 21 Feb 2014.

This implies that mitochondrial inheritance has a non-diploid genetics and so is non-Mendelian.

B chromosomes

In addition to the normal set of chromosomes, eukaryote genomes sometimes also contain chromosomes that do not follow the Mendelian law of inheritance. These chromosomes, called B chromosomes, are believed to consist of selfish genetic elements that have parasitized the genome. B chromosomes are typically supernumerary, derived from ordinary chromosomes.

Source: On the origin and evolution of germline chromosomes in songbirds.

173 pairs of legs

The story of the centipede is one of Dover's most illuminating and intriguing examples. Some centipedes (hundred-legged) have as much as 173 pairs of legs (a head segment, 173 body segments with 1 pair of legs, and two closing segments without legs), the species with the lowest number has 15 pairs of legs. Did the centipede arrive at 173 pairs of legs solely by mutation and natural selection? Are 173 pairs of legs an adaptation to a special kind of environment? Are 15 pairs of legs an adaptation to another environment? Have 16 pairs (the first step towards 173) higher survival value than 15 pairs? Even involving neutral drift requires that one (?) segment at a time spreads through an entire population by accidents of sampling in many generations until 173 pairs are established. Not a very likely possibility. Dover convincingly casts doubt on natural selection and neutral drift as the sole cause. This does not automatically prove his case for Molecular Drive as the correct explanation. However Dover suggests that properties of *Hox* genes could explain the duplication of identical segments with identical legs.



700 copies of a gene

Mendelian inheritance started with the discovery of heritable variations of a single character. For example Mendel's garden peas with round versus wrinkled seeds. Later these characters turned out to be based upon a mutation in one gene. Eye colour is an example of a genetic variant in one gene. Soon after the introduction of genome sequencing it was discovered (to the surprise of geneticists!) that genes were present in many copies. In humans 700 copies of the ribosomal RNA gene were found (1). All copies are functional. The next surprise was that the copies were identical. This is really a non-Mendelian situation. In Mendelian genetics, genes are present in just two copies. Imagine that genes for eye colour or hair colour were present in 700 copies! However this observation and many similar observations can now be found in the textbooks (2) and their existence is not doubted. That the copies are useful for producing large amounts of the gene products is not doubted either. What is controversial is the explanation of the fact that the copies are still identical. One would expect that different mutations accumulated in all those copies. But this is not observed. Dover rejects the standard explanation that natural selection ('purifying selection') is able to keep the 700 copies identical because mutated copies will be compensated for by many intact copies. They are below a threshold. He explains the observation by Molecular Drive: a collection of mechanisms that keeps copies of genes identical. The word Molecular Drive itself is not adopted in the textbooks (3), but the mechanisms behind it are. I still find homogenisation a mysterious mechanism.

Messy Design

The genome is complex, intertwined, ever changing, redundant, contaminated, in short: 'a mess, but it works'. Genomes are ten thousand or a hundred thousand times larger than necessary ... For example humans have alpha-satellite DNA, that consists of several hundred thousand copies spread in tandem arrays all over our 23 pairs of chromosomes. ... Humans carry enough DNA in each cell nucleus to code for 3 million genes. In reality we need only about 70,000 genes. ... Why are all genomes subject to such a bizarre variety of Non-Mendelian mechanisms? ('slippage', 'transposition', 'unequal crossing-over', 'gene conversion', 'homogenisation', 'concerted evolution', 'jumping genes' (4), 'mobile elements'). It is clear that nobody is supervising the evolution of the genome. It is anarchy. It is really time to clean up the mess and enforce rules. Who designed this mess? And if the genome was designed, where is the manual, because maintenance is urgently needed. Who's in charge? Is this mess 'the wisdom of the body' (5). It's really irresponsible to let ignorant DNA molecules build highly complex multicellular conscious beings!

Through Dover I realised that mutations not only produce piecemeal fine tuning of existent proteins and enzymes, but that mutations also produce and modify the bodyplan of organisms. That is a huge conceptual leap. A leap that could not be made as long as Darwinists knew nothing about the genetic control of bodyplans. As if Darwinists secretly believed that the bodyplan, the basic layout of organisms, could not evolve step by step, but must have been created in one big step (by an intelligent designer). Dover shows molecular mechanisms that could be responsible for mutations affecting the bodyplan. The new science of developmental genetics shows how the bodyplan is genetically controlled and how mutations produce modifications of the bodyplan. That is the knowledge Darwinists unknowingly (?) were waiting for.

All genes are interacting with one another. One gene can contribute to many different structures and functions, and any given structure is built by many different genes. I was amazed that Dover did not notice that this is a serious problem for evolution driven by random mutations. The critics of evolution would enthusiastically point out that it becomes very difficult to improve the whole intertwined mess by random mutations. For the impact of a mutation in a gene would not be restricted to the gene product itself but would influence many other gene products as well. The chance that a random mutation would be beneficial in all those gene products is small.

A marriage of 3

One cannot fail to notice after reading *Dear Mr. Darwin*, that Dover is an expert on the molecular structure of the genome (6). However *Dear Mr Darwin* is as much about molecular genetics, as about evolutionary biology as about developmental biology. It is impossible to separate the disciplines here. They are closely intertwined. The *Hox* gene family is a good example: they are genes (genetics) and they build an organism (development), and modifications of *Hox* genes give rise to different organisms (evolution). People have speculated about this synthesis. Dover smoothly integrates evolution, genetics and development. I got the feeling that once we know how 'development' builds a differentiated multicellular organism out of one cell, we will know how 'evolution' build the millions of organisms we observe now. Since development is controlled by genes, modifications of those genes could show us, what kind of organisms result from genetic modifications (6). Geno-Evo-Devo: that is the future of biology. An exciting future!

DNA fingerprinting

DNA fingerprinting is now one of forensic science's most reliable and potent weapons. I was amazed to find out that the fingerprinting technique is based on variation in the number of copies of a 20 base DNA sequence. The repeat number is so variable that everybody has a unique genetic fingerprint. This was discovered by Alec Jeffreys (2), just as Dover from the University of Leicester. The mechanisms responsible for this variability are unequal crossing-over and slippage. Slippage is the most frequently occurring mechanism of gain and loss of DNA in genomes. It is one of the mechanisms Dover included in 'Molecular Drive'. Thanks to Molecular Drive DNA-fingerprinting is possible.

Dear Mr Dover

Dover uses imaginary letters to Darwin, and repliers(?) from Darwin, to convince Darwin of the existence of a third force in evolution. In this way Dover created a lively discussion. At the same time there is a perfect opportunity to teach Darwin the relevant parts of genetics. Although Darwin knew everything of evolution, he knew nothing of Mendelian genetics and molecular genetics. However Dover doesn't really seize the opportunity to explain basic concepts in genetics. The task to explain progress in genetics since Darwin's time is a huge task indeed. Genetics and history-of-genetics textbooks has been written to do just that. Darwin and the reader need to do a lot of homework (2) to catch up with Dover's exposition. So this book is for the advanced (professional?) reader or determined lay reader. However Dover has important things to tell. The book is not a mess like the genome, but it is certainly complex with a lot of intertwined parts. I read it twice from cover to cover. I found subjects like gene conversion and the population genetics aspects of gene conversion insufficiently explained. I wonder if field or lab data exist about the spread of gene conversions (in humans)? Could molecular drive spread mutations faster than traditional population genetics allows for? *Dear Mr Darwin* contains an indispensable glossary. And there are black and white drawings (because the book consists of a correspondence?). The science of genetics includes a lot of technicalities and abstract mechanisms. Color illustrations could help the millions of organisms we observe now. Since development is controlled by genes, modifications of those genes could show us, what kind of organisms result from genetic modifications (6). Geno-Evo-Devo: that is the future of biology. An exciting future!

Dear Mr Darwin: Letters on the evolution of life and human nature by Gabriel Dover University of California Press 2000 hardback 268 pages. ISBN: 0-520-22790-5	
Contents:	
How to read this book	x
1. The twin peaks	1
2. The rise and fall of the mobile P gene	24
3. When is an adaption not an adaption?	38
4. The ignorant gene	49
5. Is Dawkins aware of the error of his ways?	67
6. Genetic turnover: of course, of course	77
7. Molecular Drive for advanced players	92
8. Molecular Coevolution	101
9. The mystery of meiosis	118
10. Biological Barriers	130
11. Sex - A new perspective	137
12. HOX: HOX: HOX!	148
13. Born to adapt	197
14. The unknowability of DNA	222
15. The evolution of individuality	232
Glossary, Further Reading, Index	244

Notes:

- The genome is some kind of database of sequences. From the viewpoint of information storage in relational databases it is a crazy idea to store hundreds of copies of the same item. Only unique items are stored and duplications are prevented or eliminated, because duplicate items cause a maintenance problem.
- Dan Grauer and Wen-Hsiung Li(2000) *Fundamentals of Molecular Evolution* have an extensive discussion of "concerted evolution" on p304-322, which is a good preparation for reading Dover.
- According to Kevin Davies (*Cracking the Genome*), Francis Collins believes the human genome sequence is "the language of God" (see: From the Publisher at Barnesandnoble.com).
- see for example the superbly illustrated *Exploring the Biomedical Revolution* of the Howard Hughes Medical Institute or the irresistible *Developmental Biology*, Sixth Edition published by Sinauer Associates.
- The concept 'molecular drive' is not adopted by Grauer & Li (2000). However it is discussed by Jablonka and Lamb (1995) *Epigenetic Inheritance and Evolution*. Wallace Arthur (2000) *The Origin of Animal Body Plans* has 5 references to Dover, but apparently does not see a big role for molecular drive in generating body plans.
- Walter Gehring (1998) *Master Control Genes in Development and Evolution. The Homeobox Story*. (primarily about developmental genetics, secondary but inevitable about evolution).
- Matt Ridley (2000) tells the DNA fingerprinting story in a more popular way in: *Genome*, p131-134.
- Gabriel Dover started publishing in 1990 about "Modes of genome evolution" (with W. Ford Doolittle), *Nature* Vol 288, p.646. 18/25 Dec 1990
- Gretchen Vogel (2011) *Do Jumping Genes Spawn Diversity?* *Science* 15 April 2011: "The data are clear. Transposable elements move in developing brain cells. But the question remains: Does the brain tolerate them or take advantage of them? Formally known as transposable elements, they are small bits of genetic material that can move around the genome. They have generally been seen as troublemakers, when they jump, they can land in places that cause mutations or otherwise skew the expression of important genes." "Nearly 45% of the human genome is made up of jumping genes."
- "Using an extremely sensitive PCR technique, they estimated that brain cells had about 80 additional copies of L1 compared with other tissues."
- "We were really surprised," Kazarian says, to find more L1 jumps in early embryos than even in sperm or egg cells."
- Valentino M. Gantz, Ethan Bier (2015) "The mutagenic chain reaction: A method for converting heterozygosity to homozygous mutations". *Science* 24 April 2015
- Heidi Ledford (2015) "CRISPR, the disruptor", *Nature*, 03 June 2015

Further Reading:

- Home page of Gabriel Dover. I was unable to find an email address of Gabriel Dover.
- Enrico Coen (1999) *The Art of Genes. How organisms make themselves* (my short review at amazon). This book is highly praised by Dover. It has high educational value. The emphasis is, contrary to Dover's book, on development and genetics, not on evolution. See also a review in *Nature* 398, 302-303 (25 March 1999) by John Maynard Smith.
- E.J. Steeler (2000) *The Evidence for Lamarck*. QUADRANT March 2000 No. 364 Vol XLIV Number 3 pages 47-56. This is an extremely useful and masterfully written summary of Steeler's arguments for the general reader. 'Conventional neo-Darwinian population genetics will not handle this one, nor will Gabriel Dover's molecular drive concepts.'
- John Alcock reviews *Dear Mr Darwin: What Would Darwin Think?* in *American Scientist* January-February, 2001. (John Alcock is Professor of Biology at Arizona State University). Regrettably Professor Alcock did not evaluate the role of Molecular Drive in evolution, which is the main theme in Dover's book. [link update by Chris Eilers]
- Letter from Jack Haas from the Feedback page.
- John Waller shows why Dover is wrong when stating: "Mendel's flash of inspiration was to deduce from this that the cells making up each individual pea plant contained two copies of each gene..." (p.11). See the review of *Fabulous Science. Fact and Fiction in the history of scientific discovery* (John Waller).
- Gabby Dover (1999) "Copying the evolutionary logg". *Nature* 399, 217-218, 20 May 1999. This is a very dismissive and unfair review of John Maynard Smith & Eors Szathmari (1999) "The origins of life" (see: review on this site).
- Gabriel Dover (2000) "Problems and paradigms. How genomic and developmental dynamics affect evolutionary processes", *BioEssays* Volume 22, Issue 12, Pages 1153 - 1159.
- Gabriel Dover (2001) "Anti-Darwinians" in *Alas poor Darwin*, Vintage.
- J.R. Minkel (2005) "RNA to the rescue. Novel inheritance patterns violate Mendel's laws". *Scientific American*, June 2005, 8-10. [10 Jul 2005]
- Elizabeth Pennisi (2007) "Jumping Genes Hop Into the Evolutionary Limelight", *Science*, 17 August 2007.
- Andrew Slater (2000): "Some authors continue to purvey enigmatic half-truths on the subject. Gabriel Dover, on page 12 of *Dear Mr Darwin* (2000), refers to the 'mystery' of CD supplying Mendel's name for inclusion in the hybridism entry for *Encyclopaedia Britannica*. From the foregoing, it is apparent that CD did so merely by lending a book to Romanes. He did not make a personal recommendation of any individual, and certainly not of Mendel." Source: DARWIN AND MENDEL.
- Guangbo Chen, Boris Rubinsteyn, Rong Li (2012) *Whole chromosome aneuploidy. Bio mutations drive adaptation by phenotypic leap*, *BioEssays* Volume 34, Issue 10, pages 893-900, October 2012.
- Bruce R. Conklin (2019) On the road to a gene drive in mammals, *Nature*, NEWS AND VIEWS 23 January 2019: "A method for making a version of a gene more likely to be inherited than normal, generating what is called a gene drive, might be used to control insect populations. It has now been reported to work in mammals, too." 23 January 2019

Feedback	home: wasdavidwong.com	wasdavidwong.com/korthof51.htm
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