Emerging role of artificial intelligence in nuclear medicine

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Review article

Emerging role of artificial intelligence in nuclear medicine
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The role of artificial intelligence is increasing in all branches of medicine. The emerging role of artificial intelligence applications in nuclear medicine is going to improve the nuclear medicine clinical workflow in the coming years. Initial research outcomes are suggestive of increasing role of artificial intelligence in nuclear medicine workflow, particularly where selective automation tasks are of concern. Artificial intelligence-assisted planning, dosimetry and procedure execution appear to be areas for rapid and significant development. The role of artificial intelligence in more directly imaging-related tasks, such as dose optimization, image corrections and image reconstruction, have been particularly strong points of artificial intelligence research in nuclear medicine. Natural Language Processing (NLP)-based text processing and interpretation, automated processing of textual reports, as well as other tasks that were previously thought to be the exclusive preserve of human experts. The other strong appeal of artificial intelligence is its potential to ‘mine’ from medical images in the form of ‘radiomic features’ to extract a vast number of quantitative tumor metrics from medical images, text reports and interventional data. With current generation of hybrid scanners such as PET/computed tomography (PET/CT), PET/MRI and single-photon emission tomography (SPECT)/CT scanners, every patient encounter could generate several gigabytes of data in the form of images [6]. These images are interpreted by an imaging expert and textual data are also generated in the form of diagnostic reports. A significant amount of intervention data is also generated in nuclear medicine every day in the form of clinical reports and dosimetry/postdosimetry calculations. Recently, several mathematical algorithms have been developed to extract a vast number of quantitative tumor metrics from medical images in the form of ‘radiomic features’ [7]. Nuclear medicine data generation is therefore vast, fast, diversified and also highly variable in interpretation, due to human expert subjectivity and variety of scanners from several vendors. Hence, nuclear medicine data clearly qualifies as big data, that satisfies all of the so-called ‘4 V’s’ [8].

Introduction

Artificial intelligence applied to clinical workflows and personalized cancer care has attracted immense attention in recent years [1–3]. The strong appeal of artificial intelligence is due in large part to its potential for significantly speeding up mundane clinical tasks through selective automation. In various domains of cancer care, artificial intelligence tools have shown potential to assist with automated diagnosis, segmentations of normal organs and tumor volumes, complex image transformation and interpretation, automated processing of textual reports, as well as other tasks that were previously thought to be the exclusive preserve of human experts. The other strong appeal of artificial intelligence is its potential to ‘mine’ for diagnostic or prognostic patterns (i.e. a ‘signature’) among a very large number of potential variables in order to make a reliable prediction of outcome.

In the scope of this article, artificial intelligence shall be defined as a scientific study of mathematical processes (i.e. algorithms) that are able to approximate a tightly confined aspect of human cognitive actions, without requiring constant and continuous control by a human operator. The mark of artificial intelligence is therefore computer applications that appear to demonstrate some degree of autonomy, adaptability and agnosticism towards the completion of a narrowly-defined function. Machine learning is a subfield of study in artificial intelligence involving training of mathematical and statistical algorithms, in order to generate the desired output when provided with a given set of inputs. It does so by being exposed to an extremely large number of repeated training episodes whereby the machine is conceptually ‘penalized’ for every incorrect output and ‘rewarded’ for every acceptable output. This dependency on a large number of training instances necessarily makes this type of artificial intelligence exceptionally sensitive to the volume, variety, velocity and veracity of the data on which it has to train on; these four V’s are immediately recognizable as the signature hallmarks of ‘big data’ [4,5] (Fig. 1).

Diagnostic and interventional nuclear medicine departments routinely generate huge amounts of data in the form of medical images, text reports and interventional data. With current generation of hybrid scanners such as PET/computed tomography (PET/CT), PET/MRI and single-photon emission tomography (SPECT)/CT scanners, every patient encounter could generate several gigabytes of data in the form of images [6]. These images are interpreted by an imaging expert and textual data are also generated in the form of diagnostic reports. A significant amount of intervention data is also generated in nuclear medicine every day in the form of clinical reports and dosimetry/postdosimetry calculations. Recently, several mathematical algorithms have been developed to extract a vast number of quantitative tumor metrics from medical images in the form of ‘radiomic features’ [7]. Nuclear medicine data generation is therefore vast, fast, diversified and also highly variable in interpretation, due to human expert subjectivity and variety of scanners from several vendors. Hence, nuclear medicine data clearly qualifies as big data, that satisfies all of the so-called ‘4 V’s’ [8].

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With particular attention to the big imaging data that is found in nuclear medicine, a subfield of machine learning, known as deep learning, has become an extremely powerful artificial intelligence tool for image processing and image analysis. Deep learning algorithms learn composition of data that is represented in a hierarchy of structures of simpler features as a representation of the complex data [5,9]. Deep learning neural networks exploit a vast number of extremely simple computational units, called artificial neurons, which are organized in deep stacks of interconnecting layers [9–11]. Specific deep learning architectures known as convolutional neural networks (CNNs) have been shown to be extremely adaptable to general image-based tasks such as segmentation, object detection and object classification [9–11].

Radiomics

Radiomics refers to the automated extraction and analysis of quantitative features of an image, such as a PET or a CT, in order to recognize potentially useful diagnostic or prognostic signatures [7]. Therefore, radiomics draws very heavily from technology in both machine learning and deep learning domains, to derive statistical prediction models from ultra-high dimensionality data using regression, decision trees, principal components analysis and ever increasing in recent years, deep learning. The radiomics hypothesis posits that medical images are not simply pictures for qualitative interpretation [7], but can be directly converted into minable data that could be used for personalized cancer care [12]. A typical radiomics analysis cycle is schematically summarized in Fig. 2. This involves the extraction of the region of interest on the image, preprocessing (such as digital image filters and resampling, among others), feature extraction and finally model development with validation [13].

Natural Language Processing

Natural Language Processing (NLP) employs machine learning and deep learning tools to aid in the extraction of structured data from natural language sources, such as recorded speech or written text reports or both. There are various well-established applications of NLP in diagnostic and clinical text processing in nuclear medicine such as report classification, report interpretation, sentiment analysis and text generation, etc. [14]. The NLP workflow consists of various steps and is summarized in Fig. 3. The text is first preprocessed by various processes like segmentation in which the text is segmented for sections and sentences. Then certain preposition and conjunction words are removed (stop-word removal). The remaining words are tokenized and normalized by linking to the root word (tokenization and word normalization). The text is then analyzed for grammar and part of speech (syntactic analysis). This is then subject to named entity recognition, concept recognition and relation extraction using relevant ontologies (semantic analysis). Relevant concepts maybe checked for negation using negation detection algorithms. Text features are then extracted and vectorized. The extracted features are used in model development. Models maybe rule-based, machine learning-based or a hybrid of both. The models are developed for defined outcomes and require annotated corpus for training in case of supervised learning. The trained model then undergoes validation with internal data as well as external data. These models maybe developed and applied for the classification of problems, as a part of decision support systems, treatment outcome deduction, summarization or text/report generation.

Applications of artificial intelligence in nuclear medicine

Emergence of big data in cancer care

Over the past several years, continuous efforts have been expended to improve quality of patient care and accelerate the pace of cancer research through the ever-increasing use of artificial intelligence and medical big data informatics [15]. Large volumes of routine (standard of care) data related to cancer diagnosis, treatment planning and outcomes are stored as either structured or unstructured form within the confines of the hospital or clinic in the form of medical records [16]. The conversion of such highly granular patient’s data into electronic formats has opened up a plethora of opportunities for the cancer research community.

However, realization of clinical and societal value from routine medical big data comes with attendant risks. First and foremost, patient confidentiality must be protected at all times [17,18]. Anonymization, pseudonymization and binding legal data-sharing contracts
all go some way towards protecting confidentiality, and increasingly ‘privacy-by-design’ paradigms such as distributed learning and federated machine learning [19–21] are being used, where a diverse range of artificial intelligence algorithms can be successfully trained without exposing private individuals’ identifiable information outside of the data providers’ respective communications network firewalls [22–24]. Second, data quality and provenance issues due to missing data, inexact measurements and undocumented deviations from clinical protocol can lead to serious biases, lack of generalizability and potential patient harm, when developing artificial intelligence models on such data [25]. Last but not least, big data is plagued by interoperability issues such as different formatting standards, idiosyncratic data coding, language barriers and ambiguity in clinical meaning, thus limiting the utilization of this data. The growing impetus behind adherence to Findable-Accessible-Interoperable-Reusable data stewardship standards [26] are beginning to address such problems, thus gradually rendering more big data usable by automated artificial intelligence algorithms.

**Potential benefits of applied artificial intelligence in nuclear medicine**

Several retrospective and prospective studies describe different approaches that could support earlier diagnosis and more accurate prognosis of cancer [27–29]. Various aspects of the clinical problem have been examined utilizing genomics, proteomics, radiomics and pathomics signatures [30,31]. These studies identify the potential as well as the limitations of these signatures for the prediction of cancer outcome. The vast majorities of these publications make use of one or more artificial intelligence algorithms, and further integrate data from heterogeneous sources for generating a prediction. The application of artificial intelligence to quantitative medical imaging has shown promising results in cancer diagnosis, prognosis of disease and treatment outcome prediction.

In the following sections, we will discuss the role of artificial intelligence in several parts of the nuclear medicine clinical workflow. The key area of artificial intelligence implementation in nuclear medicine may be grouped into the planning of procedure, execution of procedure, image reconstruction, image interpretation, report generation and implementation of clinical decision support system. These potential domains of application in nuclear medicine are summarized in Fig. 4 and are discussed in detail in the review.

**Planning of nuclear medicine procedure**

Nuclear medicine procedures are associated with radiation exposure to the patient and are quite expensive. These procedures require patient specific-preparation (such as fasting prior to scan, cessation of insulin if diabetic, etc.) which has been challenging and required highly skilled human resources to juggle a large number
of these requirements [32]. Diagnostic and therapeutic dosimetry planning requires imaging at multiple time points, image processing and mathematical calculations, all of which are technically demanding and widely seen as a challenging clinical task for nuclear medicine professionals [33]. Artificial intelligence algorithms can play a significant role in optimizing patient scheduling, procedure selection, patient prescan preparation and pre-imaging/treatment dosimetry planning. Ansart et al. [34] showed that artificial intelligence-based screening of patients led to an increased number of recruitments as well as reducing the number of expensive PET scans in a clinical trial. Massachusetts General Hospital developed a prediction model using multivariate logistic regression to predict the show or no-show of patient for PET/CT scan [35]. Shi et al. [36] have shown that a trained artificial neural network can reduce the individual radiation dosimetry prediction error by between factor 4 and factor 10, in comparison with population-based dosimetry approach in Lu-177-based dosimetry assessment of tumor and various organs. Xie et al. [37] have utilized 3D deep convolutional network algorithm for automated segmentation of normal organ and fetus with Dice similarity coefficient between 0.92 and 0.98, which resulted in less than 1% difference in the radiation dose calculation for normal organ and fetus for various radiopharmaceuticals by conventional and artificial intelligence-based segmentation model.
Optimization of nuclear medicine procedure

Modern generation scanners have started implementing various artificial intelligence approaches [38] that assist with optimization of image quality, scan time and radiation exposure during diagnostic nuclear medicine procedures. During therapeutic procedures, artificial intelligence-based automation can assist with pre-therapeutic and post-therapeutic dosimetry by using artificial intelligence-assisted organ and tumor segmentation, and prediction of either organ and/or tumor dose using single-point imaging, etc. [39,40]. Utilization of artificial intelligence in individualized dose assessment can assist a nuclear medicine physician to implement pre-therapeutic and post-therapeutic personalized dosimetry-based assessment, which is currently not done because the process is very lengthy, cumbersome and requires high expertise to perform the task.

During PET and SPECT examinations, we often require additional imaging or delayed imaging. For example, if a patient is scheduled for a PET scan but there is a lingering suspicion of brain metastasis, he/she may require an extended study that includes scans of the brain. This is currently a decision taken on the spot by a doctor or technologist, and image sets of such scenarios are available in the nuclear medicine department. These image sets can be used to train a deep learning artificial intelligence model to assist in decision making by predicting or scoring the likelihood of a brain metastasis actually being present.

In modern multidetector CT scanners, the tube current, and hence patient exposure, is being modulated in real-time, to minimize investigational radiation dose, but this is not patient-specific because the minimum and maximum tube currents are pre-defined such that the scanner modulates between the specified values [41]. The disadvantage of this is the dose delivered can only be known after the completion of the scan. An artificial intelligence-assisted system may allow us to perform real-time and patient-specific modulation of the tube current value to obtain the best image quality while optimally reducing radiation exposure to the patient [42].

The amount of radioisotope administered to a patient is currently calculated based on body weight, and there is ample margin for empirical (subjective) judgment. This could potentially lead to excess radiation (more than what is needed for the clinical purpose) or else to excessive noise and suboptimal image quality at lower radiation dose. An artificial intelligence-based radioisotope dosing system can be trained with the help of patient demographic data which can suggest the patient-specific required activity in a much more precise manner. This will reduce unnecessary radiation burden to the patient or avoid having to re-do suboptimal scans due to insufficient dose of activity administered.

Image reconstruction

Nuclear medicine departments have been dealing with the balancing act between radiation doses, longer imaging...
times, low count density in images, low signal-to-noise ratio, high partial volume effect and low spatial resolution since its inception [43]. Advancements in instrumentation technology and reconstruction technology have addressed these concerns to some degree, in multidetector helical CT and high field-strength MRI [44]. Image reconstruction in nuclear medicine imaging is still unable to catch up to its other radiological imaging counterparts. Nuclear medicine imaging involves various computationally intensive steps including attenuation correction, scatter correction, noise correction and partial volume correction. Reconstruction algorithms in use today range from filtered back projection to newer techniques such as 3D Ordered Subset Expectation Maximization, row-action maximum likelihood algorithm, often in combination with advanced signal processing techniques like partial volume correction and time-of-flight (TOF) correction [45]. These techniques have substantially improved image quality in nuclear medicine without necessarily leading to higher dose in the patient.

Recent literature demonstrates a paradigm shift in image reconstruction in nuclear medicine imaging from analytic predetermined approaches towards a more adaptable, heuristic and potentially more patient-specific approach. Several researchers have examined artificial intelligence-based attenuation correction, scatter correction in PET and SPECT image reconstruction and extensive research is being performed to improve image quality of low count PET scan [46–50]. Hong et al. [46] used a CNN to enhance the image resolution of PET scans. Xiang et al. [47] have shown promising results with deep learning to improve the image quality with low dose and reduced scan time reconstruction during PET imaging. Wang et al. [48] demonstrated the utility of 3D deep convolutional network to predict high-dose PET scan using low administered activity. Kim et al. [49] used a de-noising deep learning network with local linear fitting to improve image quality of PET in an iterative reconstruction algorithm. Shiri et al. [50] demonstrated the utility of a deep learning residual network to synthetically generate a full-dose myocardial SPECT image from low-dose SPECT data as well as he also demonstrated the utility of artificial intelligence in prediction of full-time acquisition SPECT image from half-time acquisition SPECT imaging data.

Multimodality fusion imaging with hybrid scanners has proven to be a significant breakthrough in medical imaging; modalities such as combined-PET/CT have now been established as the modality of choice for cancer imaging [51]. Combined-PET/MRI is another hybrid medical scanner that is rapidly growing its importance in cancer imaging, but it needed to overcome the challenges related to MRI-based attenuation correction in PET for accurate PET imaging and quantification [52]. This can be achieved in number of ways, including the use of deep learning to synthetically generate a radiation attenuation map (in effect, a synthetic CT) directly from MRI. Several analytical techniques have been tested to improve PET attenuation correction in PET/MRI scanners [52]. However, recent literature points towards a growing role of artificial intelligence [53–57] with a particularly active line of study in image corrections. Arabi et al. [53] have further shown the feasibility of deep learning to estimate the attenuation correction based solely on TOF PET emission data. Arabi et al. [54] also estimated the attenuation and scatter corrections simultaneously in a multitracer neuroimaging study, Shiri et al. [55], in a similar kind of study, used deep learning to estimate the joint scatter and attenuation correction in PET image. Hwang et al. used a deep learning neural network to apply a Maximum Likelihood reconstruction of Attenuation and Activity algorithm to jointly generate activity and attenuation maps using emission data [56]. Liu et al. [57] in another study have used the deep learning technique to generate the attenuation map for PET reconstruction using MRI images.

### Image interpretation and report generation

Image interpretation and reporting of scan are amongst the most important tasks in clinical nuclear medicine as it is in radiology. The entire process involves various steps as image reading, extraction of quantitative data, comparison of a scan with earlier scans (quantitatively and qualitatively), then reporting on the findings with (in general) a clinical interpretation of the significance of the findings. The potential of artificial intelligence here lies in the valuable time saving of nuclear medicine physicians by automatic execution of more mundane jobs in the preparation of the report. Like reading the patient’s history from the electronic medical record, the consolidation of the history in the report, quantitative data extraction and comparison often consumes lots of time and leads to reduced output from an expert nuclear medicine physician or radiologist [58–60]. Artificial intelligence tools may be utilized to reduce this burden so that the expert can do more meaningful tasks like interpretation of finding, providing expert opinion and contributing to multimodality treatment groups [58–60]. NLP is an evolving technology that can assist with nuclear medicine report processing in many ways. Several studies have shown the use of NLP in text processing in radiology. Pons et al. [14], in their systematic review of literature of NLP in radiology reporting, have identified the potential use of NLP in diagnostic surveillance, cohort building for epidemiologic studies, query-based case retrieval, and quality assessment of radiology practice and clinical support services with level of evidence. These same clinical needs exist in nuclear medicine. Pinto et al. [61], in their narrative review, have also emphasized the potential importance of artificial intelligence in processing radiology reporting requirements as well as the importance of enforcing structured reporting to increase the use of artificial intelligence for radiology data-mining.
**Clinical decision support system**

Nuclear medicine imaging plays an important role in personalized medicine. A theranostics concept is being used in radionuclide therapy to predict the efficacy of radionuclide therapy [62]. Various parameters like standardized uptake value (SUV), total lesion glycolysis (TLG) and PET Response Criteria in Solid Tumors (PERCIST), etc. have been utilized for disease outcome prognostication [63–65]. Mahadeviah et al. [66] described the use of artificial intelligence in decision support systems in oncology, but very similar concepts will also hold for nuclear medicine. The potential role of artificial intelligence is being explored by several researchers working on the development of various kinds of prediction models for outcome predictions in oncology, such as overall survival (OS), progression-free survival (PFS), loco-regional recurrence, distant metastasis, treatment outcome, toxicity and treatment selection. These studies utilize a range of qualitative, semiquantitative (SUV, TLG and PERCIST) and radiomics features extracted from nuclear medicine images [67–74].

Radiomic features derived from nuclear medicine images have been proposed as a potential digital imaging phenotype for disease-specific personalized predictions of individual outcomes [13]. Sanduleanu et al. [67] used a univariate Cox model to show that [F-18]-HX4 PET/CT hypoxia uptake was able to prognose OS and local PFS in head and neck cancer patient treated by chemotherapy [Spearman’s correlation coefficient ($\rho_s$) = 0.77] ($P < 0.001$). In a multicenter study, Betancur et al. [68] showed that a CNN model trained using polar maps of myocardial perfusion imaging (MPI) from SPECT scanners outperformed the human expert diagnosis [area under the curve (AUC): 0.81 vs. 0.78] for detecting obstructive myocardial disease. In an extended study, Betancur et al. also developed a prediction model combining clinical information and MPI data to predict the 3-year probability of major cardiac events. They demonstrated that the machine learning technique outperformed clinical decision-making by medical experts (AUC: 0.81 vs. 0.65) for predicting major cardiac events in a cohort of 2619 consecutive patients [69]. A similar study by Rios et al. [70] also suggested an improvement of radiomics biomarkers over conventional risk factors (AUC: 0.793 vs. 0.698) when predicting major adverse cardiac events. Shiri et al. [71] used a combination of radiomics with genomic sequencing to predict presence of epidermal growth factor receptor (AUC: 0.82) and V-KI-RAS2 Kirsten rat sarcoma viral oncogene homolog (KRAS) (AUC: 0.83) mutations from images of nonsmall cell lung cancer (NSCLC) patients. Kidd et al. [72] developed an FDG PET-based nomogram (SUVmaximum value and lymph node status) to predict recurrence-free survival (C-stat: 0.741), disease-specific survival (C-stat: 0.739) and OS (C-stat: 0.658) for locally advanced cervical cancer. However, some studies have also reported the poor validation performance of radiomic-based prognostic models. Foley et al. [73] performed an external validation of a prognostic model for oesophageal cancer using quantitative PET features and radiomic features but found that these features were unable to discriminate between patient groups with different OS in an independent external validation cohort. van Timmeren et al. [74] also performed a multicenter radiomic-based study for treatment response assessment in NSCLC patients; they were able to demonstrate significance of FDG-PET/CT radiomic model in their individual-institution cohort but failed to reproduce this with a multicenter study.

**Summary of review**

The emergence of artificial intelligence applications in nuclear medicine appears to be on the verge of contributing significantly throughout the clinical workflow atomization. Initial research suggests an increasing role of artificial intelligence in nuclear medicine imaging, particularly where selective automation of tasks are concerned. Artificial intelligence-assisted planning, dosimetry and procedure execution appear to be areas for rapid and significant development. The role of artificial intelligence in more directly imaging-related tasks, such as dose optimization, image corrections and image reconstruction, have been particularly strong points of artificial intelligence research in nuclear medicine. A further aspect of artificial intelligence research and implementation in nuclear medicine will be disease characterization, prognostication and potentially treatment outcome prediction. The new study of radiomics has been especially active in this field, leading towards the potential for identifying image-based digital signatures of oncological disease that can inform clinical decision-making, and therefore a stepping stone towards individualized cancer care. Artificial intelligence-assisted image interpretation based on deep learning neural networks have the potential to open up a new horizon in quantitative nuclear medicine, and could become a powerful future tool in the hands of nuclear medicine physicians and radiologists. Lastly, an aspect of artificial intelligence that still needs to develop strongly and could be of immense utility in nuclear medicine is domain-specific NLP. Complex NLP tools may help to automate the report generation and clinical interpretation tasks, and likely enrich the clinical interpretation with the aid of quantitative features from radiomics and deep learning analysis.

As a counterpoint to the promising potential future of artificial intelligence in nuclear medicine, one needs to be cognizant of the potential challenges and barriers to widespread artificial intelligence adoption in nuclear medicine clinics. Data, in vast quantities and with large dimensional variety, are the raw material for developing and testing artificial intelligence applications for nuclear medicine. While some data may be found in great abundance (e.g. PET and SPECT images), other types of data...
may be dispersed across other clinical departments (e.g. chemotherapy treatment details, or radiotherapy delineations, or long-term clinical follow-up) or have become disconnected completely from the radioisotope imaging due to suboptimal data management procedures. It is not simply the image data that needs to be curated and maintained with a view to long-term data sustainability, but it is also the metadata about the investigation that needs to be captured. For example, imaging settings, equipment and scanner software are likely to change rapidly over a short period of time, and we know that differences among vendors’ equipment and variation in imaging parameters are a major concern that directly limits the wider applicability of radiomics models [75] and also of deep learning-based image analysis models [76].

A second major barrier to artificial intelligence applicability in nuclear medicine is one of clinical acceptability. This acceptability has two distinct dimensions; clinical acceptability and psychological acceptability. The former requires a detailed and practical process of selecting, commissioning and validating an artificial intelligence tool with realistic and as locally-specific clinical conditions as reasonably possible [66]. This necessarily entails staff training, ongoing quality assurance and careful integration of emerging artificial intelligence tools into the clinical environment with oversight for safety and ethical usage. The latter degree of acceptability is much harder to quantify and define; however, it relates directly to another major topic of research in artificial intelligence, that is, the degree of ‘explainability’ (or lack thereof) in an artificial intelligence-based decision support system [77]. While a clinician can readily and eloquently supply a rationale for his/her clinical decisions to a qualified peer, to a presiding judge or to an inquiring patient, certain types of artificial intelligence are still lacking in this fundamental rationalizing power of an ‘explanation’. For instance, why exactly does an imaging-related signature lead an artificial intelligence-based system to recommend a particular course of action or to suggest a particularly unpleasant prognosis? The risk for the clinician is, either to act upon or else to over-ride such recommendations from an artificial intelligence system (particularly in a question of diagnosis, prognosis or treatment selection). Either of these feels potentially psychologically unsafe in the absence of suitable explanation or justification from an artificial intelligence-based system, leading to deep mistrust or lack of application of artificial intelligence-based nuclear medicine tools.

**Conclusion**

The role of artificial intelligence is increasing in prominence in the clinical nuclear medicine workflow, starting from patient preparation to the development of decision support system in medicine. There appears to be ample evidence of development and advancement of artificial intelligence addressing nuclear medicine needs. This is pointing to an ever-increasing role of artificial intelligence in nuclear medicine. In the coming years, the computer science, informatics and clinical specialist fields need to collaborate closely to address some major hurdles related to the long-term sustainability of high-quality data (including imaging metadata) and clinical barriers to the adoption of artificial intelligence.

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**Conflicts of interest**

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