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## Oak symbolism in the light of genomics

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### Summary

Throughout the Northern Hemisphere, human societies, political systems and religions have appropriated oaks in symbolic representations. In this review, we explore the possible associations between recent genetic and genomic findings and the symbolic representations of oaks. We first consider the ways in which evolutionary history during the Holocene has tightened links between humans and oaks in Europe, and how this may have led to symbolic representations. We then show how recent findings concerning the structure and evolution of the oak genome have provided additional knowledge about symbolic representations, such as longevity, cohesiveness and robustness.

### I Introduction

Throughout the Northern Hemisphere, human societies, political systems and religions have appropriated oaks in symbolic representations. Oaks have been present in traditional rites and celebrations for many centuries, as far back in history as the early Proto-Indo-European cultures (Chassé, 2016). Oaks have been associated with longevity, strength, fertility and robustness. Recent findings in genetics and genomics may shed new light on these symbolic representations of oaks. Intricate relationships between oaks and human populations developed following the migration of humans out of Africa, due to reciprocal benefits. Did the joint evolutionary trajectories of humans and oaks during the Holocene contribute to their “shared” history, reinforcing ties between trees and people? This review is not designed to be an exhaustive consideration of the many types of symbolism attached to oaks by humans. We address only a few symbolic elements (longevity, cohesiveness, robustness) for which the association with a genetic perspective is most marked, particularly in recent recorded European history (from Greek and Roman to Celtic societies). Our approach should be seen as a first step towards the assembly of knowledge from different fields (history, ecology, genomics and social sciences) with a view to understanding symbolic representations of plants in human societies.

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## II From early ties between humans and oaks to symbolic representations

Symbolism may have been attached to trees in early human societies on the basis of the very early ties linking people and trees and the ways in which these ties developed historically. Oaks, more than any other tree species, paved the way for the colonization of the Middle East and Europe by *Homo sapiens* some 45-50 thousand years ago (Bar-Yosef & Befer-Cohen, 2013). Oaks were an invaluable food resource for humans during the early settlement of Eurasia, as suggested by the remains of *Quercus ithaburensis* and *Q. calliprinos* acorns in caves known to have been occupied by humans (Lev *et al.*, 2005). There is increasing evidence to suggest that acorns made an essential contribution to the staple diet of the first modern humans, based on ethnographic, archeological and historical clues. Despite underestimations of acorn consumption due to the charring of remains and the decay of fossil remains (taphonomical issues), recent archeological investigations have highlighted the widespread use of acorns as a food source (Lev *et al.*, 2005; Humphrey *et al.*, 2014, Antolin and Jacomet, 2015; Morales, 2018), by various hominids, including Neanderthals (Villa and Roebroeks, 2014). This archeological element is supported by historical evidence from the writings of the earliest geographers, such as Pausanias and Strabon, philosophers, such as Platon, poets, such as Ovid, and naturalists, such as Pliny the Elder and Theophrastus, describing the regular consumption of acorns up to the Greek and Roman ages (reviewed by Chassé, 2016). Before the development of agriculture in Mesopotamia, human hunters and gatherers fed themselves with various seeds, including acorns from the oak forests of the Zagros Mountains or the oak-pistachio uplands of the Mesopotamian valley (Murphy, 2007; Scott, 2017). Ties between humans and oaks were reinforced as humans migrated to Europe and experienced the last glacial/interglacial transition about 40,000 years ago, and then during later migrations of human populations northwards. Kremer (2015) highlighted intriguingly similarities in the colonization dynamics of modern humans and oaks, in terms of velocity (500 m/year Brewer, 2002 for oak and Henn *et al.*, 2012 for humans), processes (serial founder effects, Le Corre *et al.*, 1997 for oaks and Deshpande *et al.*, 2012 for humans) and colonization routes. These similarities are too wide-ranging to be mere coincidence, suggesting that humans may have transported oaks as food reserves while migrating northwards as the climate warmed, thus contributing to rapid oak colonization (Kremer, 2015).

## III The symbolic load of oak in the humanities and science

The use of oak for food was paralleled by its use for other purposes. Acorns were used for food, but other parts of the tree were used for medicine, fuel, shelter and art, as oaks became the “tree of life” (Anderson, 2007) or the “frame of civilization” (Logan, 2006). For example, almost all prehistoric pile dwellings built along the shores of lakes during the Bronze Age were made of oak logs (Menotti, 2004). Oaks clearly facilitated the establishment of human populations in Europe, and humans returned the favor, contributing to the considerable expansion of oak populations across the continent. Subsistence relationships led to values and virtues being associated with trees, which ultimately became symbols in societies, or to the association of sacred beings with trees in religions (Logan, 2006). The shared history of humans and oaks also generated cultural and emotional relationships that translated into symbols or were incorporated into religious beliefs and

mythology. Oaks were associated with the most important gods (e.g. Zeus, Jupiter and Thor) in the mythology of ancient Greece and Rome, and were often used to represent the *axis mundi*, the center of the world, in Celtic and Germanic cultures.

Since the earliest ties between humans and oaks, a very strong symbolic image of oaks has developed, in which these trees have become associated with longevity, strength, stability, endurance, fertility, power, justice and honesty. These symbols have been widely studied in the humanities, in history (Logan, 2006), social sciences (Mazoyer and Rey, 2003; Brosse, 1989), literature (Corbin, 2005) and the arts (Farcas *et al.*, 2015). In natural sciences, the first insights into oak symbolism emerged from very early botanical writings by Theophrastus (*Historia Plantarum, Book 3, Wild trees and shrubs*) and Pliny the Elder (*Natural History*). Theophrastus provided the first taxonomic description of oaks in ancient Greece, but he also noted the longevity of these species in his scientific approach (Thanos, 2005). Pliny the Elder provided a detailed description of the ways in which druids celebrated mistletoe growing on sacred oak trees and its use to prepare remedies to cure infertility (Brunaux, 2006). Much more recently, many Northern Hemisphere countries, such as the United Kingdom, Poland, Portugal and Germany, and a number of regions have adopted the “mighty” oak as an official national emblem (Figure 1). In 2001, a nationwide popular vote took place over a period of four months, to select the official national tree of the United States of America. The people of the USA selected oaks from a list of 20 tree species (International Oak Society, 2001) and Congress then recognized the oak as the national tree of the USA in 2004.

## IV Longevity

Unlike animals, which are generally less long-lived than humans (Bozek *et al.*, 2017), tens of thousands of plant species, including many oak species, can live for hundreds of years. Majestic ancient living oaks in public squares, parks and forests are a strong element of Northern Hemisphere cultures (Farjon, 2017; Pater, 2017). Observations of the same old oaks enduring throughout a lifetime may trigger positive memories, much like Proust’s madeleine (Proust, 1913). Some famous specimens, such as the Major Oak of Sherwood Forest in England, are thought to be up to 900 years old (Farjon, 2017).

However, the widely held view that oaks have a long lifespan merits qualification. Anyone who has ever walked under a majestic oak tree will already have noticed the huge number of seedlings growing under its canopy. Total seed production and germination are difficult to evaluate, but it has been estimated that there may be 200,000 to 1,000,000 new sessile oak seedlings per year and per hectare within sessile oak-dominated forests (Jarret, 2004). Ten years later, in the absence of silvicultural disturbance, the number of living individuals per hectare has been estimated at 57,000 to 63,000, suggesting that the vast majority (68-94%) of these new individuals die within ten years due to biotic and abiotic stresses (Jarret, 2004). Some individual oaks may be long-lived, but the life expectancy of any given oak seedling is very short indeed.

But what are the genetic consequences of growing old? Several studies in primates have shown that aging has a strong impact on intergenerational heritable mutation rates,

particularly due to the number of germline cell divisions in males (e.g. Jónsson *et al.*, 2017; Thomas *et al.*, 2018). In plants, the germline cell lineage is generally assumed to segregate and differentiate from somatic cells at the end of stems or branches (but see Lanfear, 2018). Consequently, *de novo* mutations would be expected to accumulate throughout plant growth, and, might potentially be passed on to the progeny. Heritable mutations would therefore be expected to accumulate with age, particularly in long-lived species, such as oaks. Two recent independent studies tested this hypothesis, by comparing whole-genome sequence data from oak leaves or buds collected at two or three locations on 236- and 100-year-old trees (Schmid-Siegert *et al.*, 2017; Plomion *et al.*, 2018, respectively). They used very different methods and data, with potential implications for interpretation (Plomion *et al.*, 2018), but both reported only small numbers of mutations (17 and 46 SNPs over a genome of more than 750 million bases; Kremer *et al.*, 2007). These findings challenge the poetic vision of each tree as a forest in its own right (Hallé, 2005). After a detailed analysis at the interface between population genetics and philosophy, Gerber (2018) reached a similar conclusion. These findings suggest that taller plants may have low rates of mutation per unit time, consistent with the conclusions of Lanfear *et al.* (2013), based on comparisons of molecular evolution rates between herbs, shrubs and trees (i.e. plants of different statures).

## V Cohesiveness

Since ancient times, oaks have been seen as cohesive species. In Celtic cultures for example, *Dara*, which means *oak tree*, is a specific type of knot created from an endless cord forming interlaced patterns symbolising eternity and unity. However subtle morphological differences between closely related species have remained unrecognized even in recent times, as illustrated by the use of the generic word “oak” indifferently for *Q. robur* or *Q. petraea*. But beyond such symbolism, are oak species genetically cohesive units?

Over the last decade, several studies have tried to clarify oak taxonomy (Hubert *et al.*, 2014; Denk *et al.*, 2017; Hipp *et al.*, 2019). They have proposed a comprehensive backbone for the evolutionary history of the genus *Quercus*. The latest infrageneric classification clusters oak species into two subgenera (*Quercus* and *Cerris*) with eight sections (*Quercus*, *Ponticae*, *Virentes*, *Protobalanus*, *Lobatae*, *Ilex*, *Cerris* and *Cyclobalanopsis*). Hybridization between species from different sections is rare in nature, consistent with reproductive isolation (Hubert *et al.*, 2014), but hybridization within sections is widespread (Hipp, 2015). This phenomenon led Rieseberg and coworkers (2006) to include oaks in their shortlist of ‘botanical horror’ taxa. Darwin was aware of the complexity of oak species relationships (Darwin 1859), which he considered a “thorny problem”: “[...] in this country the highest botanical authorities and practical men can be quoted to show that the sessile and pedunculated oaks are either good and distinct species or mere varieties” (Darwin, 1859).

Many population genetics (e.g. Curtu *et al.*, 2007; Lepais *et al.*, 2009; Leroy *et al.*, 2017; Ortego *et al.*, 2014) and population genomics studies (Leroy *et al.*, 2019a; Ortego *et al.*, 2018) have provided empirical support for ongoing admixture and gene flow between phylogenetically related oak species, raising the possibility that oak species evolve as independent evolutionary units. Whole-genome sequencing has shown that hybridization can occur without disrupting species integrity, which is restricted to a limited part of the genome,

maintaining species barriers (Leroy et al. 2019a). Studies based on the methodology of Roux *et al.* (2016) have shown that the intensity of interspecific gene flow between divergent oak populations or species may have varied over time, with the advance and retreat of glaciation (e.g. Leroy *et al.*, 2017; 2019a; Merceron *et al.*, 2017). Genetic inferences and genome scans for differentiation have suggested that gene flow is heterogeneous over time, space and genomic regions (Lang *et al.*, 2018; Leroy *et al.*, 2017; 2019a). Barriers to mating dispersed throughout the genome may partially prevent interspecific gene flow, ensuring that interspecific differences are essentially fixed, whereas interspecific gene flow rates are high elsewhere in the genome. These recent results tell us that cohesiveness should be viewed at the level of the gene — at which species barriers are maintained — rather than at the genome level, which is largely permeable to gene flow in these outcrossing species. However, such interspecific exchanges should not be disregarded, because they can contribute to local adaptation in populations living in marginal habitats (Leroy et al. 2019b). Further population genomic studies will be required to improve our understanding of current evolutionary trajectories, and of the adaptive potential of oaks in a changing environment.

## VI Robustness

Pedunculate oak (*Q. robur*) is the most abundant oak species in Europe. The properties of its wood probably explain the Latin meaning of its name: “strength”. Figurative representations of oaks are common on military medals and decorations, including current awards for distinguished service and bravery in the United States and Germany. Artistic representations of oak trees have also been used to symbolize the strength and robustness of ideas (Figure 2). Based on the information provided by the oak genome, can we consider oaks to be robust enough to neutralize attacking enemies?

The pedunculate oak genome consortium recently revealed patterns of immune system diversification in this species (Plomion *et al.*, 2018). Their analyses support both an expansion of resistance (R) genes (accounting for 9% of the gene catalog) and a diversification of gene function driven by a long-standing co-evolutionary arms race between oaks and their natural enemies (viruses, bacteria, fungi, oomycetes, nematodes and insects). However, oaks are far from invulnerable. On the contrary, several plant pathogens are currently a major source of concern, including the causal agent of acute oak decline, which has recently spread in the UK (Brown *et al.*, 2016). The globalization of world trade has increased the dissemination of alien pests and pathogens that have not coevolved with native species, creating new threats to oaks. The accidental introduction of a powdery mildew (*Erysiphe alphitoides*) into Europe at the start of the 20<sup>th</sup> century led, for example, to high rates of mortality in Pyrenean oak (*Q. pyrenaica*) in Southwestern and Western France (Desprez-Loustau *et al.*, 2011). This pathogen is still having a deleterious impact on tree growth in European forests (Bert *et al.*, 2016). The recent emergence of *Phytophthora ramorum*, causing sudden oak death on the West Coast of the United States, is another major cause for concern (Cunniffe *et al.*, 2016). Given the vulnerability of oaks to these pathogens and the intrinsic difficulties of managing invasive forest pathogens, greater attention should be paid to preventing new introductions, thereby limiting the risk of devastating new outbreaks.

## VII Conclusions

Symbols are common literary elements transforming the complexity of reality into something easier to understand, thereby enabling writers to impart significant meaning and emotion. Some oak symbols are so ancient and powerful that they actually distort reality, influencing our lives and our way of thinking. In recent years, this gap between the symbol and reality has been widened by pseudoscientific literary essays and movies deeply rooted in this symbolism that have received considerable media and public attention, to the point of becoming worldwide bestsellers (e.g. ‘*The hidden life of trees*’, Wohlleben, 2016). Our objective in this brief excursion into oak symbolism was to call for more thorough scientific assessments and the use of scientific findings to examine the relevance of some of these symbols. We are convinced that recent academic discoveries, such as the importance of archaeological diving prospecting for studies of oak evolution through the sequencing of ancient waterlogged remains (Wagner *et al.*, 2018), will be of interest to the general public. We anticipate that the increase in genomic resources for oaks (Plomion *et al.*, 2016; 2018; Sork *et al.*, 2016; Ramos *et al.*, 2018, and this issue) will reveal a myriad of surprises, provided that analyses are data-driven and free from cultural assumptions.

This review argues for future investigations of genetic properties relating to the three symbolic representations of oaks discussed here. For longevity, future research could refine the relationship between tree aging and mutation rates, accounting for meiotic and somatic mutations. Ultimately, age-related mutation rates in different long-lived animal and plants would provide clues as to whether there is a “generation-time effect” (Moorjani *et al.*, 2016), e.g. with a lower yearly mutation rate in woody species. For cohesiveness, our conclusions are, for the time being, mostly driven by emblematic examples, such as European white oaks (but see Hipp *et al.* 2019). The cohesiveness at genome level of species traditionally considered to be taxonomic nightmares supports recent proposals for taxonomic standardization based on genomic data (see Galtier, 2018). Taxonomic decisions could be made less arbitrary by analyzing the population structure of a large set of populations from diverse geographic regions and assuming a threshold for the net amount of differentiation or divergence for oaks. Such an approach would contribute to a scientifically more rigorous definition of oak species. For robustness, we are beginning to establish an inventory of the oak leaf microbiome and rhizosphere. Combined metagenomic and expression studies would show whether the expanded R-genes in the oak genome contribute to an extended microbial phenotype and, ultimately, to greater resilience of oaks to emerging biotic and abiotic threats.

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**Figure 1. An illustrative example of the endorsement of oak trees as a symbol of human values: the Gernika tree, a pedunculate oak standing in front of the parliament of the Biscay province of the Basque Country.**

a) A third-generation offspring of the original Gernika tree in front of *la Casa de Juntas* (House of Assemblies), Basque Country, Spain. This tree became a symbol of liberty, as leaders of the Biscay province and, later, of the Basque people as a whole, swore an oath beneath the tree to safeguard the freedom of the Basque people. b) The Gernika tree depicted in a stained-glass window in the ceiling of *la Casa de Juntas* showing a Lord of Biscay swearing an oath to safeguard the freedom of the Basque people. c) Painting from *la Casa de Juntas* showing one of the first assemblies of representatives beneath the Gernika tree. General assemblies of people's representatives from villages and provinces subsequently met near the tree to pass laws, and the tree became a symbol of justice and the unity of the Basque people. The French philosopher Jean Jacques Rousseau cited these assemblies as an early form of democracy in "Le contrat social" (1762). d) Biscay coat of arms from *la Casa de Juntas*, with its representation of the Gernika tree. e) Poster of the 650<sup>th</sup> anniversary of the establishment of Gernika.



**Figure 2. Oak tree artwork as symbol of robustness of ideas (Bloomfield, 2012).**

An illustrative symbolic oak representation linking art and science was the choice for a longitudinal oak tree section as artwork for the ceiling of the Natural History Museum of London in 2009. The artwork was planned to celebrate the 200<sup>th</sup> anniversary of Darwin. A panel of art experts, museum experts and scientist selected the oak project from 10 proposals made by different artists. The work TREE proposed by artist Tania Kovats was chosen by unanimous decision. Her proposal was to embed in the ceiling a longitudinal 3 to 5 mm thick section of a 200 year old entire oak tree, from the ground to the canopy). The tree was 21 metres high and came from a managed forest of Longleat Estate in Wiltshire. The art work was meant as a replicate of the iconic drawing of the “Tree of life” by Darwin, which echoes in this contribution the “Tree of life” revered by early human populations (Andersson, 2007). As described by R.M. Bloomfield (2012) “Rendered in stalwart English oak, the artwork is a metaphor of the endurance of Darwin’s ideas, as well as Darwin’s own bravery and commitment to them”.

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