

Editorial

Advancing AI, ML, and Bioinformatics for Transformative Research Across Disciplines

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Abstract

In an era where data permeates every aspect of science and society, the demand for advanced analytical tools and intelligent computational systems has never been greater. This editorial introduces the vision and scope of the Journal of Artificial Intelligence, Machine Learning, and Bioinformatics (JAIMLB)—a platform dedicated to fostering interdisciplinary research that integrates cutting-edge developments in AI, ML, and Bioinformatics across diverse scientific fields. JAIMLB emphasizes the transformative role of intelligent systems in modern science. In agriculture, AI and big data analytics are revolutionizing precision farming, crop yield prediction, and climate-resilient practices. In health sciences, predictive modeling, deep learning for diagnostic imaging, and the integration of genomic data with electronic health records are driving personalized medicine. Life sciences and bioinformatics lie at the core of the journal’s mission, featuring research in genomics, proteomics, systems biology, and ecological forecasting—where machine learning enables the interpretation of complex biological systems and evolutionary patterns. Forensic science benefits from AI-assisted pattern recognition, digital forensics, and integrated data systems that enhance accuracy and efficiency in legal investigations. The journal also explores foundational innovations in algorithm development, explainable AI, and ethical considerations that support responsible AI/ML deployment across domains. JAIMLB is committed to bridging disciplinary divides by promoting ethical, data-driven research and facilitating cross-sector collaboration. This inaugural editorial reflects the journal’s commitment to accelerating discovery and advancing the future of scientific inquiry through AI, ML, and bioinformatics.

Keywords: Artificial Intelligence, Machine Learning, Bioinformatics, Interdisciplinary Research, Genomic Analysis, Precision Agriculture, Personalized Medicine, Digital Forensics, Systems Biology, Ethical AI.

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1. Introduction

The omnipresence of data in the modern world necessitates innovative approaches to its analysis, interpretation, and application. As Artificial Intelligence (AI), Machine Learning (ML), and Bioinformatics continue to evolve, their integration into various domains enriches scientific discovery and expands the scope of intelligent data-driven decision-making (Figure 1). This editorial outlines the vision for the Journal of Artificial Intelligence, Machine Learning, and Bioinformatics (JAIMLB), emphasizing interdisciplinary research and collaboration across Agricultural Sciences, Health Sciences, Life Sciences, and Forensic Sciences.

The transformative impact of AI, ML, and Bioinformatics is evident across fields. In agriculture, intelligent systems enhance crop yield, optimize resource use, and assess climate impacts (Kamilaris et al., 2018). In health sciences, predictive models and AI-based diagnostics are revolutionizing patient care and treatment personalization (Shilo et al., 2020). Life sciences benefit from bioinformatics techniques for genomic analysis and evolutionary modeling (Marx, 2013). Forensic science is being reshaped by AI-driven analytics that increase accuracy in criminal investigations (Casey, 2011).

JAIMLB is committed to advancing knowledge at the intersection of these fields, promoting original research, reproducible methodologies, and ethical applications of AI, ML, and Bioinformatics. Through this editorial, we present the scope and future direction of the journal as it contributes to scientific innovation and problem-solving.

2. AI, ML, and Bioinformatics in Agricultural Sciences

The urgency of addressing global food security and sustainable agricultural practices has never been greater. With the global population projected to reach 9.7 billion by 2050, ensuring adequate food supply while preserving environmental resources poses a significant challenge (FAO, 2017). AI, ML, and Bioinformatics offer critical tools to revolutionize traditional farming practices (Figure 2).

Precision Agriculture, a key component of this field, involves the use of GPS, remote sensing, and IoT devices to collect real-time data on crop conditions, soil health, and weather patterns. This data-driven approach enables farmers to make informed decisions about planting, irrigation, fertilization, and pest control, increasing productivity and efficiency (Zhang et al., 2019). Variable rate technology (VRT) allows precise input application, reducing waste and

environmental impact (Mulla, 2013).

Crop yield prediction models utilize historical data, weather forecasts, and satellite imagery to forecast production. ML algorithms analyze diverse datasets to predict the impact of various factors on yields, helping mitigate risks associated with climate change and extreme weather events (Abraham and See, 2000; Schlenker and Roberts, 2009).

Climate impact assessments integrate climate data with crop models to simulate scenarios and develop adaptation strategies. This includes identifying crop varieties resilient to drought, heat, and pests (Webber et al., 2014; Raza et al., 2019).

Future JAIMLB publications will explore big data and AI for optimizing resource management, enhancing pest and disease prediction, and advancing bioengineering. Big data analytics track pest outbreaks based on weather patterns and crop conditions (Westbrook et al., 2016). Bioinformatics supports genetically modified crop innovations that boost productivity and sustainability (Huang et al., 2002).

The socio-economic impacts of smart agriculture will also be discussed, including how technology affects smallholder farmers and addresses the digital divide (Foley et al., 2011).

3. AI and ML in Health Sciences and Biomedicine

AI and ML have revolutionized medical research, diagnostics, and treatment paradigms (Figure 2). JAIMLB will feature predictive modeling, diagnostic AI tools, integrative data platforms, and the ethical implications of data-driven healthcare innovations.

3.1. Revolutionizing Medical Research and Diagnostics

AI-driven analysis of large datasets, including genomic and clinical data, facilitates the discovery of new biomarkers and therapeutic targets (Hood and Flores, 2012). ML algorithms predict disease progression and treatment response (Obermeyer and Emanuel, 2016), especially for chronic conditions (Topol, 2019). These predictive models are being increasingly integrated into clinical decision support systems to assist healthcare providers in tailoring treatment plans. Moreover, such tools enable early intervention strategies that may significantly improve patient outcomes and reduce healthcare costs.

In diagnostic imaging, ML improves accuracy in interpreting radiology, pathology, and dermatology images (Litjens et al., 2017). CNNs, for example, detect tumors and lesions with high accuracy (Esteva et al., 2017). These technologies have shown promise in automating the detection of subtle anomalies often missed by the human eye, especially in high-throughput or resource-limited diagnostic settings.

3.2. Personalized Medicine and Integrative Health Data

By integrating electronic health records (EHRs) with genomic and biometric data, personalized medicine offers more precise treatments tailored to individual patient profiles (Collins and Varmus, 2015). Pharmacogenomics guides medication selection by identifying genetic factors that influence drug efficacy and adverse reactions (Roden et al., 2011). This enables healthcare providers to avoid one-size-fits-all prescriptions and instead adopt data-informed therapeutic strategies. Furthermore, predictive analytics can be used to stratify patients based on risk, optimizing resource allocation and follow-up care.

Wearables and mobile apps enable continuous monitoring, supporting proactive and adaptive healthcare (Wang et al., 2018). These digital tools provide real-time physiological data, allowing for early detection of potential health issues and personalized interventions before conditions worsen.

3.3. Ethical Implications and Data Privacy

Healthcare applications of AI and ML must address privacy, security, and ethical concerns to maintain patient trust and ensure regulatory compliance. Articles will explore frameworks for data security, consent, anonymization, and algorithmic fairness (Safran et al., 2007; Obermeyer et al., 2019). As these technologies increasingly influence diagnosis, treatment, and healthcare access, ensuring transparency in model development and accountability in their deployment becomes imperative. Attention must also be paid to mitigating algo-



Figure 1. Intersection between AI, ML, and Data Science.

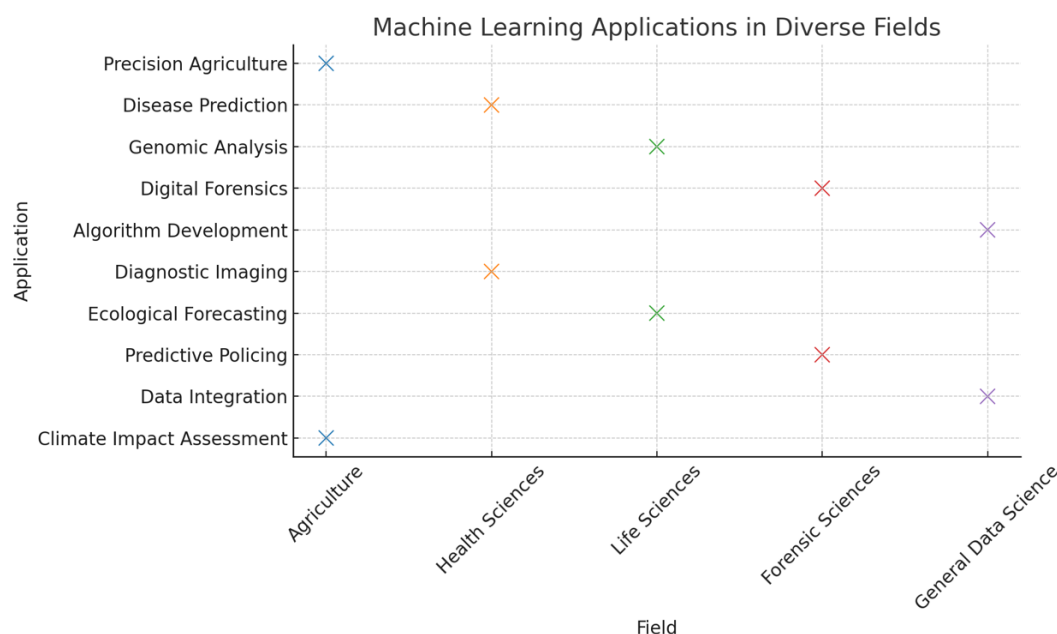


Figure 2. ML applications in diverse fields.

rhythmic bias that could exacerbate health disparities, especially among under-represented or vulnerable populations.

3.4. Innovations in Healthcare Delivery

Innovations include telemedicine, hospital resource optimization, and predictive equipment maintenance (Keesara et al., 2020; Froehle and White, 2014; Hermes et al., 2020). These technologies have become increasingly essential in improving healthcare accessibility, especially in remote or underserved areas. Predictive analytics can be used to anticipate patient surges, manage staffing needs, and reduce wait times, contributing to more efficient and responsive care delivery.

Virtual assistants and AI chatbots improve engagement and adherence by offering patients timely information, medication reminders, and symptom triage (Bickmore et al., 2018). Their integration into clinical workflows can also alleviate administrative burdens on healthcare professionals and enhance continuity of care.

4. Bioinformatics and AI in Life Sciences

This section highlights how computational tools and intelligent algorithms are revolutionizing our understanding of complex biological systems. As life sciences increasingly rely on high-throughput data and real-time measurements, bioinformatics and AI have become indispensable for managing, analyzing, and interpreting these vast datasets. Topics in this section include genomic analysis, proteomics, systems biology, biodiversity monitoring, evolutionary modeling, and ecological forecasting.

Advancements in machine learning (ML) are enabling novel discoveries in functional genomics, disease genetics, and precision agriculture by uncovering hidden patterns within multi-omics datasets. These technologies also support the integration of diverse data types—ranging from gene expression profiles to environmental sensor outputs—into unified biological models. Furthermore, the use of AI in evolutionary and ecological studies allows researchers to make robust predictions about biodiversity, adaptation, and ecosystem dynamics under changing environmental conditions. The convergence of bioinformatics and AI is thus accelerating innovation in areas such as personalized medicine, crop breeding, conservation biology, and synthetic biology, redefining how we

study and interact with living systems.

4.1. Genomic Data Analysis

Massive genomic datasets require ML for identifying variants, understanding genetic diversity, and mapping traits via QTL and GWAS (Deisboeck and Zhang, 2011; Visscher et al., 2017). These approaches facilitate the identification of causal genetic markers linked to complex traits and diseases, enabling advancements in precision medicine and crop improvement. Additionally, integrative analysis of multi-omics data—including transcriptomics, epigenomics, and metabolomics—further enhances the resolution of biological insights and supports predictive modeling of phenotype-genotype associations.

4.2. Proteomics

ML helps identify protein structures, interactions, and functions, accelerating discoveries in drug target identification and biomarker development. Advances in deep learning models, such as AlphaFold, have significantly improved the accuracy of protein structure prediction from amino acid sequences. Integrating proteomics with genomics gives a holistic view of systems biology, enabling researchers to explore regulatory networks, signaling pathways, and functional modules within the cell (Aebersold and Mann, 2016; Wilhelm et al., 2014). This integration is particularly valuable in understanding disease mechanisms and in designing targeted therapeutic interventions.

4.3. Systems Biology

Network analysis and computational modeling elucidate complex biological interactions (Kitano, 2002; Karr et al., 2012). These methods allow researchers to represent biological systems as interconnected networks of genes, proteins, metabolites, and environmental factors. Through simulations and dynamic modeling, systems biology enables the prediction of system behavior under various perturbations, helping to identify potential intervention points for therapeutic or agricultural applications. This integrative approach also facilitates the development of digital twins for cells or organisms, supporting hypothesis testing and personalized interventions in silico.

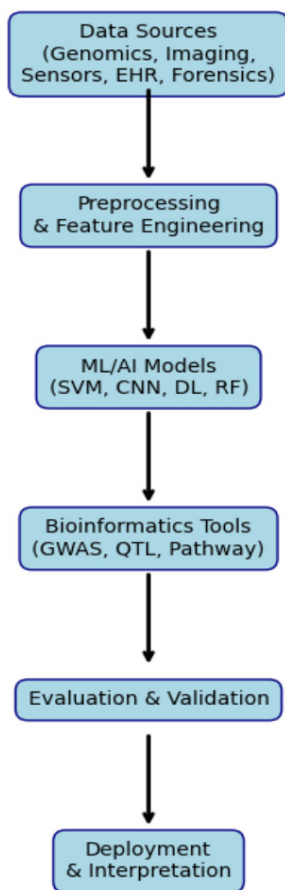


Figure 3. AI/ML and Bioinformatics Workflow in Interdisciplinary Research.

4.4. Biodiversity Conservation

Remote sensing and ML support ecosystem monitoring, species distribution modeling, and conservation planning (Pettorelli et al., 2014; Turner et al., 2015). High-resolution satellite imagery, drone-based sensors, and automated data pipelines provide real-time insights into habitat change, deforestation, and fragmentation. ML models are increasingly used to predict species range shifts under various climate scenarios and to identify biodiversity hotspots requiring urgent protection. These data-driven strategies enhance conservation prioritization and support adaptive management plans that are responsive to environmental dynamics.

4.5. Evolutionary Patterns and Ecological Forecasting

Phylogenetics and ML infer evolutionary relationships and forecast ecological outcomes by analyzing patterns in genomic, ecological, and environmental data (Yang and Rannala, 2012; Clark et al., 2001). These approaches help predict species divergence, adaptive traits, migration pathways, and ecosystem responses to climate change. They are particularly valuable in understanding biodiversity under shifting environmental pressures. Figure 3 illustrates a generalized interdisciplinary workflow showing how diverse datasets—from genomics to remote sensing—can be processed and modeled using AI/ML and bioinformatics tools to support such evolutionary and ecological predictions.

5. AI, ML, and Data Integration in Forensic Sciences

The Journal of Artificial Intelligence, Machine Learning, and Bioinformatics (JAIMLB) will showcase how intelligent systems are transforming forensic science by advancing digital forensics, enhancing pattern recognition, and

enabling integrated analysis of diverse forensic evidence. These technologies improve the accuracy, efficiency, and scope of forensic investigations and contribute to more reliable outcomes in the legal system.

5.1. Advancements in Digital Forensics

AI and ML are increasingly critical in analyzing large, complex datasets generated in digital investigations. These tools are used to detect anomalies, uncover hidden patterns, and accelerate the identification of relevant evidence (Casey, 2011; Beebe, 2009; Chandola et al., 2009). Applications include malware detection, fraud analysis, and cyber intrusion detection. Automated AI-driven tools help streamline the search for digital artifacts, thereby reducing investigative time and human error. Figure 3 also applies to this domain, representing how forensic data—from digital traces to genetic material—can be integrated into unified pipelines for efficient and intelligent evidence processing.

5.2. Pattern Recognition in Forensic Evidence

Deep learning improves fingerprint, DNA, and ballistic analysis by automating complex pattern-matching tasks that traditionally relied on human expertise (Galante and Li, 2023; Richmond et al., 2021; Jain and Feng, 2011; Tai et al., 2014). These systems are capable of extracting and comparing minute features with high accuracy and consistency, enhancing the reliability of forensic identifications. For instance, convolutional neural networks (CNNs) can classify DNA profiles or ballistic markings with reduced false positive rates. Such AI-enabled methods significantly reduce processing time while increasing the scalability of forensic casework.

5.3. New Methodologies for Data Integration in Forensic Cases

Graph databases and blockchain technology enhance the integrity and integration of evidence by allowing investigators to model complex relationships among digital and physical evidence (Stolfo et al., 2000; Angles and Gutierrez, 2008; Kumar et al., 2018). These tools can capture links between suspects, devices, communications, and locations, creating a dynamic and searchable web of forensic information. Blockchain's immutable ledger also supports transparent and tamper-proof chains of custody, ensuring legal admissibility of evidence. Combined with AI/ML, these systems can prioritize leads, detect inconsistencies, and provide investigators with decision support systems that are robust and auditable.

5.4. Ethical and Legal Implications

Ethical frameworks must ensure transparency and fairness in forensic AI to uphold justice and protect civil liberties (Geradts, 2018). As AI tools are adopted in courtrooms and law enforcement, their algorithms must be interpretable, unbiased, and validated for forensic reliability. There is a growing need to standardize practices for the use of AI in legal contexts and to implement oversight mechanisms that mitigate misuse or overreliance on automated outputs. Future JAIMLB articles will critically examine these challenges and promote best practices that align technological innovation with ethical responsibility in forensic science.

6. Core Advances in AI, ML, and Ethical Practice

JAIMLB welcomes foundational research that advances the theoretical underpinnings of Artificial Intelligence (AI), Machine Learning (ML), and Bioinformatics, as well as applied innovations that influence education and ethical implementation. This section aims to foster discourse on the technical progression and responsible adoption of intelligent systems across disciplines.

6.1. Innovative Algorithm Development

Topics include deep learning, reinforcement learning, and quantum computing (LeCun et al., 2015; Sutton and Barto, 2018; Montanaro, 2016). Advances in neural architectures, such as transformers and graph neural networks, have unlocked new applications in natural language processing, molecular modeling, and decision-making systems. Reinforcement learning has shown promise



Figure 4. Ethical AI and Explainable ML Model Flow.

in autonomous systems, robotics, and precision control, while quantum computing opens the door to solving high-dimensional optimization problems that are computationally intractable with classical approaches.

6.2. Advances in ML Techniques

This subsection highlights emerging methods such as transfer learning, generative adversarial networks (GANs), and explainable AI (Pan and Yang, 2010; Doshi-Velez and Kim, 2017). These techniques enhance the adaptability, creativity, and interpretability of AI systems. Transfer learning enables model reuse across domains with limited data, while GANs are instrumental in data augmentation, image synthesis, and drug discovery. Explainable AI is especially critical in healthcare, bioinformatics, and legal contexts, where decisions must be transparent and justifiable.

6.3. Critical Reviews of AI/ML and Bioinformatics Education

Curriculum development and interdisciplinary approaches are essential to prepare the next generation of researchers and practitioners (Baumer, 2015; Hardin et al., 2015). This section invites contributions that evaluate educational strategies, pedagogical innovations, and training programs that integrate AI, ML, and bioinformatics. Topics may include project-based learning, hybrid course delivery, partnerships with industry, and the role of ethics and reproducibility in the classroom. Emphasis will be placed on inclusive learning environments that equip students with both domain-specific and computational competencies.

6.4. Ethical Practices in AI and Data Science

As AI and ML become increasingly embedded in critical decision-making systems, ethical considerations are essential to ensure public trust and social responsibility. Key concerns include data privacy, algorithmic bias, fairness, and transparency in model design and deployment. Figure 4 outlines the foundational principles of ethical AI and ML—such as fairness, accountability, transparency, and human oversight—that guide the responsible development and application of intelligent systems.

Articles in this section will address these challenges in depth, including best practices for data anonymization, informed consent, compliance with data protection regulations, and bias mitigation strategies (Barocas et al., 2019; Tene

and Polonetsky, 2012). Research on explainable AI, diverse training data, and transparency in model architecture will also be prioritized to advance responsible and inclusive innovation.

7. Conclusion

By bridging traditional disciplines with the transformative capabilities of Artificial Intelligence (AI), Machine Learning (ML), and Bioinformatics, the Journal of Artificial Intelligence, Machine Learning, and Bioinformatics (JAIMLB) seeks to foster meaningful collaboration, share emerging knowledge, and drive innovation across diverse scientific domains. Through its interdisciplinary lens, the journal provides a vital platform for the integration of computational intelligence with life, health, agricultural, forensic, and environmental sciences.

JAIMLB's mission is to publish cutting-edge, ethical, and impactful research that not only advances scientific inquiry but also delivers practical solutions to real-world problems. The journal is committed to supporting responsible AI and data-driven discovery, promoting equitable access to technological advancements, and inspiring a new generation of researchers, educators, and practitioners. By amplifying voices across sectors and encouraging rigorous scholarship, JAIMLB aims to be a catalyst for innovation and a forum for interdisciplinary excellence in the era of intelligent science.

Call for Papers

We invite original articles, reviews, and perspectives that reflect on the challenges and opportunities at the intersection of AI, ML, and Bioinformatics across scientific and industrial domains. Visit our Author Guidelines for details.

Acknowledgments

This editorial reflects the collective vision and efforts of the JAIMLB editorial board. We thank our contributors and readers for advancing the mission of interdisciplinary AI, ML, and Bioinformatics research.

Reflect on the Learning Process

Each article in JAIMLB is a catalyst for learning and critical thinking. We encourage our readers to evaluate methods, interpret findings, and consider the broader implications of data-driven research in their fields.

References

- Abrahart RJ and L See (2000) Neural network vs. ARMA modelling: constructing benchmark case studies of river flow prediction. *Hydrological Processes* 14 (11-12): 2107-2124.
- Aebersold R and M Mann (2016) Mass-spectrometric exploration of proteome structure and function. *Nature* 537 (7620): 347-355.
- Angles R and C Gutierrez (2008). Survey of graph database models. *ACM Computing Surveys (CSUR)* 40 (1): 1-39.
- Barocas S, M Hardt, and A Narayanan (2019) Fairness and machine learning. Accessed June 20, 2024 at <https://www.fairmlbook.org>.
- Baumer B (2015) A data science course for undergraduates: Thinking with data. *The American Statistician* 69 (4): 334-342.
- Beebe NL (2009) Digital Forensic Research: The Good, the Bad and the Unaddressed. In: Peterson, G., Shenoi, S. (eds) *Advances in Digital Forensics V. Digital Forensics 2009*. IFIP Advances in Information and Communication Technology, vol 306. Springer, Berlin, Heidelberg. https://doi.org/10.1007/978-3-642-04155-6_2.
- Bickmore TW, LM Pfeifer, and MK Paasche-Orlow (2018) Using computer agents to explain medical documents to patients with low health literacy. *Patient Education and Counseling*, 101 (8): 1420-1427.
- Casey E (2011) *Digital Evidence and Computer Crime: Forensic Science, Computers, and the Internet*. ISBN 978-0-12-374268-1. Academic Press & Elsevier.
- Chandola V, A Banerjee, and V Kumar (2009) Anomaly detection: A survey. *ACM Computing Surveys (CSUR)* 41 (3): 1-58.
- Clark JS, SR Carpenter, M Barber, S Collins, A Dobson, JA Foley, DM Lodge, M Pascual, R Pielke, W Pizer, C Pringle, WV Reid, KA Rose, O Sala, WH Schlesinger, DH Wall, and D Wear (2001) Ecological forecasts: an emerging imperative. *Science* 293 (5530): 657-660. <https://doi.org/10.1126/science.293.5530.657>.
- Collins FS and H Varmus (2015) A new initiative on precision medicine. *New England Journal of Medicine* 372 (9): 793-795.
- Deisboeck TS, and L Zhang (2011) Multiscale modeling of cancer: computational approaches from the nanoscale to the whole organism. *Wiley Interdisciplinary Reviews: Systems Biology and Medicine* 3 (4): 323-336.
- Doshi-Velez F, and B Kim (2017) Towards a rigorous science of interpretable machine learning. *arXiv preprint arXiv:1702.08608*. <https://doi.org/10.48550/arXiv.1702.08608>.
- Esteva A, B Kuprel, RA Novoa, J Ko, SM Swetter, HM Blau, and S Thrun (2017) Dermatologist-level

- classification of skin cancer with deep neural networks. *Nature* 542 (7639): 115-118. <https://doi.org/10.1038/nature21056>.
- FAO (2017) The future of food and agriculture: Trends and challenges. Food and Agriculture Organization of the United Nations.
- Foley JA, N Ramankutty, KA Brauman, ES Cassidy, JS Gerber, M Johnston, ND Mueller, C O'Connell, DK Ray, PC West, C Balzer, EM Bennett, SR Carpenter, J Hill, C Monfreda, S Polasky, J Rockström, J Sheehan, S Siebert, D Tilman, and DPM Zaks (2011) Solutions for a cultivated planet. *Nature* 478 (7369): 337-342. <https://doi.org/10.1038/nature10452>.
- Froehle CM and JA White (2014) Interruption and Forgetting in Knowledge-Intensive Service Environments. *POMS*. <https://doi.org/10.1111/poms.12089>.
- Galante N, R Cotroneo, D Furci, R Cotroneo, D Furci, G Lodetti, and MB Casali (2023) Applications of artificial intelligence in forensic sciences: Current potential benefits, limitations and perspectives. *Int J Legal Med* 137: 445-458. <https://doi.org/10.1007/s00414-022-02928-5>.
- Gerads Z (2018) Digital, big data and computational forensics. *Forensic Sciences Research* 3 (3): 179-182. <https://doi.org/10.1080/20961790.2018.1500078>.
- Hardin J, R Hoerl, NJ Horton, D Nolan, B Baumer, O Hall-Holt, and MD Ward (2015) Data science in statistics curricula: Preparing students to "think with data". *The American Statistician*, 69(4), 343-353.
- Hermes S, T Riasanow, EK Clemons, M Böhm, and H Krcmar (2020) The digital transformation of the healthcare industry: exploring the rise of emerging platform ecosystems and their influence on the role of patients. *Business Research* 13: 1033-1069. <https://doi.org/10.1007/s40685-020-00125-x>.
- Hood L and M Flores (2012). A personal view on systems medicine and the emergence of proactive P4 medicine: predictive, preventive, personalized and participatory. *New Biotechnology* 29(6): 613-624. <https://doi.org/10.1016/j.nbt.2012.03.004>.
- Huang J, C Pray, and S Rozelle (2002) Enhancing the crops to feed the poor. *Nature* 418: 678-684. <https://doi.org/10.1038/nature01015>.
- Jain AK and J Feng (2011) Latent fingerprint matching. *IEEE Transactions on Pattern Analysis and Machine Intelligence* 33 (1): 88-100.
- Kamilaris A, A Kartakoullis, and FX Prenafeta-Boldú (2018) A review on the practice of big data analysis in agriculture. *Computers and Electronics in Agriculture* 143: 23-37.
- Karr JR, JC Sanghvi, DN Macklin, MV Gutschow, JM Jacobs, B Bolival, N Assad-Garcia, JI Glass, and Covert, M. W. (2012). A whole-cell computational model predicts phenotype from genotype. *Cell* 150 (2): 389-401.
- Kesara S, A Jonas, and K Schulman (2020) Covid-19 and health care's digital revolution. *New England Journal of Medicine* 382 (23), e82. <https://doi.org/10.1056/NEJMp20058>.
- Kitano H (2002) Computational systems biology. *Nature* 420: 206-210. <https://doi.org/10.1038/nature01254>.
- Kumar R, M Luthra, and RK Kesharwani (2018) Blockchain criminology: An analysis of blockchain technology in the domain of criminal investigation. In *Advances in Smart Grid and Renewable Energy* (pp. 317-324). Springer.
- LeCun Y, Y Bengio, and G Hinton (2015) Deep learning. *Nature* 521 (7553): 436-444. <https://doi.org/10.1038/nature14539>.
- Litjens G, T Kooi, BE Bejnordi, AA Setio, F Ciompi, M Ghafoorian, JAWM van der Laak, B van Ginneken, and CI Sánchez (2017) A survey on deep learning in medical image analysis. *Medical Image Analysis* 42: 60-88. <https://doi.org/10.1016/j.media.2017.07.005>.
- Lobell DB, W Schlenker, and J Costa-Roberts (2011) Climate trends and global crop production since 1980. *Science* 333 (6042): 616-620. <https://doi.org/10.1126/science.1204531>.
- Marx V (2013) Biology: The big challenges of big data. *Nature* 498 (7453): 255-260. <https://doi.org/10.1038/498255a>.
- Montanaro A (2016) Quantum algorithms: An overview. *npj Quantum Inf* 2, 15023. <https://doi.org/10.1038/npjqi.2015.23>.
- Mulla DJ (2013) Twenty-five years of remote sensing in precision agriculture: Key advances and remaining knowledge gaps. *Biosystems Engineering* 114 (4): 358-371. <https://doi.org/10.1016/j.biosystemseng.2012.08.009>.
- Obermeyer Z and EJ Emanuel (2016) Predicting the future—big data, machine learning, and clinical medicine. *New England Journal of Medicine* 375 (13): 1216-1219.
- Obermeyer Z, B Powers, C Vogeli, and S Mullainathan (2019) Dissecting racial bias in an algorithm used to manage the health of populations. *Science* 366 (6464): 447-453.
- Pan SJ and Q Yang (2010) A survey on transfer learning. *IEEE Transactions on Knowledge and Data Engineering* 22 (10): 1345-1359.
- Pettorelli N, WF Laurance, TG O'Brien, M Wegmann, H Nagendra, and W Turner (2014) Satellite remote sensing for applied ecologists: opportunities and challenges. *Journal of Applied Ecology* 51 (4): 839-848.
- Raza A, A Razzaq, SS Mehmood, X Zou, X Zhang, Y Lv, and J Xu (2019) Impact of Climate Change on Crops Adaptation and Strategies to Tackle Its Outcome: A Review. *Plants* 2019, 8, 34; <https://doi.org/10.3390/plants8020034>.
- Richmond KM (2021) AI, Machine Learning, and International Criminal Investigations: The Lessons From Forensic Science. *iCourts Working Paper Series*, No. 222, Available at SSRN: <https://ssrn.com/abstract=3727899> Or <http://dx.doi.org/10.2139/ssrn.3727899>.
- Roden DM, HL McLeod, MV Williams, GA Mensah, JF Peterson, and SL Van Driest (2011) Pharmacogenomics. *The Lancet* 380 (9852): 507-520.
- Safran C, M Bloomrosen, WE Hammond, S Labkoff, S Markel-Fox, PC Tang, and DE Detmer (2007) Toward a national framework for the secondary use of health data: an American Medical Informatics Association white paper. *Journal of the American Medical Informatics Association* 14 (1): 1-9.
- Schlenker W and MJ Roberts (2009) Nonlinear temperature effects indicate severe damages to US crop yields under climate change. *PNAS* 106 (37): 15594-15598.
- Shilo S, H Rossman, and E Segal (2020) Axes of a revolution: challenges and promises of big data in healthcare. *Nature Medicine* 26 (1): 29-38.
- Stolfo SJ, W Fan, W Lee, A Prodromidis, and PK Chan (2000) Cost-based modeling for fraud and intrusion detection: Results from the JAM project. *DARPA Information Survivability Conference and Exposition* 2: 130-144.
- Sutton RS and AG Barto (2018) Reinforcement Learning: An Introduction. MIT press.
- Tai LH, U Kruger, and EJ Jacobs (2014) Comparison of gunshot residue distribution patterns on hands and associated clothing of suicides and homicides. *Forensic Science International* 239: 7-14.
- Tene O and J Polonetsky (2012) Privacy in the age of big data: A time for big decisions. *Stanford Law Review Online*, 64, 63.
- Topol EJ (2019) High-performance medicine: the convergence of human and artificial intelligence. *Nature Medicine* 25 (1): 44-56.
- Turner W, S Spector, N Gardiner, M Fladeland, E Sterling, and M Steininger (2015) Remote sensing for biodiversity science and conservation. *Trends in Ecology & Evolution* 28 (5): 334-343.
- Visscher PM, NR Wray, Q Zhang, P Sklar, MI McCarthy, MA Brown, and J Yang. (2017) 10 years of GWAS discovery: biology, function, and translation. *American Journal of Human Genetics* 101 (1): 5-22.
- Wang L, N Pedroni, and E Zio (2018) Integrating health data into predictive maintenance of industrial systems: from failure-time distribution estimation to degradation analysis. *IEEE Transactions on Reliability* 67 (1): 283-294.
- Webber H, T Gaiser, and F Ewert (2014) What role can crop models play in supporting climate change adaptation decisions to enhance food security in Sub-Saharan Africa? *Agricultural Systems* 127: 161-177. <https://doi.org/10.1016/j.agsy.2013.12.006>.
- Westbrook JK, JD Lopez, SJ Fleischer, PD Lingren, and RL Meagher (2016) Modeling seasonal migration of fall armyworm moths. *Int J of Biometeorology* 60 (2): 255-267.
- Wilhelm M, J Schlegl, H Hahne, A Moghaddas Gholami, M Lieberenz, MM Savitski, E Ziegler, L Butzmann, S Gessulat, H Marx, T Mathieson, S Lemeier, K Schnatbaum, U Reimer, H Wenschuh, M Mollenhauer, J Slotta-Huspenina, JH Boese, M Bantscheff, A Gerstmair, F Faerber, and B Kuster (2014) Mass-spectrometry-based draft of the human proteome. *Nature* 509 (7502): 582-587. <https://doi.org/10.1038/nature13319>.
- Yang Z and B Rannala (2012) Molecular phylogenetics: principles and practice. *Nature Reviews Genetics* 13 (5): 303-314.
- Zhang C, D Walters, and JM Kovacs (2019) Applications of low altitude remote sensing in agriculture upon vertical take-off and landing (VTOL) systems: A review. *Remote Sensing* 11 (3): 320.