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Bringing Back the Dire Wolf: How CRISPR Technology is Revolutionizing De-Extinction

By

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Introduction

The prospect of resurrecting species that have vanished from the Earth is a concept that deeply stirs both scientific curiosity and public imagination. Among the array of extinct creatures, the dire wolf (Aenocyon dirus) holds a particular allure [1]. Known as prominent predators of the American Ice Age, these large canids roamed North America before their disappearance in the terminal Pleistocene [2]. The fascination with the dire wolf stems from several factors, including its formidable nature as a predator and the significant amount of fossil material available for study. Notably, thousands of wellpreserved mammalian bones, including dire wolves, have been excavated from sites like the Rancho La Brea Tar Pits in metropolitan Los Angeles, offering a remarkable window into past communities [3]. The ability to obtain genetically informative DNA from these preserved bones, particularly from locations like Rancho La Brea, raises the prospect of deeper insights [4]. Studies of the dire wolf's craniofacial

Abstract

Thanks in large part to developments in genomic technology, the idea of de-extinction the resuscitation of extinct species has quickly moved from speculative science fiction to a realistic scientific endeavor. Using the famous dire wolf (Aenocyon dirus) as a case study, this article examines the ethical, technological, and scientific aspects of de-extinction. It demonstrates how precise genome editing using ancient DNA extracted from fossil remains made possible by CRISPR-Cas9 has revolutionized the viability of this discipline. The paper highlights the distinct biological and cultural attraction of the dire wolf while examining two main de-extinction strategies: Somatic Cell Nuclear Transfer (SCNT) and CRISPR-driven gene editing. It also explores the practical difficulties of surrogate gestation, the evolutionary separation of the dire wolf from contemporary canids, and the reconstruction of extinct species, including ethical concerns, ecological risks, and the tension between techno-optimism and conservation pragmatism.

Keywords: De-extinction, CRISPR-Cas9, Dire Wolf, Ancient DNA, Genome Editing, Resurrection Biology, Somatic Cell Nuclear Transfer, Paleogenomics, Synthetic Biology

morphology reveal distinct features compared to the living gray wolf (Canis lupus), such as similarly wide zygomatic arches and a relatively longer temporal fossa, suggesting a larger temporalis muscle and greater bite strength [5]. They also possessed a larger backward projection of the inion. Furthermore, the eastern population of dire wolves was estimated to be approximately 15% heavier than the western population, which in turn was 25% heavier than the extant gray wolf. This wealth of morphological and increasingly, genetic information including the reconstruction of their nuclear paleogenome positions the dire wolf as a compelling subject when considering the possibility of bringing extinct species back to life [6]. This brings us to the concept of deextinction, also discussed under terms like species revivalism or reanimation. From a conservation biology perspective, the aim of de-extinction is not necessarily to restore the "status quo ante" [7]. Instead, it involves producing organisms that closely resemble extinct species, potentially by creating "proxy" organisms designed to fill the ecological niches of

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their extinct counterparts [8]. This ambitious undertaking has gained considerable attention, prompting discussions about its potential ethical, political, and ecological consequences. Addressing the feasibility of de-extinction is intrinsically linked to advancements in scientific technology [9]. This is where CRISPR-Cas9 has emerged as a revolutionary tool. Often described as "molecular scissors", CRISPR is a powerful genome editing tool that originated from the natural defense systems of bacteria, allowing them to program proteins to target and destroy viral DNA [10]. Scientists have been able to repurpose this fundamental property for a variety of uses, including making precise alterations to DNA. This technology relies on DNA-RNA interactions and is notably easier and more efficient than previous gene editing methods [11]. The advent of CRISPR and related techniques has dramatically increased the perceived likelihood of reviving extinct species by offering unprecedented opportunities for genetic manipulation. The ability to reconstruct a paleogenome from ancient DNA samples, and to iteratively refine this reconstruction to recover more genetic data, is a critical step enabled by modern genomic technologies [12]. CRISPR then provides the means to edit and assemble this genetic material into a viable form, fundamentally changing the approach to bringing species back from extinction [13].

The Science Behind De-Extinction

De-extinction, also called resurrection biology, is the process of creating organisms that are similar to extinct species, often to act as ecological proxies. One important technique is Somatic Cell Nuclear Transfer (SCNT), which involves inserting the nucleus of an extinct species into an enucleated egg of a living relative [14]. Although this method has produced live offspring in some species, there are still issues, such as low success rates and the impact of the cytoplasm of the egg on the resulting phenotype [15]. More recently, genetic engineering, specifically CRISPR-Cas9, has become a key component of de-extinction because it allows precise DNA edits, allowing the incorporation of traits from extinct species into living genomes. Scientists may recreate parts of the extinct genome using ancient DNA from fossil remnants, and then utilize CRISPR to modify live cells in accordance with those reconstructions [16]. Because of its ecological relevance and extensive fossil record, particularly from Rancho La Brea, the dire wolf (Aenocyon dirus) is a prime candidate for de-extinction. Dire wolves were apex predators that occupied a special place in Ice Age ecosystems; they were heavier and more resilient than gray wolves today. Scientists have been able to sequence nuclear genomes from their well-preserved bones, which has given them a workable blueprint for genetic reconstruction [17]. The dire wolf is a fascinating topic for science and culture in the pursuit of species resuscitation because of its sophisticated tools and well-preserved genetic material [18].

Enter CRISPR: A Genetic Revolution

The discovery of CRISPR-Cas9 has transformed molecular biology and transformed genetic engineering. Initially developed as a defensive mechanism for bacteria, CRISPR is known as a "molecular scissor" technique because it enables researchers to create extremely precise cuts in DNA using a guide RNA and the Cas9 protein [19]. Compared to previous technologies like ZFNs and TALENs, this precise tool makes it easier and more accurate to make targeted insertions, deletions, or alterations to the genome. Its versatility also enables modular applications like gene silencing or activation utilizing variations like dCas9, as well as multiplexing, which allows many genes to be edited simultaneously [20]. Deextinction and other complicated gene editing initiatives are now more feasible because to CRISPR's ease of use and effectiveness. It makes it possible to insert old DNA sequences into the genomes of living relatives, maybe bringing back extinct characteristics or whole species [21]. Therefore, CRISPR represents a profound change in our understanding of biological design and species restoration, not merely a genetic triumph. As depicted in Figure 1, this precise tool allows for targeted insertions, deletions, or alterations to the genome, making it easier and more accurate than previous gene editing methods.

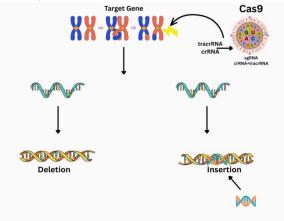


Figure 1: Simplified diagram illustrating the mechanism of CRISPR-Cas9 gene editing. The Cas9 protein, guided by a guide RNA (sgRNA, formed by tracrRNA and crRNA), targets a specific gene. This allows for precise modifications, such as deletion or insertion of DNA sequences, fundamentally changing genetic material.

Reconstructing the Dire Wolf Genome

Reconstructing the genome of an extinct species, such as the dire wolf (Aenocyon dirus), is a fundamental step in understanding its biology, evolutionary history, and the potential for de-extinction efforts [22]. This process involves recovering ancient DNA from preserved remains and using advanced genomic techniques to piece together the complete genetic blueprint. Feasibility and Challenges Obtaining viable DNA from ancient specimens presents significant feasibility and challenges. Ancient DNA is typically degraded into short fragments over time and can be chemically modified, notably through deamination resulting in cytosine to thymine (C>T) transitions. Contamination with modern DNA and the presence of environmental inhibitors, such as tar in the case of fossils from Rancho La Brea, also complicate DNA extraction and sequencing [23]. Despite these difficulties, researchers successfully recovered ancient DNA from five dire wolf specimens found in different North American locations,

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including Ohio and Idaho. These samples date from approximately 12,900 to over 72,000 years ago. Tissues like incisor roots and petrous bones were targeted as they can offer better DNA preservation. DNA extraction protocols were optimized for recovering short, degraded molecules, and in some cases, pre-treatment with bleach was used. Singlestranded DNA libraries were generated for some samples to capture more fragments and better assess damage patterns. While ancient DNA often yields low-coverage nuclear genomes, improved paleogenomics approaches allowed for higher coverage in key specimens, such as the DireGB individual, which reached 12.8× coverage. Once ancient DNA reads are obtained, they undergo extensive pre-processing to remove adapter sequences, low-quality ends, and potential contaminants [24]. A major hurdle in genome reconstruction is reference bias. This occurs when mapping ancient DNA reads to a reference genome from a living, related species (like the grey wolf) because the evolutionary divergence between the extinct and living species can lead to inaccurate mapping and inflate the apparent sequence similarity. To address this, an iterative mapping and polishing approach was employed. Reads were initially mapped to a divergent seed reference (Greenland grey wolf), a consensus sequence was generated from the mapped reads, and the reads were then remapped to this newly constructed, dire wolf-like consensus reference [25]. This process was repeated until no further reads could be added. This iterative method, combined with conservative consensus calling (masking regions with low coverage), significantly reduced reference bias and improved the overall coverage and mappability of the dire wolf genome. Multi-species pangenomes were also used as an alternative mapping strategy to minimize bias from using a single, potentially distant reference. Sequence variants, such as single nucleotide polymorphisms (SNPs) and small insertions/deletions, were then called against the reconstructed dire wolf reference genome. Wolves, Dogs, and the Dire Wolf With reconstructed dire wolf paleogenomes in hand, comparative genomics analyses were performed by comparing the dire wolf's genetic sequence to those of various living canid species [26]. These comparisons included grey wolves (Canis lupus), coyotes (Canis latrans), domestic dogs (Canis familiaris), dholes (Cuon alpinus), African wild dogs (Lycaon pictus), and African jackals (Canis anthus, C. mesomelas, C. adustus), among others. Using methods like phylogenetic analyses, species tree estimations, and D-statistics (to test for gene flow/admixture), researchers sought to clarify the dire wolf's evolutionary position and history. These comparative analyses revealed that despite sharing some morphological similarities with the grey wolf, the dire wolf was actually a highly divergent lineage within the canid family tree [27]. The dire wolf lineage split from the clade containing extant wolflike canids (such as grey wolves, coyotes, and dholes) approximately 5.7 million years ago. Importantly, the genomic data showed no significant evidence of gene flow (hybridisation) between dire wolves and North American grey wolves or coyotes since their lineages diverged. This lack of admixture, despite overlapping ranges during the Late Pleistocene, suggests that the dire wolf evolved in relative

isolation from the ancestors of modern North American canids. The analyses did, however, detect a signature of ancient admixture occurring around 3 million years ago between the lineage ancestral to dire wolves and the lineage ancestral to dholes, wolves, and coyotes [28]. Comparative genomics also allowed for the identification of genes that likely contributed to the dire wolf's specific adaptations, including genes under selection related to their large size and role as predators of large herbivores, providing insight into the molecular mechanisms behind their ecological niche. The reconstructed paleogenome is a valuable resource enabling future studies into this iconic extinct predator.

From Cells to **Species: Pathways** to Resurrection

The ambition to resurrect extinct species, or at least recover their genetic potential, fundamentally relies on bridging the gap between preserved biological material (cells) and the creation of a living organism (species) [29]. This requires sophisticated biotechnological approaches, primarily revolving around Somatic Cell Nuclear Transfer (SCNT) and the subsequent development of the resulting embryo through pathways that might involve synthetic embryology techniques and surrogacy. These methods represent potential pathways to bring the genetic information held within ancient cells back to life. SCNT is a method of cloning that aims to generate genetic copies of an individual [30]. The core technique involves transferring the nucleus from a somatic cell (any differentiated body cell, such as a skin cell or fibroblast) of the donor individual into an enucleated oocyte (an egg cell from which the original nucleus has been removed). This process was famously used to create Dolly the sheep, the first mammal cloned from an adult somatic cell, in 1996. Since then, SCNT has been successfully applied to clone at least 24 mammalian species, including various livestock, laboratory animals, and wild species such as cats, dogs, camels, and monkeys [31]. A wide variety of donor cell types can be used, and remarkably, successful cloning has been achieved even from non-viable cells or tissues stored postmortem for years without cryoprotectants, provided the nuclear material remains intact. This opens up the possibility of using preserved tissues from extinct animals. The SCNT process involves several steps: obtaining donor cells, preparing enucleated oocytes, extracting the donor nucleus, transferring it into the oocyte (often into the perivitelline space or cytoplasm), and fusing the donor cell with the oocyte (e.g., by electrofusion or using inactivated virus). The resulting reconstructed oocyte must then be activated to initiate embryonic development, mimicking the calcium release triggered by sperm in natural fertilization [32]. This activation, along with factors within the oocyte cytoplasm, begins the crucial process of nuclear reprogramming, which aims to erase the somatic cell's differentiated epigenetic memory and restore the nucleus to a totipotent, embryonic state capable of directing full development. Despite its successes, SCNT efficiency remains relatively low. A significant challenge is the incomplete or inappropriate epigenetic reprogramming of the donor nucleus, which can

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lead to high rates of embryonic and fetal mortality and developmental abnormalities [33]. When attempting to clone an extinct species using oocytes from a different, living species (known as interspecies SCNT or iSCNT), additional challenges arise, including potential incompatibilities between the donor nucleus and the recipient oocyte's cytoplasm, and the presence of mitochondrial DNA (mtDNA) from the recipient oocvte, leading to mitochondrial heteroplasmy. While iSCNT has resulted in offspring for some closely related species, outcomes have often been poor, although recent conservation efforts have seen success in species like the black-footed ferret and Przewalski's horse using Iscnt [34]. Following successful SCNT, the reconstructed embryo is typically cultured in vitro for a period, often until the blastocyst stage. This in vitro culture and manipulation can be considered a form of "synthetic embryology" in the sense of supporting and guiding embryonic development outside the body. The ability to culture embryos in vitro is essential as it allows assessment of developmental progress before implantation. More broadly, the field of cell reprogramming has led to the ability to generate induced pluripotent stem cells (iPSCs) from somatic cells, and a key recent development is the emerging capacity to differentiate these pluripotent cells in vitro into germ cells (oocytes and spermatozoa) [35]. This offers a theoretical pathway to generate reproductive potential, and thus propagate genetics, from biobanked nonreproductive tissues. Using iPSCs as nuclear donors in SCNT is also being explored to potentially improve efficiency due to reduced reprogramming requirements. However, to progress from an in vitro embryo (whether derived from SCNT or other means) to a living animal, the embryo must be transferred into the uterus of a recipient female. This female acts as a surrogate mother, carrying the pregnancy to term and giving birth to the cloned offspring. For cloning extinct species, a closely related living species would need to serve as the surrogate host [36]. Successful surrogacy requires careful synchronization of the recipient animal's reproductive cycle with the developmental stage of the transferred embryo. Establishing reliable protocols for embryo transfer and surrogacy, especially in non-domesticated and endangered species, presents significant practical and ethical challenges. Figure 2, visually outlines this cloning method, detailing how a somatic cell's nucleus is used to reprogram an enucleated egg cell, initiating the formation of an embryo.

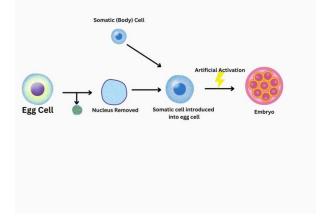


Figure 2: Illustration of the Somatic Cell Nuclear Transfer (SCNT) process. The nucleus from a somatic (body) cell is transferred into an enucleated egg cell. This reconstructed egg is then artificially activated to initiate embryonic development, leading to the formation of an embryo.

Ethical and Ecological Implications

The application of modern genetic technologies and advanced assisted reproductive technologies (aART) in conservation presents significant ethical and ecological challenges. These interventions, such as gene editing tools like CRISPR and techniques used in de-extinction efforts, represent powerful ways humans can alter the natural world [37]. Ethically, concerns arise regarding animal welfare, particularly for individuals undergoing procedures and potential surrogate animals involved in aART or de-extinction processes. The ability of tools like CRISPR to precisely alter DNA sequences raises questions about the extent of human control over evolution and the potential for unintended consequences on species themselves. There are also ethical considerations about who decides which species are targeted for intervention and the potential for actions based on "nonhuman eugenics" or "ecological xenophobia". The use of genetic technologies to eradicate species could even be likened to a form of "species genocide" from certain ethical viewpoints [38]. Engaging with diverse perspectives, including those from Indigenous communities, is important when discussing the uses, misuses, and limitations of these technologies. Furthermore, a precautionary approach to synthetic biology has been agreed upon by international bodies. Ecologically, introducing genetically modified or resurrected organisms into existing environments carries inherent risks. Such introductions could potentially act like invasive species, potentially reducing overall biodiversity despite the addition of a specific species. Predicting all negative ecological consequences is difficult due to the complex interactions within ecosystems and our incomplete understanding of how genetic changes manifest in real-world environments [39]. Introducing a species that is an approximation of an extinct one, lacking its original behaviour and ecological context, might disrupt food webs or negatively impact other species.

The concept of "de-extinction" is a prominent example of genetic technologies in conservation. Arguments for deextinction sometimes cite restitutive justice, suggesting a moral obligation to rectify past human-caused extinctions. However, making "reparations" to organisms that no longer exist is ethically complex. A stronger case for restitutive justice might be made in rare instances where human action eliminated the last reproductive partners, allowing aART to provide them. Another argument is based on forward-looking reasons, such as restoring lost ecological functions, like the proposed role of resurrected woolly mammoths in transforming Arctic landscapes [40]. It is important to recognize that current technologies are more likely to produce "proxies" rather than exact replicas of extinct species. These proxies may differ genetically and biologically from the original species and crucially lack their original microbiome,

learned behaviors (animal culture), and historical ecosystem context.

Beyond the Dire Wolf: The Future of De-**Extinction**

The concept of de-extinction is increasingly receiving attention in both popular and scientific discourse. It relies on powerful biotechnologies, including gene reading, gene synthesis, and genome editing (like CRISPR). These tools have increased the likelihood of reviving extinct species, making practices like de-extinction potentially feasible [41]. Proponents argue for de-extinction based on restitutive justice, suggesting a moral obligation to bring back species humans drove to extinction to "right a wrong". Another forwardlooking rationale is to restore lost ecological functions. For example, resurrecting the woolly mammoth is proposed for its potential role in converting Arctic tundra back to grassland, which could help mitigate climate change. Some also see deextinction as a way to shift the conservation narrative from one of loss to one of hope and excitement. Current technologies are more likely to produce "proxies" or "genetic chimeras" rather than exact replicas of extinct species, as they may differ genetically, biologically, and lack original learned behaviors and ecological context [42]. There are concerns about the "speculative ethics" around these technologies, which may overlook problems caused by reductive thinking and neglect of non-human agency. Introducing these organisms into ecosystems poses significant risks, as they could function like invasive species, disrupt food webs, or lead to other unintended consequences. The full ecological impacts may not be evident until the technology is widely deployed, leading to a potential "technology control dilemma" where adverse effects are costly or impossible to reverse. Ethical concerns about animal welfare also persist when applying assisted reproductive technologies for de-extinction. Resources invested in de-extinction might also be diverted from more traditional, proven conservation efforts like habitat protection [43]. Overall, while paleogenomic research on species like the dire wolf contributes foundational knowledge, the future of de-extinction extends to engineering "Anthropocene Organisms" using advanced genetic technologies. This presents both potential ecological benefits and significant ethical challenges, ecological risks, and uncertainties regarding ecosystem integration and unintended consequences. The debate involves differing views, including techno-optimism (ecomodernism) and techno-scepticism. A precautionary approach and collective governance are suggested for navigating these complex choices [44].

Conclusion

The hope of bringing extinct animals back to life is now based on actual scientific possibility rather than being a work of fantasy. De-extinction has gained credibility as a study area because to technologies like CRISPR and sophisticated genome reconstruction methods. An example of how a wellpreserved species might become the focus of this effort is the dire wolf, with its extensive fossil record and ecological significance. The difficulties are not just scientific, even

though the technological viability is becoming better. The use of these potent technologies must be guided by ethical, ecological, and philosophical considerations. De-extinction creates the possibility of proxy creatures that may inspire new conservation efforts or restore lost ecological roles, even if it might not produce identical reproductions. As we navigate this frontier, a cautious, interdisciplinary approach will be essential to balance innovation with responsibility.

References

- 1. Preston CJ. De-extinction and Gene Drives: The Engineering of Anthropocene Organisms. InAnimals in Our Midst: The Challenges of Coexisting with Animals in the Anthropocene 2021 Apr 30 (pp. 495-511). Cham: Springer International Publishing.
- 2. Schwarz V. Endangered Species, De-Extinction and MANipulation.
- 3. Thiele LP. Nature 4.0: Assisted evolution, deextinction, and ecological restoration technologies. Global Environmental Politics. 2020 Aug 1;20(3):9-27.
- 4 Cutter AD. Evolving Tomorrow: Genetic Engineering and the Evolutionary Future of the Anthropocene. Oxford University Press; 2023 Jun 14.
- 5. Preston CJ. The synthetic age: Outdesigning evolution, resurrecting species, and reengineering our world. MIT Press; 2019 Feb 19.
- Hirsch RE. De-Extinction: The Science of Bringing 6. Lost Species Back to Life. Twenty-First Century Books (Tm); 2017.
- 7. Greely HT. Is de-extinction special?. Hastings Center Report. 2017 Jul;47:S30-6.
- Thiele LP. Nature 4.0: Assisted evolution, de-8. extinction, and ecological restoration technologies. Global Environmental Politics. 2020 Aug 1;20(3):9-27.
- 9. Preston CJ. De-extinction and Gene Drives: The Engineering of Anthropocene Organisms. InAnimals in Our Midst: The Challenges of Coexisting with Animals in the Anthropocene 2021 Apr 30 (pp. 495-511). Cham: Springer International Publishing.
- 10. Chen YN. The Potential of CRISPR-Cas9 Genome Editing on the Woolly Mammoth Revival. Young Anthropology. 2024 Dec 31;5:15-22.
- 11. Erwin JA. Changing genes for a changing climate: genetic intervention as a tool for biodiversity conservation in the era of climate change. InResearch Handbook on Climate Change and Biodiversity Law 2024 Dec 10 (pp. 315-336). Edward Elgar Publishing.
- 12. Wray B. Rise of the necrofauna: The science, ethics, and risks of de-extinction. Greystone Books Ltd; 2017 Sep 30.

- 13. Adams WM. Geographies of conservation I: Deextinction and precision conservation. Progress in Human Geography. 2017 Aug;41(4):534-45.
- 14. El-Khatib H. HARNESSING CRISPR FOR CONSERVATION: GENE EDITING AS A TOOL FOR BIODIVERSITY PRESERVATION IN THE FACE OF CLIMATE CHANGE. Modern American Journal of Biological and Environmental Sciences. 2025 Apr 11;1(1):1-6.
- 15. Cutter AD. Evolving Tomorrow: Genetic Engineering and the Evolutionary Future of the Anthropocene. Oxford University Press; 2023 Jun 14.
- 16. Preston CJ. De-extinction and Taking Control of Earth's "Metabolism". Hastings Center Report. 2017 Jul;47:S37-42.
- 17. Seddon PJ. The ecology of de-extinction. Functional ecology. 2017 May 1;31(5):992-5.
- 18. Greely HT. Is de-extinction special?. Hastings Center Report. 2017 Jul;47:S30-6.
- 19. Perri AR, Mitchell KJ, Mouton A, Álvarez-Carretero S, Hulme-Beaman A, Haile J, Jamieson A, Meachen J, Lin AT, Schubert BW, Ameen C. Dire wolves were the last of an ancient New World canid lineage. Nature. 2021 Mar 4;591(7848):87-91.
- 20. Dundas RG. Quaternary records of the dire wolf, Canis dirus, in North and South America. Boreas. 1999 Sep;28(3):375-85.
- 21. Anyonge W, Baker A. Craniofacial morphology and feeding behavior in Canis dirus, the extinct Pleistocene dire wolf. Journal of Zoology. 2006 Jul;269(3):309-16.
- 22. Gedman G, Morrill Pirovich K, Oppenheimer J, Hyseni C, Cassatt-Johnstone M, Alexandre N, Troy W, Chao C, Fedrigo O, Hoyt SJ, Grady PG. On the ancestry and evolution of the extinct dire wolf. bioRxiv. 2025:2025-04.
- 23. Stevenson M. Dire wolf systematics and behavior. InWolf and Man 1978 Jan 1 (pp. 179-196). Academic Press.
- 24. Anyonge W, Roman C. New body mass estimates for Canis dirus, the extinct Pleistocene dire wolf. Journal of Vertebrate Paleontology. 2006 Jan 1;26(1):209-12.
- 25. Lawler DF, Widga C, Smith GK. Observations of the Acetabulum and Proximal Femur of the Dire Wolf (Canis dirus, Leidy 1854). Journal of Veterinary Anatomy. 2017 Jan 1;10(1).
- 26. O'Keefe FR, Binder WJ, Frost SR, Sadlier RW, Van Valkenburgh B. Cranial morphometrics of the dire wolf, Canis dirus, at Rancho La Brea: temporal variability and its links to nutrient stress and climate. Palaeontologia Electronica. 2014.
- 27. Hodnett JP, Mead JI, Baez A. Dire wolf, Canis dirus (Mammalia; Carnivora; Canidae), from the late Pleistocene (Rancholabrean) of east-central Sonora, Mexico. The Southwestern Naturalist. 2009 Mar;54(1):74-81.

- 28. Brannick AL, Meachen JA, O'Keefe FR. Microevolution of jaw shape in the dire wolf, Canis dirus, at Rancho La Brea. Natural History Museum of Los Angeles County Science Series. 2015;42:23-32.
- 29. Binder WJ, Van Valkenburgh B. A comparison of tooth wear and breakage in Rancho La Brea sabertooth cats and dire wolves across time. Journal of Vertebrate Paleontology. 2010 Jan 29;30(1):255-61.
- 30. Reynolds AR, Lowi-Merri TM, Brannick AL, Seymour KL, Churcher CS, Evans DC. Dire wolf (Canis dirus) from the late Pleistocene of southern Canada (Medicine Hat, Alberta). Journal of Quaternary Science. 2023 Aug;38(6):938-46.
- 31. Hoole B. Evaluating morphological characters used to distinguish dire wolves from other canids.
- 32. Brannick AL. Shape evolution and sexual dimorphism in the mandible of the dire wolf, Canis Dirus, at Rancho la Brea.
- 33. Schmökel H, Farrell A, Balisi MF. Subchondral defects resembling osteochondrosis dissecans in joint surfaces of the extinct saber-toothed cat Smilodon fatalis and dire wolf Aenocyon dirus. PloS one. 2023 Jul 12;18(7):e0287656.
- 34. Gillespy PD, PROTHERO DR, FRENCH J. Ontogenetic change in distal and proximal limb bones of juvenile Pleistocene coyotes (Canis latrans) and dire wolves (Canis dirus) from the Rancho La Brea Tar Pits, California. Fossil Record 6 Volume 1. 2018;79:225.
- 35. Haroon M, Wang X, Afzal R, Zafar MM, Idrees F, Batool M, Khan AS, Imran M. Novel plant breeding techniques shake hands with cereals to increase production. Plants. 2022 Apr 12;11(8):1052.
- 36. Sun K, Liu Y, Zhao W, Ma B, Zhang M, Yu X, Ye Z. Prokaryotic argonaute proteins: a new frontier in point-of-care viral diagnostics. International Journal of Molecular Sciences. 2023 Oct 8;24(19):14987.
- 37. Khan Z, Khan SH, Ahmad A. Challenges and future perspective of CRISPR/Cas technology for crop improvement. CRISPR Crops: The Future of Food Security. 2021:289-306.
- 38. Emrani J, Ahmed M, Newman RH, Thomas MD, Mowa CN, Teleha J. Introducing chemistry students to emerging technologies in gene editing, their applications, and ethical considerations. Journal of Chemical Education. 2020 Jun 12;97(7):1931-43.
- 39. Hamza A, Ahmad MU, Tariq AF, Din AU, Abbas ZE, Bano MI, RAFIQUE RS. CRISPR/cas9: an approach to meet global food demand. Plant Cell Biotechnology and Molecular Biology. 2021:220-33.
- 40. Enríquez P. Genome editing and the jurisprudence of scientific empiricism. Vand. J. Ent. & Tech. L.. 2016;19:603.
- 41. Koonin EV, Makarova KS, Wolf YI. Evolutionary genomics of defense systems in archaea and

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bacteria. Annual review of microbiology. 2017 Sep 8;71(1):233-61.

- 42. Chaudhuri A, Halder K, Abdin MZ, Datta A. CRISPR/Cas9: The Revolutionary Genome Editing Tools for Crop Genetic Tweak to Render It 'Climate Smart'. InCRISPR and Plant Functional Genomics 2024 May 8 (pp. 108-123). CRC Press.
- Khan Z, Ahmad S. 3D's of Genome Editing Systems: Discovery, Developments, and Diversity. InCRISPR-Based Gene Editing 2025 (pp. 1-30). Apple Academic Press.
- Chaudhuri A, Halder K, Abdin MZ, Datta A. 6 CRISPR/Cas9. CRISPR and Plant Functional Genomics. 2024 May 8:108.