# Applications of Artificial Intelligence for Pediatric Cancer Imaging

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# Pediatric Imaging · Review

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Artificial intelligence (AI) is transforming the medical imaging of adult patients. However, its utilization in pediatric oncology imaging remains constrained, in part due to the inherent scarcity of data associated with childhood cancers. Pediatric cancers are rare, and imaging technologies are evolving rapidly, leading to insufficient data of a particular type to effectively train these algorithms. The small market size of pediatric patients compared with adult patients could also contribute to this challenge, as market size is a driver of commercialization. This review provides an overview of the current state of Al applications for pediatric cancer imaging, including applications for medical image acquisition, processing, reconstruction, segmentation, diagnosis, staging, and treatment response monitoring. Although current developments are promising, impediments due to the diverse anatomies of growing children and nonstandardized imaging protocols have led to limited clinical translation thus far. Opportunities include leveraging reconstruction algorithms to achieve accelerated low-dose imaging and automating the generation of metric-based staging and treatment monitoring scores. Transfer learning of adult-based Al models to pediatric cancers, multiinstitutional data sharing, and ethical data privacy practices for pediatric patients with rare cancers will be keys to unlocking the full potential of Al for clinical translation and improving outcomes for these young patients.

Artificial intelligence (AI) algorithms are transforming the medical imaging of adult patients, facilitating image acquisition, reconstruction, quality control, segmentation, analysis, and interpretation [1–6]. However, the development of AI algorithms for pediatric applications and, in particular, for pediatric oncologic applications is limited to date. According to an overview by the American College of Radiology, only 3% of FDA-approved AI applications for medical image analysis (seven of 221 total products) are tailored for pediatric imaging [7]. None of these are specific to oncologic applications in children, highlighting a critical gap in addressing the diagnosis and treatment of pediatric cancer with AI-powered tools.

Algorithms approved for adult patients can potentially be used in children, with the likelihood of success strongest if the imaging technique and anatomy are similar in children and adults for the given context. However, to date, limited evidence has addressed the performance of adult algorithms when applied in children. One example that has been explored is the use of Al algorithms for automated lung nodule detection on chest CT [8]. However, studies have found that adult-based lung nodule detection algorithms do not perform as well in children as they do in adults [8, 9].

Many Al algorithms for pediatric cancer applications require specific development for children. However, the rarity of childhood cancers and the ongoing rapid evolution of imaging technologies limit the availability of large standardized datasets, which in turn hinders robust training of Al models for pediatric oncology applications. The rarity of childhood cancers translates to a smaller patient population; as market size is a driver of commercialization, this smaller market size of pediatric patients compared with adult pa-

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tients potentially and, in association with relatively limited financial returns, could be a disincentive to developers. Another major challenge is the lack of publicly available whole-body image datasets of children with cancer. Of 208 imaging datasets published (as of March 19, 2024) on The Cancer Imaging Archive, one of the largest public cancer imaging databases, eight are dedicated pediatric datasets [10]. The Cancer Imaging Archive collaboration with the Childhood Cancer Data Initiative of the National Cancer Institute provides a repository of high-quality CT, MR, and PET images of pediatric nasopharyngeal cancer, Hodgkin lymphoma, Wilms tumor, medulloblastoma or primitive neuroectodermal tumor, and liver cancer [10]. Another public imaging repository, the Medical Imaging and Data Resource Center, is continuously growing but does not have a dedicated pediatric oncologic context at this time [11]. The 2017 Radiological Society of North America (RSNA) database contains a large dataset of pediatric hand radiographs, which could be used as normal reference data [12]. The Children's Oncology Group has a central image registry Quality Assurance Review Center (QARC) of imaging studies of children with cancer before and after chemotherapy. The QARC image registry includes large but heterogeneous image datasets and can be accessed by interested parties through an administrative application process [13]

Many of these image registries include image datasets collected from different centers and scanners. The resulting advantage is that the registries provide data from patients with diverse demographic backgrounds. However, the heterogeneous imaging protocols and techniques of multiinstitutional image registries pose a challenge for training Al algorithms. The Children's Brain Tumor Network (CBTN) addressed this problem by generating consensus about a shared imaging protocol before collecting image data for 4900 patients in a shared database [14]. Similar efforts initiated by the Children's Oncology Group or other authoritative bodies could enhance the utility of their image registries for Al algorithm development.

Although the field of AI development is rapidly expanding, the impact of AI algorithms on the care of children with cancer has not yet been established. Today, most AI systems primarily focus on slice-by-slice or single-plane image analysis. However, imaging studies of children with cancer often require multiplanar or volumetric analysis, which are beyond the capabilities of current AI solutions. AI struggles with complex anatomic structures and pathologies spanning multiple imaging planes. Although AI holds immense promise, real-world gains have been relatively slow.

This review provides an overview of the current state of Al applications for pediatric cancer imaging, including applications for medical image acquisition, processing, reconstruction, segmentation, diagnosis, staging, and treatment response monitoring. Although some of the aspects of image data processing discussed can be applied to pediatric imaging in general, the downstream tasks related to tumor diagnoses are specific to oncologic imaging.

# **Image Data Acquisition and Processing**

Al algorithms can be developed for upstream image processing (i.e., generation of medical images before interpretation) and downstream image processing (i.e., detection and characterization of areas of interest on a medical image). Al can enable safer

# Highlights

- Al algorithms can accelerate the acquisition, reconstruction, and processing of medical images of pediatric cancers.
- Challenges in developing commercial AI algorithms for pediatric tumor segmentation include heterogeneous imaging protocols and patient anatomy, a paucity of multiinstitutional data, and limited market size.
- A critical gap exists between promising AI methods for pediatric cancer diagnosis and treatment response monitoring and their actual use in clinical practice.

image acquisition by reducing the patient's radiation exposure. Examples include augmenting low-radiotracer-dose PET images to standard-dose PET images [1, 3] and augmenting ultralow-dose CT images to standard-dose CT images [15]. Al can also accelerate imaging times by augmenting ultrafast low-resolution MR images to high-resolution images [16], enhancing image quality, and minimizing image noise. Some of these MRI augmentation algorithms are commercially available [17, 18]. However, most are trained on adult datasets and lack oncologic context.

Chaudhari et al. [3] developed a deep learning algorithm for augmenting standard-dose PET images from fourfold reduced-count whole-body PET scans of children and young adult patients. Wang et al. [1] developed a novel Al model for the augmentation of ultralow-dose FDG PET/MR images of 33 children and young adults (age range, 3–30 years) with lymphoma, which reduced patients' radiation exposure by more than 90%. Next, Wang et al. [19] compared five Al models for enhancing the image quality of ultralow-dose pediatric PET scans. SwinIR and U-Net achieved superior performance compared with three other Al algorithms for augmenting ultralow-dose input images.

Many Al algorithms focus on pretreatment scans only and do not consider posttreatment scans. To address this gap, in 2021 Theruvath et al. [2] evaluated the performance of a convolutional neural network (CNN) for augmenting the static low-dose FDG PET/MRI scans of patients with pediatric lymphoma before and after chemotherapy. Simulated low-dose scans showed increased noise and decreased tumor-to-liver contrast, hindering therapy response assessment on posttreatment scans. Although CNN augmentation improved performance at 75% and 50% reduced-dose images, achieving sensitivity and specificity of 100% for response classification at both simulated doses, significant errors persisted if augmentation was attempted for images with simulated radiotracer dose reduction of more than 50% (i.e., simulated reductions of 25%, 12.5%, and 6.25%) [2] (Fig. 1).

Other than for PET, AI applications have also been studied to enhance image data acquisition and processing of MRI and CT. Ladefoged et al. [20] compared two MRI-based attenuation correction methods for PET/MRI scans of pediatric brain tumors. Both methods were as accurate as traditional CT-based attenuation correction. However, a DeepUTE algorithm outperformed Resolute (another MRI-based method). Maspero et al. [21] used MRI-derived synthetic CT (sCT) images for dose calculation in pediatric brain tumor radiotherapy. By combining sCT examinations

from multiple planes and accounting for MRI protocol variations, the method achieved accurate dose calculations with minimal discrepancies compared with standard CT. This finding supports MRI-based sCT as a promising tool for this application (Fig. 2).

A significant challenge in the application of AI for medical image augmentation lies in striking a careful balance between noise reduction and preservation of image fidelity [22]. Noise that can hinder analysis must be removed, while crucial details essential for accurate diagnosis simultaneously must be preserved. Image processing with Al algorithms can obscure subtle yet diagnostically relevant features. Despite the widespread adoption of Al-processed medical images, such as Al-processed single-shot T2-weighted or fast spin-echo (FSE) T2-weighted sequences, concern exists regarding the potential loss of crucial diagnostic information for Al-augmented MRI scans [23]. The original images can be accessed on some scanners. However, this process requires retrieving source images that may not be routinely provided on the PACS, adding a time-consuming step to the workflow.

Al algorithms for the augmentation of CT scans have been used to reduce the radiation exposure of children undergoing CT.

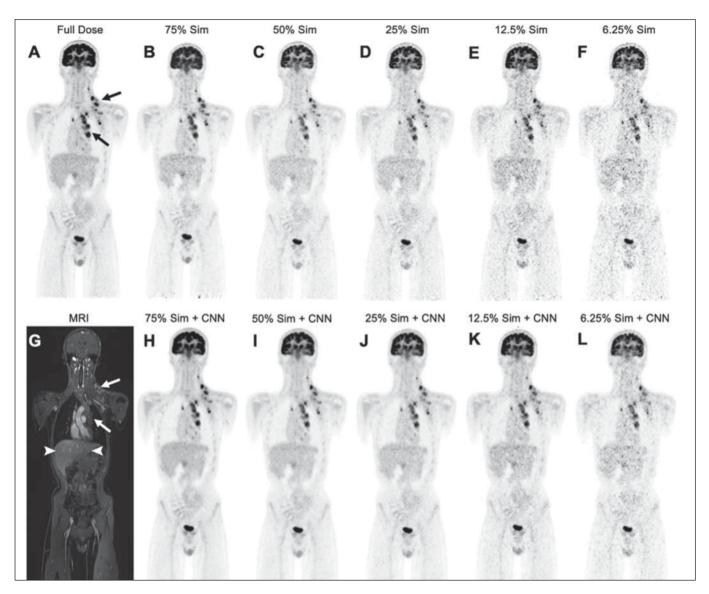


Fig. 1—14-year-old boy with Hodgkin lymphoma. (Used with permission of Radiological Society of North America, from Validation of deep learning-based augmentation for reduced 18F-FDG dose for PET/MRI in children and young adults with lymphoma, Theruvath AJ, Siedek F, Yerneni K, et al. Radiol Artif Intell, 3, 2021; permission conveyed through Copyright Clearance Center, Inc.)

A, Hypermetabolic mediastinal and left infra- and supraclavicular lymph nodes (arrows) are clearly visible on full-dose FDG PET image (3 MBg/kg).

G, Increased noise and reduced contrast between tumor (arrows) and liver (arrowheads) are also seen on corresponding coronal MR image.

B-F, Increased noise and decreased tumor-to-liver contrast are seen on coronal FDG PET images obtained using simulated (Sim) reduced radiotracer doses of 75% (B), 50% (C), 25% (D), 12.5% (E), and 6.25% (F),

H-L, Corresponding low-dose FDG PET images augmented with convolutional neural network (CNN) show reduced noise and improved contrast between tumor and liver compared with non-CNN-augmented PET images.

Brady et al. [15] achieved a radiation dose reduction of 48% with the adaptive statistical iterative reconstruction algorithm (Fig. 3).

Salman et al. [24] assessed a commercially available lung computer-aided detection (CAD) system for pulmonary nodule detection on pediatric CT scans with simulated dose reductions (75%, 50%, and 25%). Sensitivity remained low (24–27%) but was not significantly impacted by lower doses.

Pediatric studies evaluating adult lung CAD algorithms on chest CT have reported concerning results. In a study by Salman et al. [8], a pediatric CAD system for lung nodule detection showed low sensitivity (26–39%) with moderate PPV (48–62%), even with thinner slices or exclusion of smaller nodules. These findings suggest that directly applying adult-derived algorithms to pediatric CT scans may not be effective, and they highlight the need for pe-

diatric-specific AI development. A key hurdle in using existing AI models for pediatric data is the mismatch between training data and real-world applications. Models trained on adult data may underperform when encountering the distinct anatomies of children, potentially impacting diagnosis and treatment [25].

# **Image Segmentation**

In medical imaging, segmentation involves the identification, localization, and isolation of predefined structure(s) within the medical image(s) to identify tumors, organs, or other structures for image processing and interpretation. Encoder-decoder architectures, such as the U-Net architecture [26, 27], have shown good overall performance in segmenting anatomic structures in adult patients, such as the abdominal organs [28, 29], cardio-

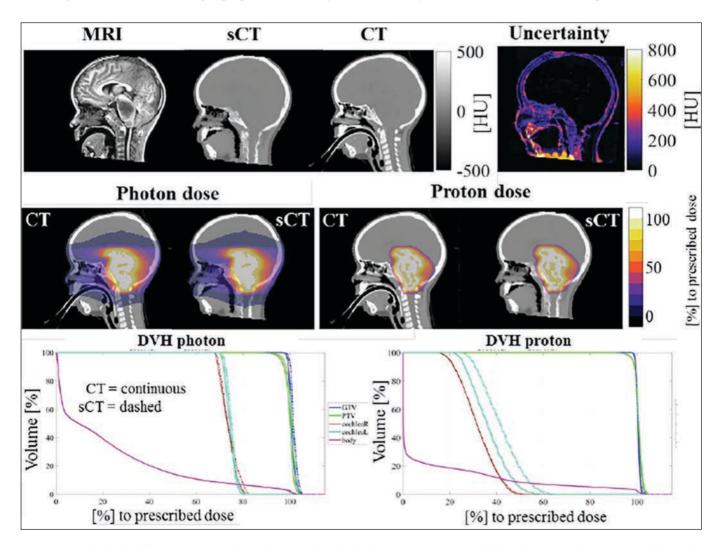


Fig. 2—5-year-old girl with diffuse intrinsic pontine glioma that was irradiated with palliative intent using prescribed dose of 39 Gy in 13 fractions. Top row (*left* to *right*) shows MR image, simulated CT (sCT) image, CT image, and uncertainty map. Middle row shows photon dose plan (*left*) and proton dose plan (*right*), as calculated on CT (solid lines) and sCT (dashed lines) as shown on respective dose-volume histograms (DVHs) (bottom row). MRI was acquired without gadolinium-based contrast media at 1.5 T. This patient had largest relative difference in photon dose in study sample, with difference of -0.8% for high-dose region (dose > 90% of prescription). MRI and CT were performed 3 days apart. Patient was scanned while under anesthesia, and tubes used for anesthetic administration are visible on both modalities. Quality of sCT is lower in neck region, toward end of MRI FOV; uncertainty is larger in same region. GIV = gross inner volume, PIV = planned inner volume, cochleoR = right cochlea, cochleoL = left cochlea. (Reproduced from Maspero M, Bentvelzen LG, Savenije MHF, Guerreiro F, Seravalli E, Janssens GO, van den Berg CAT, Philippens MEP. Deep learning-based synthetic CT generation for pediatric brain MR-only photon and proton radiotherapy. *Radiother Oncol* 2020; 153:197, © 2020, with permission under Creative Commons Attribution 4.0 International Public License [creativecommons.org/licenses/by/4.0/legalcode], including disclaimer of warranties in Section 5)

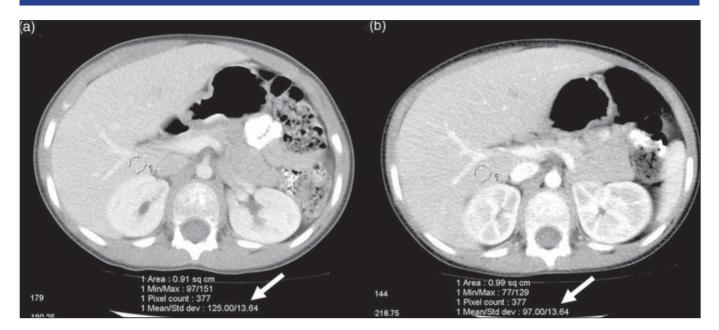


Fig. 3—Patient weighing 23 kg underwent two CT examinations performed approximately 30 days apart. Both images show same SD of attenuation of 13.64 HU (arrow) in 1-cm<sup>2</sup> ROI (circle labeled "1") posterior to right portal vein. Min = minimum, max = maximum, Std dev = SD. (Used with permission of John Wiley & Sons-Books, from Characterization of adaptive statistical iterative reconstruction algorithm for dose reduction in CT: pediatric oncology perspective, Brady SL, Yee BS, Kaufman RA, Med Phys, 39, 2012; permission conveyed through Copyright Clearance Center, Inc.)

A, Axial image from first CT examination, performed without adaptive statistical iterative reconstruction (ASiR).

B, Axial image from second CT examination, performed with ASIR at level of 40%. Use of ASIR increases noise index by 25% and reduces minimum milliamperessecond for automatic tube current modulation. Second examination had dose reduction of 41%. ASIR is variant of model-based iterative reconstruction algorithm that uses advanced computational and statistical techniques integral to artificial intelligence.

vascular structures [29, 30], and neuroanatomy [29, 31]. However, there is a scarcity of segmentation algorithms for pediatric datasets. Recently, a small number of pediatric segmentation datasets have been published for automated CT-based segmentation of up to 29 anatomic organ structures per patient on 359 abdominopelvic CT examinations (with possible inclusion of the chest) [32], segmentation of six organ structures (liver, lungs, bladder, kidney, bones, and brain) on 140 CT images [33], and segmentation of the heart on 64 cardiac MR images [30].

Due to the relatively high standardization of brain MRI protocols, segmentation of brain tumors has been widely studied in adult and pediatric populations [31, 34-39]. The BraTS (Brain Tu-

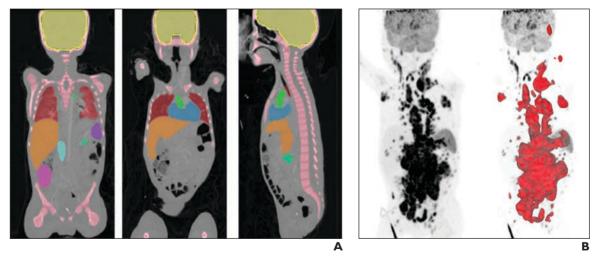


Fig. 4—5-year-old boy with posttransplant lymphoproliferative disorder (PTLD), evaluated by FDG PET/CT. A, Coronal (left and middle) and sagittal (right) fused PET/CT images show automated multiorgan segmentation using publicly available U-Net deep learning model, including segmentation of brain (yellow), bones (pink), lungs (red, left and middle), trachea (red, right), heart (blue), aorta (green), liver (orange), and spleen (purple). Model was trained on adult population of patients with cancer [69].

B, Coronal PET images show automated segmentation of PTLD tumor lesions using available U-Net deep learning model. Model was trained in adult population of patients with lung cancer, lymphoma, and malignant melanoma [46].

mor Segmentation) [34] challenge was one of the first and most widely recognized biomedical machine learning challenges. Based on methodologic advances in brain tumor segmentation in adult populations, studies have recently described automated segmentation of pediatric brain tumors (glioma, ependymoma, and medulloblastoma) in MRI and PET/MRI data using deep learning techniques [35, 40].

Fully automated Al algorithms for the detection of pulmonary nodules on chest CT require two steps: lung segmentation and pulmonary nodule segmentation. For well-defined tasks such as pulmonary nodule segmentation on chest CT, automated machine learning tools are currently commercially available and have shown good performance in children [41]. Li et al. [41] achieved high accuracy (0.9479), Dice score (0.9678), precision (0.9711), and recall (0.9715) performance.

Tumor segmentations could be either automated or user guided. User-guided segmentation is more attainable but requires a dedicated viewer that supports AI algorithms and appropriate editing tools. Rickard et al. [42] described a semiautomatic segmentation algorithm for measuring tumor and kidney volumes on CT images of children with Wilms tumors before, during, and after treatment. This technique could potentially aid surgical planning, treatment response assessment, and prediction of long-term kidney function [42]. In 2023, deep learning-based Al algorithms have been used by Veiga-Canuto et al. [43], for neuroblastoma detection and segmentation on MRI, and by Klimont et al. [44], for automated segmentation and volumetry of pediatric lymphoma on contrast-enhanced CT.

Segmenting tumor lesions in whole-body imaging data presents unique challenges, particularly for pediatric patients. First, the high variability in imaging protocols across different institutions and scanners makes consistent lesion identification difficult. Second, the inherent complexity of multiorgan anatomy in whole-body images adds another layer of difficulty. Third, the diverse range of shapes and sizes in pediatric patients, spanning neonates to adolescents, further exacerbates these segmentation challenges. Deep learningbased tumor segmentation algorithms for whole-body CT, PET/CT, and MRI have been tested in adult patients with varying success [45, 46]. Deep learning algorithms for organ and lesion segmentation in adult patients can be cross-trained using transfer learning for applications in pediatric imaging. Figure 4 shows an example of U-Netbased segmentation algorithms, pretrained on adult CT and PET data and cross-trained for whole-body organ segmentation on CT and detection of tumor lesions on FDG PET.

# **Tumor Diagnosis and Treatment Response**

Diagnosing tumors heavily relies on the ability of radiologists to assess multiple features such as location, size, border characteris-

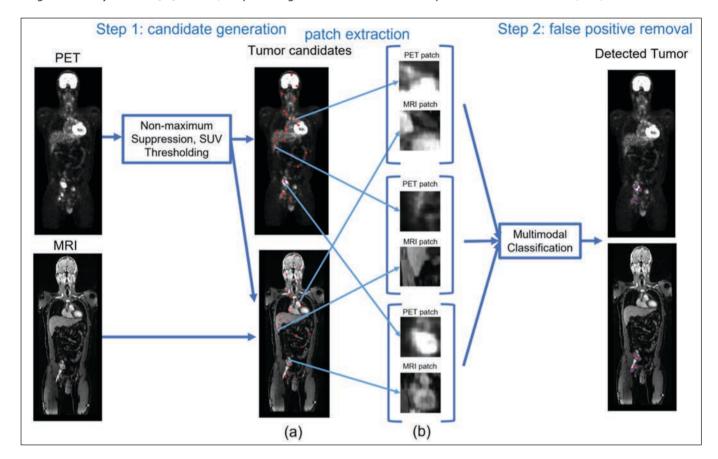


Fig. 5—Two-step method for detecting pediatric lymphoma. (Reproduced from [64], with permission) A and B, Initially, potential tumor sites are identified by tumor candidate generation process using PET images, with nonmaximum suppression used to identify areas where SUV exceeds 2.0. Second step involves eliminating false-positive results. This process is achieved by use of deep learning-based algorithm and four experimental approaches that analyze patches extracted from both PET and MRI examinations through technique that integrates multiple imaging modalities.

tics, internal variation, and contrast material uptake. Researchers are actively developing Al and radiomic tools to assist radiologists in diagnosing pediatric tumors. Radiomics involves extracting and analyzing handcrafted mathematically defined image features, whereas Al encompasses a broader range of techniques, including deep learning methods, that may or may not use these features. Combining these approaches holds promise for improving patient care. However, limited pediatric data hinder the development and implementation of these tools in clinical settings.

#### Musculoskeletal Tumors

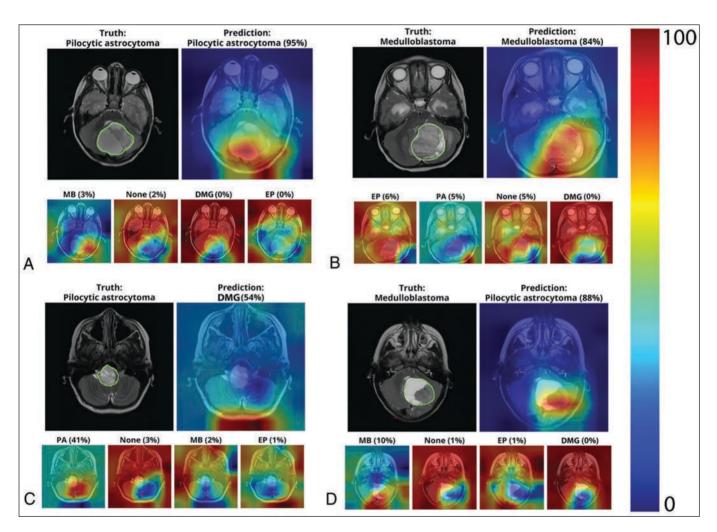
Osteosarcoma—Pereira et al. [47] built a machine learning-based radiomics model to predict lung metastases in patients with osteosarcoma using CT scans of the primary tumor. Their model, trained on data from 81 patients, achieved accuracy of 73% in identifying patients at risk for pulmonary metastases.

Further validation on larger, diverse, and prospective datasets is needed to confirm the clinical utility of such approaches.

Teo et al. [48] developed a deep learning classification model for assessing the chemotherapy response of high-grade osteosarcoma on multimodal MRI in 10 children. They achieved an average accuracy of more than 90% in differentiating necrotic (dead) and viable tumor areas using conventional MR images such as T1-weighted, STIR, and postcontrast images. Diffusion-weighted MRI and dynamic contrast enhancement parameters further improved accuracy.

Huang et al. [49] compared different MRI techniques for assessing tumor response to chemotherapy in patients with osteosarcoma. Their machine learning model, which incorporated T1-, T2-, and diffusion-weighted MRI data, achieved higher accuracy than ADC alone in differentiating necrotic and viable tumor tissue.

Jeong et al. [50] investigated a machine learning approach using baseline FDG PET to predict response to neoadjuvant chemo-



**Fig. 6**—Class activation maps produced by multiinstitutional MRI-based deep learning model developed by Quon et al. [6]. In each panel, top left image shows tumor areas manually outlined on T2-weighted image; top right image shows class activation map overlay of most-confident prediction made by model. Four images in bottom row show less-confident predictions. MB = medulloblastoma, DMG = diffuse midline glioma, EP = ependymoma, PA = pilocytic astrocytoma. (Used with permission of American Society of Neuroradiology, from Deep learning for pediatric posterior fossa tumor detection and classification: multiinstitutional study, Quon JL, Bala W, Chen LC, et al., *AJNR*, 41, 2020; permission conveyed through Copyright Clearance Center, Inc.) **A** and **B**, Model correctly predicts PA (A) and MB (**B**).

**C** and **D**, Model incorrectly predicts DMG in patient with PA (**C**) and PA in patient with MB (**D**).

therapy in 70 patients with osteosarcoma. They analyzed both traditional PET metrics (SUV<sub>max</sub> and tumor volume) and image-derived textural features. Although traditional metrics showed no significant difference between responders and nonresponders, machine learning models using textural features achieved better prediction accuracy (AUC up to 0.82). Analyzing the changes in FDG heterogeneity derived from PET using machine learning offered improved prediction accuracy (AUC up to 0.863). This study was limited by its small sample size. Consequently, the authors recommended a need for data from a larger patient cohort to validate the findings and develop a more robust predictive model.

Ewing sarcoma—Consalvo et al. [51] developed and validated an algorithm for the detection of bone tumors on 182 radiographs of 58 children. The algorithm achieved a diagnostic accuracy of 94.4% and 90.6% for lesion detection and 90.3% and 86.7% for the differentiation of osteomyelitis and Ewing sarcoma in validation and test data, respectively [51].

Gitto et al. [52] compared the predictive value of 3D and 2D MRI radiomics to assess response to neoadjuvant chemotherapy in 30 patients with Ewing sarcoma. The 3D radiomic feature model achieved sensitivity of 85% and specificity of 87% in predicting treatment response [52].

These results were not compared with those of human readers, who typically achieve sensitivities and specificities greater than 90% [53] given that Ewing sarcomas of bone usually show a significant decrease in size in response to chemotherapy [54, 55].

#### **Abdominal Tumors**

To date, limited research on the development of AI algorithms for the diagnosis of solid pediatric abdominal tumors on MRI and CT has been performed.

Wilms tumor—Zhu et al. [56] developed a CNN-based deep learning algorithm to differentiate 269 Wilms tumors from 95 non-Wilms tumors on the basis of the preoperative triphasic CT images of 364 pediatric patients, with histopathology used as the reference standard. Al achieved significantly higher sensitivity (78.1%) for identifying non-Wilms tumors than did the human experts (13.3–20%). Ma et al. [57] developed a support vector machine (SVM) model with 15 radiomic features to distinguish early- and advanced-stage Wilms tumors on preoperative CT in 118 children. The model showed accuracy, sensitivity, and specificity of 79%, 87%, and 69%, respectively [57]. Further development is needed to distinguish all four Wilms tumor stages (1–4) for more precise clinical guidance. This path reflects the incremental nature of Al development, whereby future work builds on initial successes.

In Europe, where preoperative chemotherapy for Wilms tumors is common, Sharaby et al. [58] developed a computer-aided system to predict treatment response in 63 patients with 46 chemotherapy-responsive and 17 nonresponsive Wilms tumors. Their model achieved accuracy of 95.24%, sensitivity of 95.65%, and specificity of 94.12% for prediction of chemotherapy response. That study did not mention the reference standard. Resectability involves many features beyond the size, shape, and texture of the primary tumor, such as tumor thrombus presence and extent, infiltration of adjacent structures, and tumor rupture.

Neuroblastoma—Liu et al. [59] compared the performance of six machine learning algorithms to predict clinical outcomes in patients with neuroblastoma based on CT. An artificial neural

network performed well for most outcomes, such as mortality, presence of metastasis, and grade of neuroblastic differentiation (mean AUC,  $0.79\pm0.045$  [SD],  $0.83\pm0.034$ , and  $0.80\pm0.047$ , respectively) [59].

Veiga-Canuto et al. [43] tested an Al algorithm (nnU-Net) for automatically detecting and outlining neuroblastomas on MRI in children with neuroblastoma. Al achieved a success rate of 94% for tumor segmentation and was significantly faster than human-derived manual segmentations (mean time, 7.9 vs 124 seconds) [43].

Chen et al. [60] developed a CT-based machine learning model to extract radiomic features that may be linked to prediction of *MYCN* oncogene amplification in pediatric neuroblastoma. Random forest, SVM, and logistic regression models achieved similar high AUCs (0.851–0.909), significantly outperforming the Bayes model (AUC, 0.729) [60]. Feng et al. [61] built three machine learning models to predict the mitosis-karyorrhexis index (linked with response to chemotherapy) based on the FDG PET/CT and clinical data of 102 children with neuroblastoma. Future studies could correlate the mitosis-karyorrhexis index with MIBG scans.

# **Pulmonary Nodules**

For the detection of pulmonary nodules, some Al algorithms provide encouraging results [62], whereas others report inferior performance of algorithms trained on adult datasets for the detection of lung lesions on the chest radiographs [63] or CT scans [8, 9] of pediatric patients. Ni et al. [62] compared a deep CNN model (3087 detections and 278 misses) with the readings of junior physicians (2442 detections and 657 misses) for identifying pulmonary nodules in patients with osteosarcoma (109 nodules on 675 CT examinations). The deep CNN model achieved significantly higher sensitivity (92.3% vs 90.8%), specificity (55.2% vs 35.1%), and AUC (0.795 vs 0.687), indicating better detection and potentially greater accuracy. Additionally, the deep CNN model accelerated reading times compared with human interpretations. Shin et al. [63] used an adult-trained ResNet34 CNN for detection of lung lesions on 2273 pediatric chest radiographs and achieved an overall accuracy of only 87.5% (sensitivity, 67.2%; specificity, 91.1%). Exclusion of children younger than 2 years old and patients with cardiomegaly significantly improved the model's performance (accuracy, 96.9%; sensitivity, 86.4%; specificity, 97.9%). Hardie et al. [9] found that two CAD systems trained on adult data had lower sensitivity for detection of pulmonary nodules on chest CT in children (FlyerScan, 68.4%; Medical Open Network for Artificial Intelligence [MONAI], 53.1%) than in adults (FlyerScan, 83.9%; MONAI, 95.5%), highlighting the need for pediatric-specific training data for accurate diagnoses.

# Lymphoma

Although substantial attempts have been made to develop Al algorithms for detecting lymphoma on FDG PET/CT in adults, few applications to date have focused on pediatric patients. Wang et al. [64] pioneered Al-based detection of pediatric lymphoma using FDG PET and T1-weighted MRI. Their two-stage approach (50 training and 20 testing datasets) focused on identifying regions of high radiotracer uptake on PET (likely representing tumors), followed by false-positive removal. A CNN-based multimodal fusion method incorporating PET and MRI data addressed these false-positives, highlighting Al's potential for this application [64] (Fig. 5).

# **Brain Tumors**

Al has facilitated the diagnosis, segmentation, and detection of brain tumors [6, 65, 66]. In 2022, a systematic review reported that 22 studies on Al applications for pediatric brain tumor imaging identified tumor diagnosis as the dominant application, followed by tumor segmentation and detection [67]. Although five of six studies comparing Al to human experts favored Al for tumor diagnosis, none reported real-world clinical implementation of Al for this purpose [67].

Quon et al. [6] established a multiinstitutional MRI-based deep learning model for the detection and pathology classification of posterior fossa tumors in 600 pediatric patients. The model achieved an overall tumor detection and classification accuracy that was comparable with the performance of four board-certified radiologists [6] (Fig. 6).

Pisapia et al. [68] investigated an MRI-based machine learning model for predicting progression of optic pathway gliomas (OPGs). The model, trained on data from 19 progressing and 19 nonprogressing tumors, incorporated manual segmentation of optic nerves and diffusion tractography for optic radiations. Analyzing features derived from various MRI sequences, including diffusion tensor imaging (DTI), the model achieved high accuracy (86%), sensitivity (89%), and specificity (81%) for the prediction of OPG progression. A key predictive feature was the fractional anisotropy of the optic radiations on DTI (AUC, 0.83).

Zhou et al. [65] developed an automatic machine learning model based on MRI in 288 patients with posterior fossa tumors. A radiomic model of automatic machine learning with the tree-based pipeline optimization tool (TPOT) outperformed expert radiologists [65].

# **Conclusion and Future Directions**

Machine learning algorithms hold promise for pediatric oncology imaging, offering faster, cheaper, and potentially safer medical imaging methods for children with cancer. Challenges include the relative scarcity of pediatric cancer imaging data, inhomogeneity of multiinstitutional datasets, data distribution shifts, and lack of commercially available Al tools for pediatric imaging applications. Additional impediments to clinical translation include the diverse anatomies of growing children and the presence of nonstandardized imaging protocols.

Possible solutions include standardized imaging protocols, improved data harmonization techniques, formation of multiinstitutional consortia for data sharing, and generation of data repositories for transfer learning from datasets that include adult and pediatric cases. Increased efforts such as those of The Cancer Imaging Archive, the Children's Oncology Group, and the CBTN are required to ensure that data availability will not be a major hurdle for developing Al algorithms. Al algorithms trained on adult data alone may not perform well on children due to anatomic and physiologic differences, highlighting the need for transfer learning and the generation of pediatric-specific models. The initial results from machine learning-based radiomics models are encouraging, but prospective validation is necessary to ensure generalizability and real-world effectiveness. Additional opportunities include leveraging reconstruction algorithms to achieve accelerated low-dose imaging and automating the generation of metric-based staging and treatment monitoring scores.

Despite current limitations, AI offers a compelling path forward for pediatric oncology imaging. In contrast to human experts, who have a limited learning period and eventually retire, AI models hold the promise of perpetual learning. This continuous improvement across vast datasets, spanning generations of human expertise, can lead to significant performance gains over decades to come. As AI integration in clinical practice accelerates, robust safeguards are essential to ensure continued clinician autonomy in the event of system downtime. Transfer learning of adult-based AI models to pediatric cancers, multiinstitutional data sharing, and ethical data privacy practices for pediatric patients with rare cancers will be keys to achieving robust validation and unlocking AI's full potential for clinical translation and improved outcomes for these young patients.

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(Editorial Comment starts on next page)

# Editorial Comment: The Radiologist's Role in Artificial Intelligence for Pediatric Oncologic Imaging

Artificial intelligence (AI) is growing at an exponential pace and impacting our lives in myriad ways. From planning the most efficient morning commute to providing book recommendations for leisure reading, AI is designed to augment many of our experiences in the world. As radiologists, we have an opportunity to incorporate elements from the burgeoning AI revolution into the imaging-based care we provide for our patients.

Much like Al's application in our personal lives, the arc of radiology's potential Al application is broad. This article nicely summarizes the many roles Al may have in imaging pediatric patients with cancer [1]. Although some applications, such as Al-aided image reconstruction, are today's realities, other applications, such as imaging-based outcomes prognostication for children with cancer, are concepts we hope to see in practice in the not-too-distant future.

The road to achieving Al's full potential for application in pediatric oncologic imaging will not be without obstacles. Prior articles have discussed the unique impediments to progress in Al development for pediatric patients, yielding a dearth of FDA-approved algorithms and a health equity deficit for this vulnerable population [2, 3].

While reading this review article, I hope you ask: "What is my role in bringing AI to imaging care for pediatric patients?" There are many possible answers, but I believe two important ones are to be curious (i.e., stay informed about relevant AI technology and critically appraise how its application may allow improved care for your patients) and be an advocate (i.e., encourage and, where feasible, participate in the development of safe, effective, and accessible AI tools designed and developed for pediatric patients).

Al holds the promise of an exciting future in the care of pediatric patients with cancer—one where we as radiologists remain integral members of a health care team that offers efficient, effective, precise, and personalized care.

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