Artificial Intelligence in Musculoskeletal Medicine: A Scoping Review

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n recent years, artificial intelligence (AI) has blossomed via the booming increase in computing power, access to large data sets and the development of novel data science techniques and architectures. This has led to a great interest in the clinical deployment of AI in healthcare. In this review, we look particularly at the field of musculoskeletal medicine and have curated the latest progress in AI and machine learning in the areas of inflammatory arthritis, vasculitis and connective tissue diseases, osteoporosis, Ehlers–Danlos syndrome, osteoarthritis and musculoskeletal surgical interventions. We also highlight the challenges and pitfalls as clinical AI moves forward.

Keywords

Arthritis, artificial intelligence, inflammatory, machine learning, musculoskeletal, osteoporosis, rheumatology

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There is much excitement about the deployment of artificial intelligence (AI) in healthcare, and the musculoskeletal field is no exception. In this article, we introduce some of the latest developments relating to osteoarthritis (OA), osteoporosis, rheumatoid arthritis (RA) (as an example of inflammatory arthritis), connective tissue disease (CTD), Ehlers–Danlos syndrome (EDS) and musculoskeletal surgical interventions.

Artificial intelligence

Al is the ability of machines to imitate human behaviour. Machine learning (ML) is a branch of Al that uses computer algorithms to automatically learn from input data and through the identification of patterns to make accurate predictions with minimal human intervention. ML algorithms do not rely on a predetermined equation as a model but adaptively improve their performance as more data are provided.

ML is increasingly being used in medical applications to independently and efficiently carry out specified tasks, such as image processing in radiology and histopathology and the identification of disease markers, and in the field of drug discovery.

The process by which ML algorithms learn is called 'training', and the training methods are classified as supervised, unsupervised or reinforcement learning. Supervised learning involves training algorithms to predict future values through learning patterns from labelled input data, for example, accurately predicting the opacification on a chest X-ray through training with images that are labelled as normal or abnormal.

Supervised learning algorithms use classification, regression and forecasting approaches. In classification models, the algorithm must draw a conclusion from the observed values and determine to which category new observations belong. In regression tasks, the ML algorithm must estimate and understand the relationships among variables. Forecasting is the process of making predictions about the future based on past and present data and is commonly used to analyse trends.

In unsupervised learning, the input data are not assigned or labelled. Instead, the algorithm infers the underlying patterns and relationships within the input data, which is useful, for example, in finding patterns in blood biomarkers that are associated with a particular disease state. Clustering is a type of unsupervised learning that involves grouping sets of similar data based on the defined criteria. It is useful for segmenting the data into groups and performing analysis on each data set to find patterns. Dimension reduction is another type of unsupervised learning that gradually reduces the number of variables being considered to find the exact information required. In reinforcement learning, the algorithm achieves a specified goal by interacting with an environment through trial

Figure 1: Schematic representations of machine learning and deep learning methods



predictions based on how a previous decisions were made.

LASSO = least absolute shrinkage and selection operator.

and error. This type of method is commonly applied in the field of robotic surgery.

Examples of ML algorithms are shown in *Figure 1* and described in greater depth in the glossary (online supplementary file 1).

Inflammatory arthritis

Inflammatory arthritides are characterized by joint inflammation, leading to clinical signs, such as pain, swelling, heat and erythema. This collection of rheumatological conditions encompasses those causing inflammation due to various aetiologies, including autoimmunity, crystals and reactive processes. In this review, the focus will be on RA as a vital example of inflammatory arthritis.

RA is a chronic, systemic autoimmune disease leading to inflammatory arthritis, typically involving the small joints that can result in joint destruction and severe functional impairment.¹ RA is a heterogeneous disease with a wide variation in the age of onset, degree of joint involvement and severity. The diagnosis is based on clinical criteria, and there is no definitive diagnostic test available to differentiate RA from other inflammatory joint diseases.² The aim of treatment for RA is to reduce inflammation and prevent joint destruction, and it includes corticosteroids, conventional disease-modifying anti-rheumatic drugs (DMARDs) and an ever-expanding list of targeted biological therapies.³ Patients have variable response and remission rates to different treatment strategies, and identifying responders from non-responders is difficult. However, early diagnosis and prompt initiation of treatment are important in the prevention of inflammatory joint destruction. AI and ML are increasingly being used to address the unmet needs in the diagnosis, monitoring and treatment of RA, and in this section, we review the latest developments.⁴

Machine learning in inflammatory arthritis diagnosis

AI and ML for diagnostic purposes use the data from electronic health records (EHRs), diagnostic biomarkers and imaging modalities to identify patients with RA at an early stage of the disease.

Electronic health records

The phenotypic identification of RA using data from EHRs was first carried out in 2010 using a support vector machine (SVM) model, which accurately predicted RA disease activity from the data containing specific terms (area under the curve [AUC] >0.90).⁵ The comparison of different ML methods using a 10-fold cross-validation from thousands of EHRs showed that an SVM model performed best in identifying patients with RA in <7 s (an AUC of 0.98 and a positive predictive value [PPV] of 0.94).⁶ The model compared favourably with manual methods.⁷ A random forest method and a decision tree model identified the best predictors of RA, such as local coding for RA; clinical features, such as nodules, seropositivity and arthropathy and drugs, such as prednisolone, methotrexate, sulfasalazine and leflunomide, from EHRs, with a PPV of 85.6%, specificity of 94.6% and sensitivity of 86.2%.⁸

Molecular markers

Random forest, logistic regression and other statistical models have differentiated patients with RA from controls using gene expression signatures, with an AUC of 0.96.^{9,10} Artificial neural networks (ANNs) have successfully distinguished RA from OA and psoriatic arthritis (PsA, another form of inflammatory arthritis) using serum protein profiles identified through mass spectrometry and immunoassays, although the AUC varies significantly.^{11–13}

Imaging techniques and histopathology

Supervised and unsupervised algorithms have been used, together with imaging techniques, for RA diagnosis. ANN models trained and tested on hand radiographs accurately classified RA with an AUC of 0.97, a sensitivity of 90.7% and a specificity of 92.6%.^{14,15} Larger data sets have allowed the effective differentiation of RA and OA.¹⁶ ANNs also identified synovial proliferation in joint ultrasound with an AUC of 0.863–0.886, depending on the joint area sampled.^{17,18} A systemic review and meta-analysis of computer-assisted magnetic resonance imaging (MRI) showed a good correlation with manual methods in RA diagnosis.¹⁹ Thermal imaging of diurnal joint temperatures has been studied, with quantum SVM models used to improve the accuracy.^{20,21} Scoring algorithms based on features, such as inflammatory infiltrates and hyperplasia in stained synovial tissue, have demonstrated considerable efficacy in differentiating RA, OA and PsA from normal tissue (AUC 0.87–0.98).^{22,23}

Disease progression and disease activity

Once RA is diagnosed, an ongoing assessment is vital to monitor the disease activity. An ordinal regression-based ML model developed to estimate the Clinical Disease Activity Index (CDAI) scores was trained and validated on 11,985 patient notes with CDAI scores from the OM1 RA registry (OM1[®] real-world data and AI for research [OM1, Boston, MA, USA).²⁴ The model had a PPV of 0.80, a negative predictive value of 0.84 and an AUC of 0.88 when evaluating the performance using the binarized (negative CDAI score ≤10 or positive CDAI score ≥10.1) version of the outcome.

Predicting treatment response

ML techniques have shown promise in the evaluation of treatment response to DMARDs and biologics. Least absolute shrinkage and selection operator and random forest methods showed that patients with RA with a low baseline Disease Activity Score in 28 joints (DAS28)erythrocyte sedimentation rate (ESR) score, positive anti-citrullinated protein antibody and the Health Assessment Questionnaire ≤2 were most likely to respond well to methotrexate (AUC of 0.68 and 0.79, respectively).²⁵ Conversely, a DAS28-ESR score (>3.2) at 3 months postmethotrexate treatment predicted a poor therapeutic response with different methods.²⁶ Elastic net models demonstrated a 3.6-fold better response to anti-tumour necrosis factor (TNF) biologics in males, and using genetic data, carriers of the resistin gene rs3219177 polymorphism had a sixfold better response than non-carriers. In a subsequent study, carriers of the TLR9 gene rs352139 polymorphism were found to have a fivefold better response to anti-TNF than non-carriers.^{27,28} ML using biomarker signatures in blood or synovial tissue has reliably predicted responses to other biologics (e.g. rituximab, tocilizumab and sarilumab).^{29–31}

Vasculitis and connective tissue disease

Vasculitis is a complex group of rare rheumatic diseases characterized by inflammation of blood vessels that present considerable diagnostic and therapeutic challenges. Recent studies using AI techniques have attempted to support earlier diagnosis, treatment optimization and prognosis prediction.³²⁻³⁴

The diagnosis of vasculitis relies upon the recognition of characteristic clinical features often alongside the use of imaging or biopsy. ML techniques show great promise for supporting this process, with studies using natural language processing to differentiate vasculitis from clinical notes, deep learning algorithms (such as *U-Net*) to detect features

on medical imaging and supervised learning approaches to identify predictive biomarkers detectable from blood samples. $^{\rm 35\text{-}41}$

Treatment typically includes induction through high-dose corticosteroids followed by maintenance with 'steroid-sparing' agents. However, variability in treatment response combined with common side effects makes management fraught with difficulty. ML has been used to address treatment challenges through supervised algorithms (including *random forest* and *light gradient boosting models*) to predict relapse after glucocorticoid tapering in giant cell arteritis and the development of models predicting resistance to intravenous immunoglobulin therapy in Kawasaki's disease.^{32,42,43}

The risk of morbidity following a diagnosis of vasculitis is significant, and understanding the prognostic risk factors is essential to tailor management strategies. Emerging ML-based tools show promise, especially in predicting eye diseases from clinical data (using ensemble algorithms, such as *XGBoost*) or angiography imaging and risk of renal damage in immunoglobulin A (IgA) vasculitis based on patients' clinical features.^{44–46}

With only 26 primary research studies published over the last 20 years using AI methods in vasculitis, and the majority investigating Kawasaki's disease (n=12), there is still a dearth of available evidence despite many studies showing highly promising results.^{47–57}

Autoimmune CTDs are systemic in nature and characterized by a variety of phenotypes. ML techniques have begun to revolutionize the study of these varied diseases, providing tools for improving diagnostic approaches, the classification of disease subtypes and predicting factors associated with treatment response and the development of disease-related complications.

CTDs present a diagnostic challenge for clinicians, and many studies using ML focus on demonstrating accurate tools to support the diagnostic process. There are encouraging studies investigating gene expression data as novel biomarkers in CTDs, but perhaps, the approaches closer to clinical implementation are those using deep learning to support and automate the interpretation of clinical data (including point-of-care tests, radiological imaging and biopsies).^{58–71} An interesting example includes the use of deep learning to differentiate systemic sclerosis from primary Raynaud's syndrome using a low-cost digital nailfold capillaroscope, an underused tool in which AI may support more widespread use.⁶³

The spectrum of CTD presentations highlights the importance of classification, subtype definition and tailored treatment strategies. Data mining and supervised approaches have been used to discriminate subtypes of Sjögren's syndrome and lupus based on clinical data, in addition to multiple studies investigating the classification of myositis using electromyography and MRI data.^{72–76}

Many of the Al-based studies on CTDs aim to predict outcomes and treatment responses to inform a personalized management approach. Multiple ML approaches have been used to predict complications and comorbidities in lupus (including nephritis), pulmonary involvement in systemic sclerosis and response to immunomodulatory treatments.^{77–97}

Osteoporosis

Osteoporosis is characterized by reduced bone mineral density (BMD), leading to a predisposition to a fragility fracture. The diagnosis is complicated by the absence of symptoms or clinical signs and is reliant

on measures of BMD. BMD is primarily derived from dual-energy X-ray absorptiometry (DXA) as a single parameter, and it is this measurement that informs the densitometric diagnosis of osteoporosis, defined as a BMD *T*-score of \leq -2.5. Al techniques have been used to interrogate 'omic' data sets to identify biomarkers and therapeutic targets in osteoporosis, as they have in most major disease areas; however, the focus of this review will be on the deployment of Al on skeletal imaging (computer vision) in the field of osteoporosis.^{98,99}

The major aims of computer vision in osteoporosis are as follows:

- use DXA images to improve fracture prediction beyond BMD
- opportunistically measure BMD from routinely performed imaging
- opportunistically identify vertebral fractures from routinely
 performed imaging
- derive bone microarchitecture features.

It is worth noting that, in the field of osteoporosis, the outcome of interest can be either the diagnosis of osteoporosis or the assessment of fracture risk. Although the former was vital for defining the disease and developing the field, the measures of the latter (fracture risk) are used to adjudicate treatment thresholds via fracture risk-prediction tools, including The Fracture Risk Calculator (FRAX[®]), which provides a 10-year estimation of major osteoporotic risk and hip fracture risk.¹⁰⁰ This is, in part at least, as over half of fragility fractures occur at a normal BMD and so a pure focus on the diagnosis of osteoporosis would miss a large proportion of those at risk of fracture. Indeed, fractures are associated with high morbidity and mortality, surpassing simply the densitometric diagnosis of osteoporosis.

As mentioned earlier, the DXA scan is performed to provide a measure of BMD (via levels of beam attenuation), but the rest of the DXA image (of the hips and anterior–posterior spine) is neglected unless a lateral image of the spine is acquired as a vertebral fracture assessment. These acquired, but underused, DXA images are ripe for computer vision analyses.

Indeed, a meta-analysis of seven studies that attempt to improve the diagnosis of osteoporosis via hip DXA images found a pooled sensitivity of 0.844 (95% confidence interval [CI], 0.79–0.89), a pooled specificity of 0.781 (95% CI, 0.73–0.82) and a summary of receiver operating characteristic AUC of 0.878.¹⁰¹ Although these results are encouraging, DXA imaging is limited by quality due to artefacts of soft tissue and degenerative sclerosis and, in a clinical setting, is only performed to assess BMD. Plain radiographs and computed tomography (CT) are imaging modalities that are used for a plethora of clinical indications and often contain skeletal elements. These images could all contribute to the assessment of skeletal health opportunistically.

This has led to studies showing that abdominal CT can predict lumbar spine BMD (AUC 0.96–0.97), chest CT scans (performed for lung cancer screening) can discriminate osteoporosis from normal participants (AUC 0.97) and lateral spine X-rays can identify osteoporosis, though with a lower degree of accuracy (AUC 0.85).^{102,103}

Vertebral fractures are often missed on routine imaging, and hence, models have been developed using deep learning (amidst other methods) to identify vertebral fractures from lateral spine radiographs (AUC 0.93) and sagittal reconstructions of CT images.^{104,105}

Bone is composed of two compartments, cortical and trabecular, and information regarding these compartments can provide more detailed 'bone microarchitectural' measures of bone strength and resistance to fracture than simply BMD alone. This microarchitecture can be derived from specialist scans currently largely used in research settings, including high-resolution peripheral quantitative CT (HR-pQCT). Computer vision analyses of HR-pQCT have demonstrated excellent discrimination for prevalent fractures with an AUC of 0.85–0.97 using a gradient boosting machine and statistical shape mapping.^{106–108} What is more, there are efforts, using AI, to derive some microarchitectural features (though only from the trabecular compartment) from the DXA imaging.¹⁰⁹

In conclusion, computer vision analyses have improved the identification of fracture risk on DXA images, developed the opportunistic screening of BMD and vertebral fractures and shown promise for deriving bone microarchitectural properties from clinically applicable imaging modalities.

Ehlers–Danlos syndrome

EDSs are a group of inherited disorders of connective tissue. Within this group, there are 14 recognized EDS subtypes, 13 of which are genetically characterized. The clinical presentation commonly includes features such as joint hypermobility, skin hyperextensibility, abnormal wound healing and widespread pain. Similar to connective tissue disorders, EDSs are not solely disorders of the musculoskeletal system but are complex multi-system disorders due to the role of connective tissue throughout the body. Al tools in development may eventually assist in the diagnosis of EDS, with two current examples being video assessment of joint hypermobility and the classification of rare diseases via analysis of patient-created pain drawings.

A thematic clinical feature of EDS is generalized joint hypermobility. The clinical finding of joint hypermobility is a core diagnostic feature measured through the Beighton examination completed using a goniometer. However, the use of a goniometer is uncommon in actual clinical practice, and evidence of inter-rater reliability in the Beighton score reporting is limited or conflicting.¹¹⁰ Mittal et al. have devised a method to apply computer vision to smartphone camera footage of patients, with the goal of creating a more objective, validated and scalable method for assessing hypermobility.¹¹¹ This provides an excellent case study to highlight the design of research in the field of musculoskeletal AI.

The footage will be analysed using ML algorithms from the existing opensource human pose-estimation models, which use the video input to create an output of the location and angles of joints in the body. Poseestimation models lack training data on generalized joint hypermobility, potentially creating *out-of-distribution errors* for the pose libraries. The authors plan to address this by additional fine-tuning of the models with manually annotated examples of hyperextended joints. The method is planned to be validated compared with the ground truth of expert clinicians using the goniometer-validated assessment.

The outcomes of the pre-registered trial are awaited in 2024 (Assessing the Feasibility of a Smartphone-based, Machine Learning Visual Imaging Application for Assessment of Hyperextensibility of Peripheral Joints in Ehlers Danlos Syndrome; ClinicalTrials.gov identifier: NCT05366114), and the authors anticipate possible use cases as a triage tool for specialized EDS clinic referrals.¹¹²

Chronic pain is another thematic clinical feature of EDS. There are many aetiologies of chronic pain, including other rare diseases, and

management is informed by the underlying cause. Emmert et al. developed an algorithm trained to distinguish between patterns of chronic pain caused by different rare diseases.¹¹³ The input data to the algorithm were pain drawings. Pain drawings are made by patients who mark where they experience pain on a 2D outline of a human body to create a visual representation of their pain (instead of relying on language). The authors used an open-source platform 'Pain2D' to collect the data from patients with specific known rare disease diagnoses involving chronic pain: EDS, Guillain-Barré syndrome, facioscapulohumeral muscular dystrophy, proximal myotonic myopathy and a control group with nonspecific chronic pain. An overall pain profile for each disease category was created using this platform by calculating the frequency with which each pixel in the pain drawing was marked over all instances of each diagnostic category. Pain profiles summarizing different diagnostic categories were then compared with the known cases (leave-one-out cross-validation) by a mathematical coefficient of similarity - the highestscoring pain profile was chosen as the output classification. This method classified EDS among the five diagnostic categories with a sensitivity of 64.4% and a specificity of 88.7%. Significant limitations include the absence of pain-drawing data from other diagnostic categories of chronic pain and small data sets within the existing categories that precluded the use of deep learning by convolutional neural networks (CNNs).

It is anticipated that these study areas will be developed and built upon but provide good examples of AI study designs that have the potential to be used across musculoskeletal disorders.

Osteoarthritis

OA is a common joint disease affecting millions of people throughout the world but is plagued by a disconnect between clinical and radiographic diseases. In recent years, ML and AI techniques have shown promise for enhancing OA diagnosis, care and management.¹¹⁴ This section seeks to give readers an overview of the most recent developments in ML and AI applications in the field of OA.

Early diagnosis and disease progression detection

Early diagnosis and detection algorithms use a variety of data sources, including patient-reported outcomes and medical imaging (X-rays and MRI scans), to detect the patterns and signals linked to the disease. This can take different approaches, but one study combined clinical, imaging and demographic features using a *multi-modal feature integration method* using an L-1 normalization approach to minimize the number of irrelevant features from each modality.¹¹⁵ Early OA symptoms can be recognized so that therapies can start sooner, possibly slowing the disease's progression.

To develop predictive models for OA progression, ML algorithms have been trained on big data sets combining patient demographics, clinical data and imaging results. These models determine the chance that an illness will progress and assist physicians in selecting the best course of action, potentially improving patient outcomes via a clinical decision support system. One research presented a CNN model referred to as the *YOLOV3* tiny-based object identification approach to automatically detect and categorize knee OA according to the Kellgren–Lawrence (KL) classification scheme.¹¹⁶The strategy used two models, one for classifying severity (progression) and the other for distinguishing between healthy and osteoarthritic knees. The CNN model sought to assist radiologists and orthopaedic surgeons by correctly identifying and classifying knee OA across various stages. Another research introduced *DenseNet169* as a model for the effective diagnosis of knee OA.¹¹⁷ It compared *DenseNet169*'s performance with five other deep learning systems and highlighted its efficacy in finding knee OA and determining its severity through multi-classification (accuracy of 95%, sensitivity of 88.77%, specificity of 95.41% and precision of 87.08%) and binary classifications (accuracy of 93.78%, sensitivity of 91.29%, specificity of 87.57% and accuracy of 89.27%). The suggested methodology showed potential in relieving radiologists' burden, enabling early diagnosis and saving time and expense related to knee OA diagnosis. It used a weighted training approach and incorporated gradient-weighted class activation mapping, a heatmap-based algorithm that deep learning models used to visualize and decipher the significant areas of an image (using gradients in the final convolutional layer) that influence the model's prediction.

Treatment plans for people with OA have been improved using ML and AI techniques. Algorithms can recommend personalized treatment programmes that are suited to each patient's needs by examining patient data, including medical history, genetic factors and lifestyle variables. By identifying the most efficient interventions and minimizing the process of trial and error while choosing treatments, this strategy seeks to improve patient outcomes.

One study found that AI-aided clinician diagnostic ratings have a stronger correlation with the total KL scores and Knee Injury and Osteoarthritis Outcome Scores than those by unaided physicians, increasing the consistency and precision of OA diagnosis.¹¹⁸ The KL score appears to be a limited tool on its own, with only a modest relationship between clinical severity and radiographic assessments. AI assistance has the potential to address the disconnect between radiological OA severity and clinical symptoms, resulting in more precise detection and better patient care.

The use of wearable and sensor technologies in conjunction with ML algorithms enables remote monitoring of patients with OA. These devices can monitor joint motion, levels of activity and other important factors. Clinicians can provide real-time feedback and useful insights into patient situations by analysing the acquired data, enabling proactive management and intervention. For monitoring human movement, wearable sensors provide an affordable and practical alternative to optical motion capture.¹¹⁹ Large amounts of real-world data can be provided by them, enabling remote monitoring, resolving time restrictions in the delivery of remote treatment and encouraging patient interaction. For movement analysis, accelerometers, gyroscopes and magnetometers are frequently combined into *inertial measurement units* for cost-efficiency and usability in large cohorts. For people with hip or knee OA, including those who have had joint replacement surgery, wearable sensors hold great promise for improving research and care.

Another study validated a remote patient monitoring system for 25 patients with OA undergoing total knee arthroplasty using a wearable knee sleeve and smartphone applications.¹²⁰ The sleeve accurately measured mobility, range of motion, patient-reported outcome measures, opioid compliance and home exercise compliance. Such studies demonstrate the potential of AI-augmented remote patient monitoring for evaluating hip and knee arthroplasty recovery and rehabilitation.

In OA management, persistent challenges remain around the quality and accessibility of data. Obtaining comprehensive data sets spanning diverse patient profiles, multiple data modalities and longitudinal records remains a hurdle, essential for training reliable ML models. Another critical challenge lies in the interpretability and explainability of these models; understanding the rationale behind their diagnoses or treatment suggestions is crucial for building trust among healthcare professionals and patients. Additionally, ensuring the ethical use of sensitive medical information while maintaining patient privacy poses an ongoing concern. Moreover, integrating these advanced technologies safely into clinical workflows and healthcare systems presents a significant challenge for wider adoption and practical implementation in real-world settings.

Musculoskeletal surgical interventions

The sphere of surgical intervention has notably benefitted from the introduction of AI-aided technologies, which present the potential to amplify several aspects of surgical care.^{121–128} The purview of these tools encompasses diagnosis and planning to intraoperative guidance.^{121–128} A salient advantage rests in their capacity to aid clinicians in the early detection of musculoskeletal disorders, such as OA and soft tissue pathologies.^{123,127,129,130} Furthermore, the competency of AI in the precise identification of fractures and distinguishing between diverse types of spondylitis parallels that of experienced radiologists potentially enabling faster management.^{131–133}

In the planning phase, AI resources are able to deliver patient-specific preoperative plans, thereby minimizing the necessity for surgeon-delivered modifications to the surgical plan.^{121,123,134} These resources can also construct 3D models of patient anatomy, providing orthopaedic surgeons with a holistic and detailed visual representation of the surgical area.^{124,125} In the operating theatre, AI's role expands to offering real-time feedback and assistance in navigating complex anatomical structures.^{124,126,135,136}

Postoperative care has similarly been affected by the advent of AI, aiding in the identification of implant positioning and type, forecasting potential implant failure and assisting in planning for future surgical interventions.^{122,137,138} AI-assisted tools have manifested as an efficacious solution to the common difficulties in postoperative CT and MRI associated with metal-related image artefact and distortion.¹³⁹ AI can also help in determining follow-up recommendation plans based on analysing the radiological report.¹⁴⁰

Challenges

The field of musculoskeletal AI is filled with optimism, but there are substantial challenges that must be addressed to enable safe, unbiased and equitable deployment in clinical practice.^{141–143}

Most current ML methods use supervised methods using large data sets, which need manual checks and large amounts of data. This means that the accuracy of the models relies on the quality of the input data. Quality control measures are, therefore, required, and these need timeconsuming human input.

The integration of AI algorithms into the clinical workflow must be performed carefully and thoughtfully and be congruent with the existing information technology infrastructure.^{141–143} The replication of AI studies on 'real-life' data remains problematic due to the sporadic availability of training data and code and must be addressed moving forward and is sometimes neglected in favour of model performance.¹⁴²

The data for future AI research must be suitably anonymized with consideration of patient privacy and ethical considerations.¹⁴² The patients' preference for human experts' diagnoses accentuates the importance of cultivating appropriate levels of trust in AI applications. Additionally, considerable challenges persist in data availability and model validation, especially for rare musculoskeletal disorders and diseases (including vasculitis and CTD). The regulation in AI is developing and is vital to ensure that these tools are created, maintained and used appropriately in clinical practice.^{142,144,145}

Conclusions

Al within the musculoskeletal field harbours immense benefits, and as detailed earlier, a great deal of work has been performed and is ongoing. As research and development progress, Al has the potential to become an increasingly vital component of musculoskeletal care. However, the

above challenges must be considered of utmost importance as the field moves forward.¹⁴⁶ While AI embodies the promise of enhanced patient care, cognizance of the related challenges and ethical considerations is indispensable for wider acceptance and implementation of AI in the musculoskeletal field. \Box

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