

A lesson from sex:

Abundant variation can be worth its high price.

David G King, Southern Illinois University Carbondale

Blind, undirected mutational variation is mostly deleterious, but without it evolution would grind to a halt. Blind, undirected variation from sexual reproduction is vital for maintaining population fitness, though it comes at a very steep price. In between these two contrasting ideas -- that replication errors are bad but inevitable while sex is good though expensive -- lies a seldom appreciated reality: Quite generally, sources of suitably constrained variation can be favored by natural selection in spite of seemingly exorbitant cost.

To be sure, sex generates an especially conservative style of variation, guaranteeing a diversity of genotypes simply by rearranging pre-existing alleles. But sex also imposes a huge burden: In addition to the many hazards of mating, sex entails a 50% reduction in fitness relative to the efficiency of asexual reproduction, and meiotic recombination separates favorable alleles just as readily as it brings them together. Quantifying benefits sufficient to balance these high costs remains an elusive goal. Nevertheless, the prevalence of sexual reproduction among eukaryotes proves that at least one source of blind, undirected variation can be worth an enormous price. This understanding of sexual reproduction should prompt us to consider that some mutational sources of variation might similarly confer benefit sufficient to outweigh cost, even if their benefit cannot yet be clearly appraised.

The idea that selection could favor an elevated frequency for any type of mutational variation has long been dismissed: "[N]atural selection of mutation rates has only one possible direction, that of reducing the frequency of mutation rates to zero" (GC Williams, 1966). But such sweeping denial is based on a simplistic argument whose underlying assumptions do not apply to several highly-constrained mutational mechanisms, including expansion and contraction of simple sequence repeats, transposition of mobile elements, gene duplication, horizontal gene transfer, localized hypermutability, and phase switching. Such mechanisms bias the styles and sites for resulting mutations, thereby offering an opportunity to shift the balance between harm and benefit. Abundant evidence that these mechanisms have contributed to adaptive evolution should suggest that their associated constraints might constitute "protocols" for generating advantageous variation. If the obvious harm from deleterious mutation does not exceed the stunningly high cost of sex, then natural selection might also deem the benefits from any such source of variation (including an emergent potential for innovative exploration) as being well worth the price. Website: www.siu.edu/anatomy/KingCoS/index.htm

Further reading

Arber W (2005) Gene products with evolutionary functions. *Proteomics* 5: 2280-2284.

Barry JD (2006) Implicit information in eukaryotic pathogens as the basis of antigenic variation. In: LH Caporale, ed., *The Implicit Genome*, pp. 91-106. Oxford University Press, Oxford.

Bayliss CD, Moxon ER (2006) Repeats and variation in pathogen selection. In: LH Caporale, ed., *The Implicit Genome*, pp. 54-76. Oxford University Press, Oxford.

Bell G (1982) *The masterpiece of nature: The evolution and genetics of sexuality*. Croom Helm Ltd, London. (*"The paradox of sex is the queen of problems in evolutionary biology."*)

Bridges CB (1919) Specific modifiers of eosin eye color in *Drosophila melanogaster*. *J Exp Zool* 28(3): 37-384. (*Defines "mutation" and establishes the expectation that most mutations are deleterious.*)

Caporale LH (1999) Chance favors the prepared genome. In: LH Caporale, ed., *Molecular Strategies in Biological Evolution*, *Ann N Y Acad Sci* 870: 1-21.

Caporale LH (2000) Mutation is modulated: Implications for evolution. *BioEssays* 22: 388-395.

Caporale LH. (2003a) Natural selection and the emergence of a mutation phenotype: An update of the evolutionary synthesis considering mechanisms that affect genomic variation. *Ann Rev Microbiol* 57: 465-485.

Caporale LH (2003b) Foresight in genome evolution. *Amer Sci* 91: 234-241.

Caporale LH (2006) An overview of the implicit genome. In LH Caporale, ed., *The Implicit Genome*, pp. 3-22. Oxford University Press, Oxford.

Caporale LH, Doyle J (2013) In Darwinian evolution, feedback from natural selection leads to biased mutations. *Ann N Y Acad Sci* 1305: 18-28.

Carja O, Liberman U, Feldman MW (2014) The evolution of phenotypic switching in subdivided populations. *Genetics* 196: 1185-1197.

Csete M, Doyle J (2002) Reverse engineering of biological complexity. *Science* 295: 1664-1669. (*Introduces the "protocol" metaphor.*)

Darwin, CR (1859a) Letter to TH Huxley, Nov. 25, 1859. *Darwin Correspondence Project*. <http://www.darwinproject.ac.uk/entry-2553>.

Darwin, CR (1859b) *On the Origin of Species by Means of Natural Selection*. John Murray, London.

Doyle J, Csete M (2007) Rules of engagement. *Nature* 446: 860.

Doyle J, Csete M, Caporale LH (2006) An engineering perspective: The implicit protocols. In: LH Caporale, ed., *The Implicit Genome*, pp. 294-298. Oxford University Press, Oxford.

Earl DJ, Deem MW (2004) Evolvability is a selectable trait. *Proc Natl Acad Sci USA* 101: 11531-11536. (*"Life has evolved to evolve."*)

Gemayel R, Cho J, Boeynaems S, Verstrepen KJ (2012) Beyond junk - Variable tandem repeats as facilitators of rapid evolution of regulatory and coding sequences. *Genes* 3: 461-480.

Gemayel R, Vences MD, Legendre M, Verstrepen KJ (2010) Variable Tandem Repeats Accelerate Evolution of Coding and Regulatory Sequences. *Annu Rev Genet* 44: 445-477.

Goldschmidt R (1940) *The Material Basis of Evolution*. Yale University Press, New Haven. (*Before DNA, a premature inquiry into the nature of mutation protocols.*)

Jurka J, Kapitonov VV, Kohany O, Jurka MV (2007) Repetitive sequences in complex genomes: structure and evolution. *Annu Rev Genomics Hum Genet* 8: 241-259.

Jurka J (2007) Conserved eukaryotic transposable elements and the evolution of gene regulation. *Cell Mol Life Sci* 65: 201-204.

Kashi Y, King DG (2006a) Simple sequence repeats as advantageous mutators in evolution. *Trends Genet* 22: 253-259.

Kashi Y, King DG (2006b) Has simple sequence repeat mutability been selected to facilitate evolution? *Isr J Ecol Evol* 52: 331-342. (*Includes an explanation of indirect selection.*)

King DG, Kashi Y (2007a) Mutability and evolvability: Indirect selection for mutability. *Heredity* 99: 123-124.

King DG, Kashi Y (2007b) Mutation rate variation in eukaryotes: Evolutionary implications of site-specific mechanisms. *Nature Rev Genet* 8 (doi: 10.1038/nrg2158-c1).

King DG (2012) Indirect Selection of Implicit Mutation Protocols. *Ann N Y Acad Sci* 1267: 45-52. (*Most appropriate citation for this poster presentation.*)

King DG, Soller M (1999) Variation and fidelity: The evolution of simple sequence repeats as functional elements in adjustable genes. In: S.P. Wasser, ed., *Evolutionary Theory and Processes: Modern Perspectives*, pp. 65-82. Kluwer Academic Publishers, Dordrecht. (*Includes an explanation of indirect selection.*)

King DG, Trifonov EN, Kashi Y (2006) Tuning knobs in the genome: Evolution of simple sequence repeats by indirect selection. In: LH Caporale, ed., *The Implicit Genome*, pp. 77-90. Oxford University Press, Oxford.

Kirschner M, Gerhart J (1998) Evolvability. *Proc Natl Acad Sci USA* 95: 8420-8427.

Martincorena I, Luscombe NM (2013) Non-random mutation: The evolution of targeted hypermutation and hypomutation. *BioEssays* 35: 123-130.

Mihola O et al. (2009) A mouse speciation gene encodes a meiotic histone H3 methyltransferase. *Science* 328: 373-375. (*A protocol for speciation?*)

Oliver KR, Green WK (2009) Transposable elements: Powerful facilitators of evolution. *BioEssays* 31: 703-714.

Otto SP, Lenormand T (2002) Resolving the paradox of sex and recombination. *Nature Rev Genet* 3:252-261.

Otto SP (2008) Sexual reproduction and the evolution of sex. *Nature Education* 1:182.

Shapiro JA (1983) Variation as a genetic engineering process. In D.S. Bendall, ed. *Evolution from Molecules to Men*, pp. 253-270. Cambridge University Press, Cambridge.

Shapiro JA (1997) Genome organization, natural genetic engineering and adaptive mutation. *Trends Genet* 13: 98-104.

Sniegowski PD et al. (2000) The evolution of mutation rates: Separating causes from consequences. *BioEssays* 22: 1057-1066. (*Contrary viewpoint, reiterates the classical argument that natural selection favors minimal mutation rates.*)

Sniegowski PD, Murphy HA (2006) Evolvability. *Current Biology* 16: R831-R834. (*Contrary viewpoint, argues that evolvability is not an adaptation.*)

Sturtevant AH (1937) Essays on evolution. I. On the effects of selection on mutation rate. *Q Rev Biol* 12: 464-467. (*"Mutation are accidents."*)

Thaler D (1994) The evolution of genetic intelligence. *Science* 264: 224-225.

Verstrepen KJ et al. (2005) Intragenic tandem repeats generate functional variability. *Nature Genet* 37: 986-990.

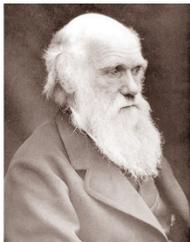
Vences MD et al. (2009) Unstable tandem repeats in promoters confer transcriptional evolvability. *Science* 324: 1213-1216.

Volff JN (2006) Turning junk into gold: Domestication of transposable elements and the creation of new genes in eukaryotes. *BioEssays* 28: 913-922.

Weissmann A (1889) *Essays Upon Heredity*. Oxford, Clarendon Press.

Williams GC (1966) *Adaptation and Natural Selection*. Princeton Univ. Press, Princeton. (*A classic text, with a strong argument that natural selection must always prefer minimal mutation rates.*)

Williams GC (1975) *Sex and Evolution*. Princeton Univ. Press, Princeton.



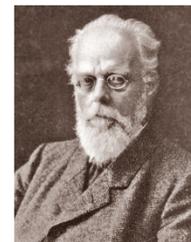
Some authors believe it to be as much the function of the reproductive system to produce individual differences ... as to make the child like its parents.

Charles Darwin 1859

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David G. King Depts. of Anatomy and Zoology, Southern Illinois University Carbondale

What the devil determines each particular variation? What makes a tuft of feathers come on a cock's head, or moss on a moss-rose? Charles Darwin (in a letter to TH Huxley, 1859)



The object [of sexual reproduction] is to create those individual differences which form the material out of which natural selection produces new species.

August Weismann 1889

Introduction

This poster invites readers to consider an analogy between sex and mutation.

Sex and mutation share fundamental similarities, but their differences are usually emphasized:

Sex is understood as an evolved function. Despite the high cost of reproducing sexually, selection favors the allele-shuffling of meiotic recombination. Some undirected but highly constrained variation is evidently more advantageous than exact genome duplication.

Mutations are understood as accidental errors. Selection for minimal mutation rates is believed to be limited only by the cost of replication fidelity.

Nevertheless, this distinction may not be as great as generally conceived. Fundamentally, both sex and mutation are sources of genetic variation.

In essence, sexual recombination is a source of mutation, creating novel DNA sequences.

Certain mutational mechanisms can, like sex, yield highly constrained styles of variation which can be employed for adaptive advantage.

Both sexual reproduction and several styles of mutation should be conceptualized together, as "protocols" that can facilitate stochastic but adaptively useful variation.

Background: History

Darwin (1859) famously anticipated future inquiry into "the causes and laws of variation."

Sex as a source of variation

That sexual reproduction functions as a source of variation seemed evident by the late 1800s.

However, by the mid-1900s sex had been designated as "the queen of problems in evolutionary biology" (G Bell 1982).

Because organisms reproducing sexually must produce twice as many offspring to compete effectively against asexuals, identifying benefit sufficient to overcome such a huge selective disadvantage had become a major theoretical challenge.

Nevertheless, recent theoretical models have finally been validating the old view:

"August Weismann [1889] might have been right all along in arguing that sex evolved to generate variation" (SP Otto, 2008).

In other words, sex really is "a parental adaptation to the likelihood of the offspring having to face changed or uncertain circumstances" (GC Williams 1975).

Mutation as a source of variation

Unfortunately, in contrast to such recent developments concerning sex, theoretical modeling of mutability has lagged far behind growing knowledge of mutational mechanisms.

The prevailing explanation for the existence of mutation remains largely as developed in the mid 20th century, that "mutations are accidents, and accidents will happen" (Sturtevant 1937).

"[N]atural selection of mutation rates has only one possible direction, that of reducing the frequency of mutation to zero. ... So evolution takes place, not so much because of natural selection, but to a large degree in spite of it" (GC Williams 1966).

But this view properly applies only to a mutator allele which reduces the genome-wide fidelity of DNA replication without remaining linked to any resulting mutations, and only when the "vast majority" of mutations are deleterious.

In spite of such limited applicability, this view that mutations are accidents is commonly wielded against the idea that any style of mutation could be advantageous.

For many well-known sources of mutation, circumstances for positive selection may closely parallel those for selective maintenance of sex.

Mutations are often deleterious, but -- as with sex -- immediate advantages for certain styles of constrained mutability could consistently outweigh the cost.

Sexual recombination: Cost vs. benefit

Meiotic recombination assures unique allele combinations in each individual offspring.

Cost

Sexual reproduction imposes a profound 50% cost relative to asexual reproduction, in the efficiency of transmitting genes to the next generation.

Furthermore, meiotic recombination can separate favorable combinations of alleles. And the act of mating is time-consuming, effortful, and dangerous.

Specific benefits sufficient to overcome such huge costs are not readily apparent.

"The widespread occurrence of sex, despite its seemingly overwhelming costs, is known as the paradox of sex" (SP Otto 2002).

Benefit

Recent models that avoid unrealistic assumptions have suggested that (SP Otto 2008):

- Sex evolves when selection changes over time.
- Sex evolves when selection changes over space.
- Sex evolves when organisms are less-well adapted to their environment.
- Sex evolves when populations are finite.

Apparently, variation from sex can be beneficial under conditions experienced by most natural populations, in spite of "seemingly overwhelming costs."

Mutation protocols: Cost vs. benefit

Mutation protocols are conceived as bet-hedging strategies which promote advantageously constrained styles of DNA variation.

Cost

Deleterious mutations are an inevitable result of all known mutational processes.

Classically, the "vast majority" of mutations are deleterious, no matter what the cause.

Some deleterious mutations can be severely deleterious.

Yet for some mutational mechanisms, both the proportion of severely deleterious mutations and the overall cost remain quite low, even with extremely high mutation rates.

No mutational process carries an "overwhelming cost" as high as that paid for sexual reproduction.

Benefit

Several mutational mechanisms could represent beneficial bet-hedging strategies. Thus, just as for sex:

- Mutation protocols might evolve when selection changes over time.
- Mutation protocols might evolve when selection changes over space.
- Mutation protocols might evolve when organisms are less-well adapted to their environment.
- Mutation protocols might evolve when populations are finite.

One or another mutation protocol might be favored under conditions commonly encountered in nature.

A sampling of mutation protocols

Bet-hedging strategies for DNA variation range from sophisticated to simple, with corresponding differences in style of mutation, mutation rate, and risk of deleterious effect. Several of these protocols are well-established as the means for rapid and effective microbial adaptation. There is no reason (apart from over-confidence in simplistic theory) to doubt that complex eukaryotes also exploit such protocols.

- Meiotic recombination** is supported by an astonishing array of anatomical, physiological, and behavioral adaptations. But this protocol is typically excluded, by definition, from the concept of "mutation."
- Phase switching** shuffles genes in and out of active sites by programmed gene arrangement, without necessity for sex.
- Horizontal gene transfer** offers access to potentially-advantageous alleles previously evolved by other members of the local community.
- Tuning-knob sites** based on tandem repeats allow reversible, incremental adjustment of most gene functions, including site-specific adjustment of mutation rate.
- On/off switching** (also based on tandem repeats) allows stochastically reversible variation in the expression of "contingency genes."
- Transposable elements** implement copy-and-paste of functional modules. Although seemingly selfish, appropriately domesticated TEs (JN Volt 2006) provide opportunities for adaptive innovation and diversification, especially in times of stress.
- Targeted hypermutation** concentrates single-nucleotide mutation at mutation hotspots where variation has proven especially advantageous in the past.
- Epigenetic modification** offers heritable variation without altering DNA sequence and may provide a substrate for subsequent, site-specific mutation.
- Whole-genome duplication** creates a variety of opportunities for diversifying variation.
- Conventional mutation** (e.g., non-site-specific alterations of single nucleotides) may be the only style of mutation that is adequately modeled by conventional theory, such that mutation rate is minimized rather than optimized by selection.

This is but a partial list of special modes of mutation that are available for exploitation as protocols for generating variation. As with sexual reproduction, each of these protocols may be favored, suppressed, or regulated, depending on a population's circumstances.

Background: Indirect selection

Indirect selection for mutation protocols (DG King 2012) occurs when favorable variants arise within constraints that are themselves heritable and linked to those variants.

An example: The potential for indirect selection is most clearly illustrated by site-specific elevation of mutation rate, as represented by tandem repeats. When favorable variants arise, they retain the site-specific mutation rate by which they arose. Selection for the favorable variant then also indirectly but inevitably favors the locally elevated mutation rate for this particular style of mutation, thus facilitating future variation under similar constraints.

Indirect selection should be expected to shape and maintain any mechanism of mutation whose utility offers even a fraction of the adaptive value provided by sexual reproduction.

Although natural selection cannot directly favor genomic patterns which facilitate propitious styles of variation, indirect selection can nevertheless shape mutation protocols just as effectively as natural selection can shape phenotypic adaptation.

What next?

Several concepts merit further exploration.

Genomes have evolved to evolve. They exploit a wide range of protocols to manage the potential advantages as well as the risks of genetic variation.

Sexual reproduction with meiotic recombination is perhaps the most sophisticated (and expensive) of these protocols.

The surprising prevalence of several mutational mechanisms suggests that they too should be understood as implicit protocols for stochastic production of variation rather than as flaws in replication fidelity. Resulting changes in DNA sequence are better viewed not as "mistakes" or "accidents" but as products of these protocols.

If variation from sexual recombination can offer generation-by-generation advantage sufficient to outweigh its "seemingly overwhelming" cost, then perhaps other mechanisms for producing variation can also be maintained by positive selection.

As long as the burden of deleterious mutation does not exceed the 50% cost of sex, positive selection for a protocol should be considered as plausible.

Mutation protocols can thereby be integrated, together with sexual recombination, into patterns of "genetic intelligence" (DS Thaler 1994).

Mutation protocols complement physiological and epigenetic mechanisms for responding to environmental variation, while offering emergent opportunities for evolutionary innovation.

Mutation protocols form the basis for creative bet-hedging in a complex and inconstant world. The selective value of mutation protocols, although difficult to measure in nature, should be addressed through modeling of indirect selection (cf. O Carja et al. 2014).

Understanding the genetic basis for evolvability, especially for evolutionary innovation in complex adaptive behavior, may well depend on appreciating the role of implicit mutation protocols.

Abstract and references

Please see the handout for references as well as a copy of poster text and abstract. If handouts are not available, please contact the author at dgking@siu.edu.

