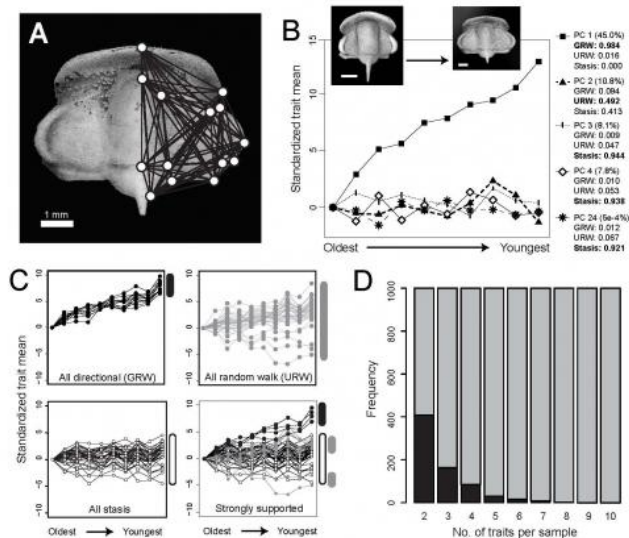


# The more things change: Trait variance provides evidence of pervasive mosaic evolution

14 December 2012, by Stuart Mason Dambrot



Results from simulated trilobite cranium shape data. (A) Landmarks representing overall shape of cranium (required only from half because of bilateral symmetry) and all possible length measurements between landmarks. (B) Time series of selected PC axes scores. (Upper Left) Image shows example of typical morphology from oldest sample; (Upper Right) image shows typical morphology from stratigraphically youngest sample (scale bars = 1 mm). Change in morphology is dominated by an expansion and rotation of palpebral lobes relative to rest of the cranium. (C) Time series of individual length: length measurements, divided up into four panels showing those that were best characterized by directional change (Upper Left) (black), unbiased random walk (Upper Right) (gray), stasis (Lower Left) (white), and strongly supported trends of all types (Lower Right). Bars on right show range of means for each mode of evolution at the end of the sequence. (D) Stacked histograms showing distribution of samples of randomly selected length: length measurements where all traits in the sample show the same evolutionary mode (black) and where evolutionary mode varies across traits (gray). Number of samples taken at each sample size = 1,000. Copyright © PNAS, doi:10.1073/pnas.1209901109

(Phys.org)—Despite evidence that phenotypic change does not always occur uniformly across all species members and lineages, single size or shape traits are often used to represent species-level change in its entirety. Recently, however, scientists in the Department of Geology at the Field Museum of Natural History in Chicago and at the Museum für Naturkunde in Berlin asked a deceptively simple question – *Are (or when are) single traits adequate representations of species-level change?* – and answered it by asking another: *How often do single traits show conflicting patterns in the same sequence?* To answer the second, they examined trait variation frequency in fossils, finding that within most lineages, evolutionary mode (the pattern of evolution, as opposed to its rate) does indeed vary across traits – and the probability of these patterns conflicting within a given lineage patterns increases in proportion to the number of traits analyzed. Moreover, the evolution of single traits may vary despite a particular mode being dominant. The researchers suggest that their findings provide evidence that widespread mosaic evolution – the tendency for different parts within species to evolve in different ways or at different rates – has occurred throughout our planet's history.

[Postdoctoral researcher Melanie J. Hopkins](#) (Museum für Naturkunde) and Associate Curator [Scott Lidgard](#) (Field Museum of Natural History) encountered a range of challenges when conducting their investigation. "The main challenge in testing the validity of this [assumption](#) that single [phenotypic traits](#) are adequate representations of species-level change is that, due to time and computational constraints, researchers are limited in just how much they can measure when describing an organism," Hopkins tells *Phys.org*. "Therefore, it's essentially impossible to know all aspects of change within a species through time –

making it very difficult to test the assumption that a single trait, or even several traits, adequately represents change across the species."

When it came to tallying the frequency with which traits vary in evolutionary mode within fossil species lineages, Hopkins adds, the main challenge here was twofold: tabulating the data in a form that could be consistently analyzed; and determining if the dataset or methods used biased the results in any way.

The researchers addressed these challenges through a series of insights, innovations and techniques. "In reading previous work," Hopkins notes, "we realized that traits were often analyzed separately – even in the same study – and that the traits did not always show the same mode of evolution. This implied that our understanding of the evolution of species might depend critically on what traits we were looking at." Once having this insight, the scientists wanted to know how widespread this conflict was across different groups of [organisms](#) and through time. They then decided to undertake a systematic review of all of the traits that had been measured in the fossil record to date.

"Past studies on within-lineage trends in the fossil record were conducted at different levels of spatial and temporal resolution and using different methods of analysis," Hopkins continues. "As a result, it has been difficult to synthesize findings without simply relying on the particular interpretation on any given study. Instead, we analyzed the original data from as many studies as we could find using a consistent analytical protocol – in this case, the model-selection based method developed in 2006 and 2007 by Gene Hunt at the Smithsonian." The model-selection technique is very powerful because it gave the scientists a way to compare numerous studies in a consistent and rigorous way.

One of the study's findings was that within the majority of [lineages](#), evolutionary mode varies across traits and the likelihood of conflicting within-lineage patterns increases with the number of traits analyzed. "Some traits show directional change – for example, an increase in tooth size over time – while others traits change randomly, show little

change or stasis, Hopkins explains. "However, traits do not always show the *same* patterns of change – they frequently do not change in a coordinated fashion, even within the same lineage. Therefore, the more traits researchers measure from a single lineage, the more likely they are to see conflict in patterns of change amongst traits."

Another important result was that single traits may show variation in evolutionary mode even in situations where the overall morphological evolution of the lineage is dominated by one type of mode. "Sometimes evolutionary change within a species is dominated by a particular mode of evolution – for example, overall, a species may stay relatively the same in size and shape through time," says Hopkins. "However, there are often situations where a few traits change even though the rest do not."

Perhaps the study's key finding is that its results validate the idea that morphological patterns of mosaic evolution are pervasive across groups of organisms throughout Earth's history. "Our results show that the frequency for different parts within species to evolve in different ways is high. Because this dataset included representatives from all different types of organisms that were sampled throughout the last 500 million years of the [fossil](#) record, including mammals, fish, mollusks, and single-celled marine plankton, our results imply that mosaic evolution is ubiquitous."

Moving forward, Hopkins concludes, "We're currently looking in more detail at data collection methods and expanding our inquiry to include additional and more complicated models of evolution."

**More information:** Evolutionary mode routinely varies among morphological traits within fossil species lineages, *PNAS* Published online before print November 26, 2012, [doi:10.1073/pnas.1209901109](https://doi.org/10.1073/pnas.1209901109)

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