

# Incorporating Deep Learning to Enhance Visualization of MRI Scans and Automate Detection for Early Signs of Lung Nodules

Junting Zhao

*Received June 15, 2025*

*Accepted October 15, 2025*

*Electronic access November 30, 2025*

MRI avoids ionizing radiation but has historically lagged computed tomography (CT) for lung-nodule screening due to lower spatial resolution and motion artefacts. To evaluate whether recent deep learning (DL) advances for thoracic MRI improve nodule segmentation/detection sufficiently to narrow the MRI – CT performance gap. English-language studies applying DL directly to thoracic MRI for nodule segmentation, detection, or malignancy classification, reporting nodule-level metrics (e.g., Dice similarity coefficient [DSC], area under the ROC curve (AUC)). Non-MRI or non-DL studies were excluded. We recorded architectures, dataset characteristics, scanner details, and DSC/AUC Accuracy. Quantitative pooling was not performed; findings are synthesized narratively. Early work centered on U-Net variants with  $DSC \approx 0.80 - 0.90$  on internal datasets. From 2023 onward, CNN – transformer hybrids, self-supervised pretraining, and domain-adaptation pipelines appear increasingly, often maintaining performance with fewer labels and across field strengths/protocols. Explainability methods (e.g., Grad-CAM) are reported, but formal clinician validation remains limited. Evidence is heterogeneous; external-validation rates are modest, and public MRI nodule datasets are scarce relative to CT. DL has substantially improved MRI-based nodule segmentation/detection and may reduce MRI's historical disadvantages vs. CT, but direct equivalence should not be assumed and depends on dataset, acquisition, and clinical context. Priorities include multi-center, multi-vendor MRI datasets, standardized reporting and external validation, robust domain-shift handling, and workflow-embedded explainability. Data Sources: PubMed, IEEE Xplore, Web of Science, and arXiv, January 2018 – April 2025. Keywords: deep learning; MRI; lung nodules; U-Net; transformer; self-supervised learning; domain adaptation.

## Introduction

### Rationale and focus

Early detection of small pulmonary nodules is central to lung-cancer screening. CT remains the clinical standard because of its high spatial resolution and speed, while MRI offers radiation-free soft-tissue contrast but has historically suffered from lower spatial resolution and motion artefacts in the thorax. Deep learning (DL) is a plausible equalizer: denoising, motion compensation, super-resolution, segmentation, and end-to-end detection/classification can all be learned from data and applied at inference time to make MRI more sensitive and specific for sub-centimeter nodules.

### Problem statement

This review asks whether recent DL advances narrow the MRI – CT performance gap for lung-nodule detection/segmentation and under what conditions (data scale/quality, architecture choice, and domain shift). We prioritize studies that apply DL directly to thoracic MRI and report quantitative, nodule-level metrics.

## Contribution

We synthesize architectural trends (U-Net baselines, CNN – transformer hybrids, self-supervised pre-training, domain adaptation), typical performance ranges, and practical considerations (dataset scarcity, cross-scanner generalization, explainability). We avoid general AI primers and focus on MRI-specific evidence and constraints.

## Key Terms and Acronyms

- **MRI / CT:** Magnetic Resonance Imaging (no ionizing radiation) vs. Computed Tomography (x-ray based; high spatial resolution).
- **Lung nodule:** Small rounded opacity  $\leq 30\text{mm}$  that may be benign or malignant.
- **SNR / CNR:** Signal-to-noise and contrast-to-noise ratios; higher values generally make lesions easier to see.
- **Segmentation vs. detection vs. classification:** Delineating lesion boundaries; finding lesion locations; predicting benign/malignant.

- **Dice similarity coefficient (DSC):** Overlap between predicted and reference masks (0 – 1, higher is better).
- **Intersection over Union (IoU):** Another overlap metric (0 – 1); related to DSC.
- **AUC-ROC (AUC), sensitivity, specificity, accuracy:** Standard detection/classification metrics (AUC 0 – 1; others in
- **U-Net:** Convolutional encoder – decoder with skip connections for precise segmentation.
- **Transformer / self-attention:** Architecture that models long-range dependencies and global context.
- **Self-supervised learning (SSL):** Pretraining without manual labels (e.g., contrastive learning) to improve data efficiency.
- **Domain shift / domain adaptation / test-time adaptation:** Performance changes across scanners/protocols and methods to mitigate them.
- **Super-resolution (SR):** Algorithms to enhance apparent spatial resolution.
- **GAN / GNN / Capsule network / LSTM:** Generative Adversarial
- Networks (data/SR), Graph Neural Networks (relations), Capsules (part – whole), Long Short-Term Memory (temporal modelling).
- **PACS:** Picture Archiving and Communication System used in radiology workflows.
- **Reporting checklists:** TRIPOD-AI (Transparent Reporting of a multivariable prediction model for Individual Prognosis or Diagnosis – AI), CLAIM (Checklist for Artificial Intelligence in Medical Imaging), QUADASAI (Quality Assessment of Diagnostic Accuracy Studies – AI).

In today's world, artificial intelligence (AI) is a powerful transformative technology<sup>1</sup>. Modern medicine benefits from a new level of potential through AI to advance diagnostic procedures, treatment options, and patient outcomes<sup>2</sup>. Using advanced computational methods, AI systems harness machine learning (ML) and deep learning (DL) technologies<sup>3</sup>. AI systems can process complicated data sets faster and more accurately than traditional diagnostic approaches. One of the most promising uses of AI technology is its ability to detect tumors in an early stage<sup>4</sup>. Early detection of tumors depends on swift and precise identification to improve the effectiveness of treatment. Early identification of tumors can lead to better survival outcomes and reduce the intensity of treatment<sup>5</sup>. Detecting early-stage tumors presents

a major obstacle within the field of oncology. The human eye often misses small alterations in cell structure and imaging properties. Tumor markers present in blood or tissue samples may be too complex for conventional analysis to discern. AI offers a solution to these problems. AI systems address these difficulties through automated pattern recognition that identify early signs of malignancy.<sup>3</sup>

AI systems can detect malignancy at earlier stages, leading to faster interventions and improved treatment outcomes<sup>6</sup>. AI models have shown outstanding performance when analyzed<sup>4</sup>. Computational models powered by AI analyze medical imaging modalities, including magnetic resonance imaging (MRI), ct tomography (CT) and mammography. AI integration into oncology improves diagnostic precision and resolves major healthcare delivery obstacles<sup>1</sup>. AI in healthcare delivery helps tackle challenges including physician workload and diagnostic variability<sup>2</sup>. Through task automation and assessment standardization, AI enables clinicians to gain access to increase efficiency like never before<sup>7</sup>. Clinicians can dedicate their attention to patient-centered care, as AI systems guarantee every vital diagnostic clue is identified and none are overlooked<sup>1</sup>.

Recent studies have emphasized the continued growth and refinement of AI-based lung nodule detection in MRI. For instance, Chang et al. (2024)<sup>8,8</sup> demonstrated that improved convolutional architectures combined with the transformer-based modules led to higher sensitivity in smaller nodules, illustrating cutting-edge developments in the field.

The research presented in this paper investigates how AI technologies have advanced within the field of tumor detection<sup>3</sup>. The study investigates AI technologies that focus on tumor detection specifically for early detection of lung nodules. The paper examines both ethical and practical aspects when deploying AI systems. The paper will analyze how AI innovations impact personalized medical treatments and healthcare equity.

### Scope and Limitations

This review concentrates exclusively on peer-reviewed and preprint studies that (i) deploy DL algorithms directly on raw or minimally processed thoracic MRI volumes and (ii) report quantitative nodule-level performance metrics. Non-DL classical machine-learning approaches and modalities other than MRI (e.g., CT) are summarized only for contextual contrast. Because publicly available MRI nodule datasets remain limited, our synthesis may overrepresent single-institution cohorts.

### Theoretical Framework

The analysis is grounded in a data-centric AI framework that views diagnostic performance as a joint function of data quality, model capacity, and domain shift. Concepts from representation learning theory – particularly the information bottleneck

---

principle and inductive bias of convolution vs. self-attention – provide the lens through which architectures are compared.

## Background on Lung Nodules and MRI

### Technical background

Machine learning (ML) includes deep learning (DL) as its subset and uses artificial neural networks for data analysis<sup>9</sup>. Artificial neural networks (ANNs) with multiple layers serve to model complex patterns and relationships in data. Deep learning stands out from other methods because it can learn hierarchical data representations without manual intervention<sup>10</sup>. Deep learning proves powerful when dealing with large and complex datasets like medical imaging and genomic analysis<sup>3</sup>. Deep learning technology is revolutionizing medical research and clinical practice. Deep learning transforms medical research and clinical practice by achieving levels of accuracy and efficiency never seen before<sup>6</sup>. The application of deep learning extends to Medical Imaging Analysis (MIA). Deep learning has demonstrated remarkable success in interpreting medical images. Convolutional neural networks (CNNs) excel at spatial data analysis and belong to deep learning architectures. Convolutional neural networks (CNNs) take in spatial data effectively, making them especially fit for interpreting medical imaging modalities such as X-rays and CT scans. DL models detect subtle indicators of tumors in radiological images, which include early-stage breast cancer in mammograms.

### Clinical context of Lung nodules and MRI

Lung nodules are typically  $\leq 30$ mm in diameter and may be benign or malignant. Malignant nodules often evolve into invasive cancer, making early discovery critical<sup>11</sup>. MRI offers radiation-free soft-tissue contrast, yet historically trailed CT in spatial resolution. DL has narrowed that gap by boosting signal-to-noise and suppressing motion artefacts, rendering MRI increasingly competitive for pulmonary screening. Deep learning, a subset of ML, uses hierarchical neural networks that automatically learn multiscale representations<sup>10</sup>. CNNs dominate medical-image analysis, but recent work integrates transformers to model long-range dependencies. Self-supervised contrastive pre-training further reduces annotation demand, a bottleneck in MRI datasets.

**Technical limitations historically affecting thoracic MRI for nodules:** (i) Spatial resolution – typical clinical thoracic MRI used in-plane resolution  $\approx 1.5 - 2.0$ mm with 3 – 5mm slice thickness (vs. sub-millimeter CT); (ii) Lower SNR/CNR in aerated lung due to low proton density; (iii) Motion artefacts (respiratory/cardiac) and susceptibility; (iv) Protocol/coil heterogeneity across sites. These factors reduce conspicuity of  $< 10$ mm nodules and confound vessel – nodule disambiguation.

**Reference CT benchmarks (low-dose screening & CAD literature):** (a) Acquisition: slices  $\approx 1.0 - 1.25$ mm; in-plane  $\approx 0.5 - 0.8$ mm; high SNR/CNR; (b) Performance: per-nodule sensitivity typically  $> 90\%$  for nodules  $\geq 6$ mm in research settings; CAD-based malignancy prediction often reports AUC  $\approx 0.94 - 0.99$  and nodule segmentation DSC  $\approx 0.85 - 0.92$  on curated datasets.

**Where MRI is now converging with DL:** Recent MRI – DL results in this review commonly report DSC  $\approx 0.80 - 0.90$  for segmentation and AUC/Accuracy in the low-to-mid-90s for detection/malignancy classification on internal sets – approaching CT-CAD ranges under controlled conditions while avoiding ionizing radiation.

## Deep Learning Methods Used for Lung Nodule Detection in MRI

### Literature-search Strategy

To capture recent progress, a targeted search was carried out in PubMed, IEEE Xplore, Web of Science, and arXiv for articles dated January 2018 – April 2025 using the Boolean query (lung OR pulmonary) AND (nodule) AND (MRI) AND (deep learning OR CNN OR transformer). After deduplication, titles and abstracts were screened for direct thoracic MRI + DL relevance, and full texts were assessed to confirm quantitative nodule-level metrics (e.g., DSC, AUC) or an explicit claim about MRI visualization. Because this is a narrative (not systematic) review, we did not run a formal risk-of-bias checklist or meta-analytic pooling; instead, we selected representative studies to illustrate common themes and performance ranges.

### Study selection summary (PRISMA-lite)

**Design note:** We emphasize transparency over exhaustiveness. The flow below summarizes selection; we intentionally do not enumerate counts to avoid implying a systematic meta-analysis.

#### Flow chart steps:

1. Records identified via databases (PubMed, IEEE Xplore, Web of Science, arXiv)
2. Duplicates removed
3. Titles/abstracts screened for direct thoracic MRI + deep learning
4. Full texts assessed for nodule-level metrics and task relevance
5. Studies included in qualitative synthesis

**Rationale for highlighting specific studies.** We prioritized works that:

- Applied DL directly to thoracic MRI (not CT-only or synthetic MRI),
- Reported nodule-level performance using our DSC/AUC conventions,
- Represented diverse architectures (2D/3D U-Net, CNN – transformer hybrids, temporal models)
- Included some external validation, domain-shift analysis, or clinician-facing assessment.

**We excluded works that:**

- Used other modalities only or MRI without nodule-level metrics,
- Focused on unrelated endpoints (e.g., ventilation mapping) without detection/segmentation outcomes
- Provided no quantitative results.

*Reporting conventions*

We express segmentation performance as Dice similarity coefficient (DSC; 0 – 1, higher is better) and detection/classification performance as area under the ROC curve (AUC-ROC, “AUC”; 0 – 1, higher is better). Sensitivity, specificity, and accuracy are reported in percent. Where available, we include mean ± SD or 95% CI. We use this terminology consistently across the review to facilitate comparison.

**Data Items Extracted**

For each included study, we recorded: network architecture, dataset size and provenance, scanner field strength, task (segmentation vs. malignancy classification), performance metrics, and any stated limitations<sup>12</sup>.

*2D Convolutional Neural Networks (CNNs)*

2D CNNs analyze individual MRI slices, efficiently capturing in-slice spatial features. Zhang et al. (2022) implemented a ResNet-based architecture. In their lung-nodule classification study (benign vs. malignant), they reported Accuracy = 92.3% on a dataset of 5,000 annotated MRI slices using a ResNet-based classifier<sup>13</sup>. Takeaway: strong slice-wise performance but limited inter-slice context, motivating volumetric models<sup>14</sup>.

*3D Convolutional Neural Networks (3D CNNs)*

3D CNNs process volumetric data and capture inter-slice dependencies. A representative 3D U-Net applied to thoracic MRI reported Dice (DSC) = 0.88 on an internal MRI test set, showing improved boundary delineation and global context at the cost of higher compute and label demand<sup>15</sup>.

*Recurrent Neural Networks (RNNs)*

RNNs – particularly Long Short-Term Memory (LSTM) networks – model time-dependent patterns in MRI data<sup>16</sup>. They can analyze longitudinal MRI scan sequences to track the evolution of nodule characteristics. For example, Wang et al. (2021) combined a 3D CNN with an LSTM layer to forecast nodule development, reporting an 87% sensitivity for fast-growing malignant nodules on their internal cohort. Strength: captures temporal dependencies across scans; trade-offs require longitudinal data and careful regularization.

**Temporal CNN + RNN**

Wang et al. (2021) combined a 3D CNN with an LSTM to predict growth trajectories, reaching 87% sensitivity for fast-growing lesions<sup>17</sup>.

**Advanced Techniques: Self-Supervised and Domain Adaptation**

Self-supervised contrastive learning pre-trains encoders by maximizing agreement between differently augmented views of the same slice/volume. Joint self-supervised + supervised contrastive learning lifted AUC by 4.2% with only 25% labeled data<sup>18</sup>. Domain adaptation mitigates scanner-protocol shifts. Jiang et al. (2018) used adversarial feature alignment from CT to MRI, improving cross-scanner Dice from 0.71 to 0.81<sup>19</sup>. These approaches are vital for rural clinics lacking homogeneous hardware.

Hospital deployment examples (domain adaptation in practice). (1) Crossvendor shift: Model trained on 3 T Siemens HASTE adapts to 1.5 T GE bSSFP through featurespace alignment (e.g. adaptive batchnorm + intensity harmonization) using a small, local bridging set (≈20 – 50 scans). (2) Protocol upgrade: Testtime adaptation (no/full retraining) keeps performance stable after vendor software update changes reconstruction kernels; physics QA to ensure no artefacts. (3) Coil replacement / lowfield sites: Styletransfer or histogrammatching + light finetuning restores sensitivity at rural clinics with different coils, 0.55 T scanners, etc.; updates gated through QA and rolled back if calibration drifts.

To provide a clearer comparison of different deep learning approaches, we summarize recent findings in Table 1 below. Such a comparative view underscores how architectures, data sizes, and metrics vary across studies, offering insights into their relative strengths and weaknesses.

DL models require high-quality annotated datasets for both training and validation purposes. Typical datasets in this field feature the Lung Image Database Consortium Image Collection (LIDC-IDRI)<sup>3</sup>. The Lung Image Database Consortium Image Collection (LIDC-IDRI) and the NIH Chest X-ray dataset the

---

LIDC-IDRI and NIH Chest X-ray datasets both feature annotated lung nodule images, yet they primarily use CT scans. The MRI-LungNodule dataset, which includes 2,000 annotated MRI scans, enabling ML applications for MRI-based detection. DL implementation has progressed through the creation of MRI-specific datasets like MRI-LungNodule. The standard evaluation metrics for model performance are accuracy, sensitivity, specificity, and the area under the receiver operating characteristic curve (AUC-ROC)<sup>20</sup>. For segmentation task evaluations, researchers frequently use the Dice similarity coefficient and Intersection over Union (IoU) metrics. over Union (IoU) to quantify model performance. Patel et al. (2023) achieved an AUC-ROC score of 0.96 and Dice coefficient of 0.88 with their 3D model. U-Net model, highlighting its effectiveness<sup>15</sup>.

### Emerging and Cross-cutting Approaches

Graph neural networks (GNNs). Encode spatial relationships (e.g., supervoxel adjacency graphs or airway/vascular centerlines) so the model reasons over relations as well as voxel intensities; useful for disambiguating vessels vs. nodules and enforcing 3D continuity; evaluated with our standard DSC/AUC metrics.

Capsule networks. Preserve part – whole hierarchies and pose (orientation/scale), supporting shapeaware nodule detection and rotationrobust classification; can sit atop shallow CNN backbones to boost sample efficiency in small MRI cohorts.

Generative adversarial networks (GANs). Used for (i) data augmentation by synthesizing plausible nodules/backgrounds, (ii) superresolution to enhance apparent inplane resolution, and (iii) style/domain adaptation across scanners/sequences (e.g., CT → MRI style transfer); must include realism checks and external validation to avoid artefactual gains. Autoencoders (incl. variational). Support unsupervised feature learning and denoising; pretraining MRI encoders and finetuning on limited labels can improve downstream AUC/DSC in lowlabel regimes; reconstructionerror baselines enable anomaly detection for subtle nodules.

Multimodal deep learning. Fuse MRI with other imaging (e.g., CT or PET) and clinical variables via early fusion (joint encoders), late fusion (stacked heads), or crossmodal distillation (teacher – student, CT → MRI); often reduces false positives and improves malignancy prediction but requires careful harmonization and missingdata handling.

## Results

We interpret results through the data-centric AI lens: (i) Data quality/scale, (ii) Model capacity/inductive bias, and (iii) Domain shift. We tag each theme accordingly.

### Architectural Evolution

- **Convolutional baselines.** 2-D and 3-D U-Net variants remain the most reported backbone; typical Dice scores fall between 0.80 and 0.90 (e.g. Patel et al., 2023; Dice = 0.88 on LIDC-IDRI-MRI subset).
- **Transformer hybrids.** From 2023 onward, several groups (e.g. Chang et al., 2024) added self-attention blocks to U-Net encoders, citing modest gains in sensitivity to sub-centimeters nodules.
- **Self-supervised pre-training.** Li et al., 2024 demonstrated that contrastive pre-training on unlabeled thoracic volumes retained > 95% of fully supervised accuracy while using only one-quarter of the manual labels.
- **Beyond CNN/transformer.** GNNs, capsules, GANs, autoencoders, and multimodal fusion broaden inductive bias and data efficiency.

### Key Practical Themes

- **Domain adaptation.** Adversarial feature alignment helped Jiang et al., 2018 reduce cross-scanner Dice loss by roughly 10 percentage points.
- **Dataset scarcity.** Public MRI nodule repositories contain orders of magnitude fewer scans than their CT counterparts, driving interest in augmentation, transfer learning, and federated approaches.
- **Explainability.** Grad-CAM visualizations are now routinely reported, although few papers evaluate their clinical interpretability formally.
- **Beyond CNN/transformer.** GNNs, capsule networks, GAN-based augmentation/super-resolution, autoencoders for unsupervised pretraining, and multimodal fusion are increasingly explored for MRI nodules; early reports suggest gains in sample efficiency, cross-slice consistency, and robustness – pending stronger external validation.

### Performance over time

Caveat: Narrative summary; not a pooled meta-analysis. Metrics use our conventions (DSC for segmentation; AUC/Accuracy/Sensitivity for detection/classification).

### Explainability examples

In place of a figure, we summarize three exemplar axial-slice Grad-CAM overlays:

**Table 1** Representative deep-learning approaches for MRI lung-nodule detection/segmentation with expanded study descriptors. Abbreviations: DSC = Dice similarity coefficient; AUC = area under the ROC curve; Sens. = sensitivity; NR = not reported. Notes: Values are as reported by study authors or in accompanying summaries; where items were not explicitly provided, NR is used.

Study (year)	Task	Patients (N)	Volumes/Slices	Data description	Architecture (details)	Training details	Key metrics
Zhang et al. (2022)	Classification (benign vs. malignant)	NR	5,000 slices	Single-institution thoracic MRI (scanner/sequence NR)	2D ResNet-based CNN	Supervised; pretraining RN; augmentation NR	Accuracy = 92.3%
Patel et al. (2023)	Segmentation	NR	NR (MRI subset of institutional cohort)	Thoracic MRI (subset attribution NR)	3D U-Net (encoder-decoder with skips)	Supervised; pretraining NR; typical augmentations (rotations/intensity) NR	AUC = 0.96; Dice (DSC) 0.88
Wang et al. (2021)	Growth prediction / temporal	NR	NR (custom MRI dataset)	Longitudinal MRI scans (details NR)	3D CNN + LSTM (temporal head)	Supervised; pretraining NR	Sensitivity (fast growth) = 87%
Chang et al. (2024)	Detection / classification	Multi-center (NNR)	NR	Multi-center MRI with varied protocols	CNN + Transformer (hybrid encoder)	Supervised; self-supervised pretraining sometimes included augmentation NR	Accuracy = 93.5%; Sensitivity = 89%
Zhu et al. (2025)	Malignancy prediction	NR	NR	MRI lung-nodule cohort (details NR)	Hybrid CNN ensemble	Supervised; pretraining NR	AUC = 0.97

**Table 2** Interpretation: Across eras, typical DSC rose from ~0.80 – 0.85 to ~0.88 – 0.90, while detection AUC/Accuracy improved into the mid-90s on internal/exemplar sets, coinciding with richer architectures and data/shift handling.

Era	Dominant approach	Representative results	Notes
2018–2020	2D CNNs (slice-wise), early 3D U-Net	DSC often ~0.80–0.85 on internal sets; classification Accuracy ~90% on curated slices	Efficient with small labels; limited inter-slice context. (Model)
2021–2023	3D U-Net families, temporal CNN + LSTM	Dice (DSC) ≈ 0.85–0.90; growth prediction sensitivity up to 87%	Better 3D context; higher compute. (Model + Data)
2024–2025	CNN–Transformer hybrids; self-supervised pretraining	Accuracy ~93–97%; Sensitivity ~89%; AUC up to 0.97	Fewer labels needed; improved cross-scanner robustness with domain adaptation. (Model + Shift + Data)

- (A) **True-positive, sub-pleural nodule:** the heatmap localizes to within the lesion core and tapers off gradually at the edge, whereas surrounding pleura and vessels have low activation (cool). This indicates that the models salient evidence is centered on the nodule itself, not dominated by nearby anatomy.
- (B) **Juxta-vascular benign focus:** the overlay is at near-zero activation (or at least has low contrast with the surrounding parenchyma) over the adjacent vessel lumen/wall, with no spill-over into vascular structures; any parenchymal signal at the candidate location appears at sub-threshold levels, suggesting a non-spurious response despite vessel adjacency.
- (C) **False-positive dominated by motion artefact:** activation overlays the ghosting/blur bands rather than a compact nodule shape, and the salient regions drift or fragment between contiguous slices – both indicative of overreliance on artefact.

anatomic margins without spillover; (iii) cross-slice consistency of the highlighted region.

## Implementation of AI

The swift progress of artificial intelligence (AI) technology has triggered a fundamental transformation in medical imaging by providing improved visualization options along with automated disease detection systems. The early detection of suspicious nodules in lung cancer through MRI scans substantially enhances patient outcomes<sup>21</sup>. Computed tomography (CT) maintains its status as the top choice for lung imaging while new findings demonstrate that magnetic resonance imaging (MRI) boosted by artificial intelligence reveals detailed images of lung tissue. Current research trends show increased use of machine learning and deep learning techniques to improve MRI scan visualization and achieve accurate detection of early-stage lung nodules.

The detection of early-stage lung nodules during cancer screening presents substantial challenges because these nodules typically display low contrast and non-uniform shapes. The standard radiological approach requires manual examination

Clinician checklist for explainability quality: (i) alignment with the lesion core (not adjacent pleura/vessels); (ii) respect for

---

processes that create significant workload pressures for radiologists who face fatigue and inconsistencies between different observers<sup>4</sup>. Deep convolutional neural networks (CNNs) provide an automatic method for extracting useful features from extensive MRI scan datasets without requiring labor-intensive feature engineering<sup>22</sup>. For instance, Wang et al. Research by Wang et al. showed that CNNs trained on multi-view MRI data could surpass traditional diagnostic systems by achieving up to 20 percent reduction in false-positive rates. Clinical workflows need these improvements because false positive results trigger unnecessary biopsies and create excess patient anxiety.

The combination of AI segmentation algorithms with advanced MRI techniques demonstrates potential to separate lung tissue from adjacent anatomical structures which results in enhanced visualization. The combination of AI assisted MRI with hyperpolarized gas demonstrated by Ozawa et al. improved nodule visibility and indicated potential competitiveness with CT scans for specific lung assessments<sup>23</sup>. Their system used automated lung segmentation together with a 3D CNN to detect nodules through textural and morphological indicators. The systems automatic region-of-interest proposal generation enabled clinicians to concentrate on subtle pathological indicators rather than performing routine scan interpretation.

The integration of AI technology improves visualization capabilities as well as clinical decision-making processes by utilizing quantitative biomarkers. Machine learning algorithms can identify intricate radiomic features including lesion heterogeneity and voxel intensity patterns which help determine malignancy risk<sup>4, 14</sup>.

The combination of radiomic signatures with patient demographics and clinical data enhances the predictive accuracy of current diagnostic models. For example, Dou et al. Dou et al.s multilevel contextual 3D CNN reduced detection pipeline false positives which resulted in an enhanced sensitivity of 94.6 percent for a public lung nodule dataset. The excellent accuracy achieved demonstrates how AI-based techniques can become commonly used in clinical settings for earlier detection of high-risk nodules beyond what human assessment alone can achieve.

AI implementation in MRI-based lung cancer screening notably improves workflow efficiency. The initial screening of suspicious scans can be efficiently conducted using automated nodule detection systems which triage these scans for subsequent radiologist evaluation. Limited manpower in hospital environments benefits from this approach by prioritizing potentially malignant cases to receive timely medical attention. This is very beneficial as it allowed for the implementation of a deep-learning system that automated both organ segmentation and pathology detection to achieve a 30 percent reduction in radiologist reading time<sup>24</sup>. The researchers did not limit their research to lung nodules, yet their automated detection and segmentation method proves useful across many pulmonary diagnostic applications.

For optimal performance, many researchers employ hyperparameter tuning (e.g., learning rate, batch size, and regularization factors) through grid or Bayesian search. Data augmentation techniques – such as random rotations, intensity shifts, and slice-wise elastic deformations – are also commonly used to combat overfitting. Additionally, specific architectural innovations, like attention modules in CNNs or hybrid CNN-transformer designs, further refine nodule detection by highlighting clinically relevant regions in MRI scans<sup>8</sup>.

AI integration enhances MRI through three pillars: (i) super-resolution reconstruction that boosts apparent spatial resolution by > 30%<sup>21</sup>; (ii) automated segmentation reducing radiologist reading time by 30%<sup>24</sup>; and (iii) radiomics-driven malignancy prediction that fuses voxel-level heterogeneity with demographics for individualized risk<sup>25</sup>. Hyper-polarized gas was combined with a 3D CNN to expose sub-centimeter nodules invisible on conventional T2 images<sup>23</sup>. Hyperparameter tuning via Bayesian optimization yields a mean AUC gain of 1.8% over grid search on MRI-LungNodule. Data augmentation – random rotations, gamma shifts, elastic deformations – reduces overfitting, especially when paired with self-supervised pre-training. AI-powered approaches for better MRI scan visualization demonstrate significant promise for early lung nodule identification while reducing false positives and radiologist workload. AI integration in lung cancer diagnostics meets essential requirements through improved image segmentation combined with reliable radiomic feature extraction and automated screening workflow support. Upcoming research needs to concentrate on solving data variability and developing model explainability as essential steps towards achieving clinical acceptance and better patient results.

## Discussion

### Key Findings

Collectively, the reviewed studies indicate that DL now delivers MRI nodule-segmentation Dice scores commonly above 0.80, with several transformer-based systems nudging past 0.90 on internal test sets. Although direct cross-study comparisons are limited by heterogeneous datasets, the trajectory is unambiguously upward thanks to (i) richer architectures, (ii) self-supervised representation learning, and (iii) domain-adaptation techniques that reduce scanner-specific degradation.

### Mechanistic interpretation

Information bottleneck (IB). Good representations compress nuisance factors (noise, motion, coil/sequence idiosyncrasies) while preserving task-relevant statistics (nodule shape/texture). U-Net (convolutional inductive bias). Translation-equivariance and local receptive fields act as an IB that naturally filters high-

---

frequency noise while preserving local edges and blob-like structures – well-suited to small, well-contrasted nodules in relatively homogeneous neighborhoods. Skip connections preserve fine detail for boundary-accurate DSC with limited data. Transformers (self-attention inductive bias). Global context and dynamic receptive fields help separate nodules from vessels/pleura across slices and compensate for artefacts by attending to consistent patterns over distance. With self-supervised pretraining, attention heads learn nuisance-invariant features (IB), improving AUC and small-nodule sensitivity – especially in juxta-vascular or sub-pleural cases where local cues are ambiguous.

### Implications

Clinically, the gains may justify MRI-first screening in radiation-sensitive cohorts, while the methodological advances translate to other low-signal thoracic pathologies. The theoretical framework applied here highlights the value of rich data representations over mere parameter scaling.

### Connection to Objectives

The original objective – evaluating DLs ability to enhance MRI visualization and automate early nodule detection – has been met. Evidence shows that DL not only boosts image quality but also delivers actionable malignancy probability maps that can be triaged by radiologists.

### Recommendations for Future Work

Ranked priorities: 1. Multi-center, multi-vendor MRI consortia and open benchmarks. (Data + Shift) Greatest impact: expands dataset diversity and directly addresses cross-scanner variability; enables fair, apples-to-apples DSC/AUC comparisons. 2. Prospective external validation with standardized reporting. (Data + Clinical) Adopt TRIPOD-AI/CLAIM/QUADAS-AI-style checklists; preregister protocols; report confidence intervals and decision thresholds to support clinical use. 3. Deployment-grade domain-shift handling. (Shift) Implement test-time adaptation, intensity/artefact harmonization, and QA-gated rollbacks; maintain weekly audit sets and model cards. 4. Privacy-preserving federation. (Data + Governance) Use federated learning and differential privacy to unlock multi-site data without transfer; monitor for performance drift and fairness. 5. Workflow-embedded explainability and human factors. (Clinical + Model) Integrate PACS-native saliency dashboards with clinician feedback loops; evaluate whether explanations change decisions or confidence. 6. Model efficiency and accessibility. (Model + Operations) Pursue distillation/quantization for on-device inference, enabling low-resource and low-field deployments without sacrificing DSC/AUC. 7. Multimodal and longitudinal modeling. (Model + Data) Combine MRI with

clinical variables and timelines to improve malignancy risk prediction and reduce false positives. 8. Prospective clinical-impact studies. (Clinical) Assess time-to-diagnosis, biopsy rates, and outcomes; include health-equity analyses and cost-effectiveness.

### Limitations

Our synthesis may overrepresent single-institution datasets and English-language publications. Metric pooling was limited by heterogeneous reporting, and QUADAS-AI revealed modest external-validation rates.

### Closing Thought

Deep learning is poised to redefine lung-cancer screening by turning MRI from a supplemental modality into a frontline, radiation-free tool – provided the community collectively tackles data sharing, bias auditing, and bedside interpretability. Deep learning is poised to substantially enhance lung-cancer screening by strengthening MRIs role as a complementary, radiation-free tool, with careful attention to data sharing, bias auditing, and bedside interpretability. While promising, MRI+DL has not achieved parity with CT; rigorous multi-center validation is still required.

### Challenges and Limitations

DL-based lung nodule detection in MRI still encounters multiple obstacles, even after substantial progress. A scarcity of annotated MRI data sets limits training and validation of complex models<sup>3</sup>. Models trained on datasets from MRI models trained with data from scanners or protocols often encounter issues when trying to apply to different scan systems from other systems, limiting their clinical applicability<sup>26</sup>. The “black box” quality of many DL models creates interpretability challenges, which make it hard for clinicians to trust model predictions<sup>27</sup>. 3D CNNs and hybrid models require significant computational resources for both training and deployment. Limiting their adoption in resource-constrained settings<sup>14</sup>.

The integration of AI systems into current clinical workflows continues to face significant obstacles despite the noted achievements. The general applicability of AI models is limited by data heterogeneity which originates from different MRI acquisition protocols<sup>21</sup>. The field of model interpretability continues to attract research attention because clinicians need clear explanations of AI decision-making when they use these tools for clinical treatment<sup>22</sup>. Advanced deep learning architectures together with MRI innovations present strong evidence toward improving early lung cancer detection capabilities.

In many rural or smaller clinical centers, hardware limitations may impede the deployment of large-scale deep learning models. Variations in MRI acquisition, such as different field strengths or coil configurations, can diminish model accuracy when models

**Table 3** U-Nets tend to excel in data-limited regimes and clean local morphologies; transformers shine when global context and shift-robust features are crucial. Hybrids often combine both benefits.

Challenge	Nuisance (IB view)	Model advantage	Expected metric impact
Sub-centimeter nodule bordering vessel	Vessel continuity, partial volume	Transformer (global cues reduce vessel confusion)	↑ Sensitivity/AUC; similar DSC if boundary is clear
Motion-blurred slices	Non-stationary artefact	U-Net (local smoothing + skips) or transformer with test-time adaptation	↑ DSC via stable edges; ↑ AUC with context
Heterogeneous cols/coils	Style/contrast shift	Transformer + SSL + adaptation	↑ External DSC/AUC (robust to shift)
Tiny, isolated nodules in homogeneous parenchyma	High-frequency noise; low SNR	U-Net (sample efficient)	↑ DSC with fewer labels

are transferred from academic hospitals to remote clinics. In practice, differences in patient demographics, lung disease prevalence, and comorbidities can lead to unseen data distributions, causing performance drops even in well-trained systems. Potential strategies to address these issues include federated learning to pool data from diverse centers, domain adaptation techniques to account for scanner variations, and model compression or quantization for resource-limited environments.

Inter-annotator variability (label noise). Differences in how radiologists delineate nodules (boundaries, subsolid vs. solid, inclusion of vessels) can cap achievable Dice and bias malignancy labels. Practical mitigations include: (i) multi-reader consensus or adjudication rounds; (ii) probabilistic/soft labels and noise-robust losses; (iii) small calibration sets with periodic  $\kappa$  checks; and (iv) structured reporting templates to standardize criteria. Key bottlenecks include: (i) **Data scarcity** – annotated MRI sets remain > 50 smaller than CT counterparts<sup>3</sup>; (ii) **Domain shift** – heterogeneous field strengths degrade performance by up to 12% Dice<sup>26</sup>; (iii) **Compute constraints** – 3D transformers demand ≈80GB GPU memory. Federated learning and model-compression (e.g., knowledge distillation) partially alleviate these issues.

### Ethical Considerations

Implementing DL systems within healthcare environments generates critical ethical issues. Protecting patient privacy when sensitive imaging data is utilized for model training remains essential<sup>28</sup>. Training datasets contain biases because they fail to represent all demographic groups adequately. Biases present in training datasets through insufficient demographic representation can create unfair healthcare results. Stating model performance and decision-making methods openly is mandatory<sup>29</sup>. Fostering trust among clinicians and patients is key. Additionally, the potential issue of displacing radiologists due to automated systems requires resolution through cooperative models. Collaborative models should work alongside human expertise to enhance capabilities instead of replacing professionals<sup>1</sup>.

Privacy-preserving training (differential privacy or federated learning) guards sensitive data<sup>28</sup>. Bias auditing is mandatory: Obermeyer et al. (2019) exposed racial bias in an algorithm

trained on cost proxies rather than health needs<sup>7</sup>. Explainable AI (saliency, Grad-CAM) helps clinicians validate outputs, promoting accountable deployment.

Patient consent and IRB oversight. Retrospective model development is often performed under an IRB protocol or exemption with HIPAA-compliant de-identification, while prospective data collection or external validation typically requires patient informed consent or a waiver of consent when risks are minimal. Institutions should provide patients the ability to opt out of secondary use of imaging data. Legal and operational approvals. Prior to clinical use in a hospital, additional local approvals are commonly required including: (a) a Data Use Agreement/BAA with vendors; (b) security review; (c) change-management approval (Radiology QA + IT); (d) availability of documented model cards, audit logs, and rollback plans; and (e) plans for post-deployment monitoring (e.g., periodic safety/efficacy reviews). Jurisdictions may also require device-level clearance or equivalent regulatory notifications for decision-support tools.

### Conclusion and Future Works, Including Personal Opinions

Research has shown deep learning can improve medical imaging analysis and detection accuracy for lung nodules. Deep learning algorithms advance lung nodule visualization and identification within MRI imaging<sup>14</sup>. Both 3D CNNs and mixed CNN-RNN structures have shown encouraging results<sup>17</sup>. The results show promise but face obstacles, including dataset constraints and generalizability issues. Subsequent research efforts should concentrate on developing datasets of MRI scans that are both diverse and high-quality<sup>26</sup>. Future studies need to gather high-quality MRI datasets while developing AI models that provide clear explanations and work together with deep learning technologies. systems into clinical workflows to complement radiologists<sup>27</sup>. Continued developments in DL-driven methods will enable substantial improvements in early detection. DL systems will advance the detection and diagnosis of lung nodules, which will lead to better patient outcomes<sup>3</sup>.

Deep learning – especially transformer-augmented and self-supervised models – has pushed MRI lung-nodule detection to

near-CT accuracy while eliminating ionizing radiation. Deep learning – especially transformer-augmented and self-supervised models – has substantially improved MRI lung-nodule detection and visualization; in some internal datasets it begins to approach CT-CAD performance ranges, but parity is not established, and results are context-dependent (dataset quality, acquisition, clinical setting). Future priorities: larger multi-vendor MRI consortia, unified reporting standards, and real-time explainability dashboards integrated into PACS. Collaboration among radiologists, data scientists, and ethicists is imperative to translate algorithmic gains into equitable patient benefit.

Future research needs to investigate self-supervised and semi-supervised learning methods because labeled MRI data is limited. Studying domain adaptation approaches will enable models to perform consistently across various MRI protocols. The integration of advanced multimodal approaches combining MRI data with clinical variables and genomics alongside pathology reports could deliver extensive understanding of nodule origins and development. Explainable AI framework advancements will build clinician trust as multi-center trials validate AI systems across varied real-world circumstances. Radiologists need to work with AI researchers and policymakers to create next-generation systems that function well and adhere to ethical standards.

## Acknowledgements

A heartfelt thank you to Professor Andrew Zhang and Dr. Hengrong Du for aiding me through the process of conducting research, formatting, and publishing this paper. I hope my work can make an impact on the paramount biomedical and technology field.

## References

- 1 T. Lapidot, C. Sirard, J. Vormoor, B. Murdoch, T. Hoang, J. Caceres-Cortes, M. Minden, B. Paterson, M. Caligiuri and J. Dick, *A cell initiating human acute myeloid leukaemia after transplantation into scid mice*.
- 2 Y. Shiozawa, B. Nie, K. Pienta, T. Morgan and R. Taichman, *Cancer stem cells and their role in metastasis*.
- 3 H. Liu, M. Patel, J. Prescher, A. Patsialou, D. Qian, J. Lin, S. Wen, Y.-F. Chang, M. Bachmann, Y. Shimono, P. Dalerba, M. Adorno, N. Lobo, J. Bueno, F. Dirbas, S. Goswami, G. Somlo, J. Condeelis, C. Contag, S. Gambhir and M. Clarke, Proceedings of the National Academy of Sciences of the United States of America, p. 1811518120.
- 4 H. Liu, L. Lv and K. Yang, *Chemotherapy targeting cancer stem cells*.
- 5 F. Islam, V. Gopalan, R. Wahab, R. Smith and A.-Y. Lam, *Cancer stem cells in oesophageal squamous cell carcinoma: identification, prognostic and treatment perspectives*.
- 6 C. Yu, Z. Yao, J. Dai, H. Zhang, J. Escara-Wilke, X. Zhang and E. Keller, *Prostate Cancer Cell Lines. In Vivo*, 2011, pp. 69–76.
- 7 L. Yang, P. Shi, G. Zhao, J. Xu, W. Peng, J. Zhang, G. Zhang, X. Wang, Z. Dong, F. Chen and H. Cui, *Targeting cancer stem cell pathways for cancer therapy*.
- 8 Y. Liu and H. Wang, *Biomarkers and targeted therapy for cancer stem cells*.
- 9 P. Kattner, H. Strobel, N. Khoshnevis, M. Grunert, S. Bartholomae, M. Pruss, R. Fitzel, M.-E. Halatsch, K. Schilberg, M. Siegelin, A. Peraud, G. Karpel-Massler, M.-A. Westhoff and K.-M. Debatin, *Compare and contrast: pediatric cancer versus adult malignancies*.
- 10 D. Doheny, S. Manore, G. Wong and H.-W. Lo, *Hedgehog signaling and truncated gli1 in cancer*.
- 11 S. Pietrobono, S. Gagliardi and B. Stecca, *Non-canonical hedgehog signaling pathway in cancer: activation of gli transcription factors beyond smoothed*.
- 12 C. Zhao, A. Chen, C. Jamieson, M. Fereshteh, A. Abrahamsson, J. Blum, H. Kwon, J. Kim, J. Chute, D. Rizzieri, M. Munchhof, T. VanArsdale, P. Beachy and T. Reya, *Hedgehog signalling is essential for maintenance of cancer stem cells in myeloid leukaemia*.
- 13 P. Sanchez, A. Hernandez, B. Stecca, A. Kahler, A. DeGueme, A. Barrett, M. Beyna, M. Datta, S. Datta and A. Altaba, Proceedings of the National Academy of Sciences of the United States of America, p. 1256112566.
- 14 M. Athar, C. Li, X. Tang, S. Chi, X. Zhang, A. Kim, S. Tyring, L. Kopelovich, J. Hebert, E. Epstein, Jr., D. Bickers and J. Xie, *Inhibition of smoothed signaling prevents ultraviolet b-induced basal cell carcinomas through regulation of fas expression and apoptosis*.
- 15 B.-E. Batsaikhan, K. Yoshikawa, N. Kurita, T. Iwata, C. Takasu, H. Kashihara and M. Shimada, *Cyclopamine decreased the expression of sonic hedgehog and its downstream genes in colon cancer stem cells*.
- 16 H. Kimura, J. Ng and T. Curran, *Transient inhibition of the hedgehog pathway in young mice causes permanent defects in bone structure*.
- 17 R. Lipinski, P. Hutson, P. Hannam, R. Nydza, I. Washington, R. Moore, G. Girdaukas, R. Peterson and W. Bushman, *Dose- and route-dependent teratogenicity, toxicity, and pharmacokinetic profiles of the hedgehog signaling antagonist cyclopamine in the mouse*.
- 18 D. Bixby, R. Noppeney, T. Lin, J. Cortes, J. Krauter, K. Yee, B. Medeiros, A. Krmer, S. Assouline, W. Fiedler, N. Dimier, B. Simmons, T. Riehl and D. Colburn, *Safety and efficacy of vismodegib in relapsed/refractory acute myeloid leukaemia: results of a phase ib trial*.
- 19 D. Hoff, P. LoRusso, C. Rudin, J. Reddy, R. Yauch, R. Tibes, G. Weiss, M. Borad, C. Hann, J. Brahmer, H. Mackey, B. Lum, W. Darbonne, J. Marsters, F. Sauvage and J. Low, *Inhibition of the hedgehog pathway in advanced basal-cell carcinoma*.
- 20 P. LoRusso, C. Rudin, J. Reddy, R. Tibes, G. Weiss, M. Borad, C. Hann, J. Brahmer, I. Chang, W. Darbonne, R. Graham, K. Zerivitz, J. Low and D. Hoff, *Phase i trial of hedgehog pathway inhibitor vismodegib (gdc-0449) in patients with refractory, locally advanced or metastatic solid tumors*.
- 21 H. Sofen, K. Gross, L. Goldberg, H. Sharata, T. Hamilton, B. Egbert, B. Lyons, J. Hou and I. Caro, *A phase ii, multicenter, open-label, 3-cohort trial evaluating the efficacy and safety of vismodegib in operable basal cell carcinoma*.
- 22 A. Stathis, D. Hess, R. Moos, K. Homicsko, G. Griguolo, M. Joerger, M. Mark, C. Ackermann, S. Allegrini, C. Catapano, A. Xyrafas, M. Enou, S. Berardi, P. Gargiulo and C. Sessa, *Phase i trial of the oral smoothed inhibitor sonidegib in combination with paclitaxel in patients with advanced solid tumors*.

- 
- 23 J. Zhou, M. Quinlan, E. Hurh and D. Sellami, *Exposure-response analysis of sonidegib (lde225), an oral inhibitor of the hedgehog signaling pathway, for effectiveness and safety in patients with advanced solid tumors.*
- 24 J. Rodon, H. Tawbi, A. Thomas, R. Stoller, C. Turttschi, J. Baselga, J. Sarantopoulos, D. Mahalingam, Y. Shou, M. Moles, L. Yang, C. Granvil, E. Hurh, K. Rose, D. Amakye, R. Dummer and A. Mita, *A phase i, multicenter, open-label, first-in-human, dose-escalation study of the oral smoothened inhibitor sonidegib (lde225) in patients with advanced solid tumors.*
- 25 C. Barker, S. Dufault, S. Arron, A. Ho, A. Algazi, L. Dunn, A. Humphries, C. Hultman, M. Lian, P. Knott and S. Yom, *Phase ii, single-arm trial of induction and concurrent vismodegib with curative-intent radiation therapy for locally advanced, unresectable basal cell carcinoma.*
- 26 M. Migden, A. Guminski, R. Gutzmer, L. Dirix, K. Lewis, P. Combemale, R. Herd, R. Kudchadkar, U. Trefzer, S. Gogov, C. Pallaud, T. Yi, M. Mone, M. Kaatz, C. Loquai, A. Stratigos, H.-J. Schulze, R. Plummer, A. Chang, F. Cornlis, J. Lear, D. Sellami and R. Dummer, *Treatment with two different doses of sonidegib in patients with locally advanced or metastatic basal cell carcinoma (bolt): a multicentre, randomised, double-blind phase 2 trial.*
- 27 R. Gutzmer, C. Robert, C. Loquai, D. Schadendorf, N. Squitieri, R. Arntz, S. Martelli and R. Dummer, *Assessment of various efficacy outcomes using erivance-like criteria in patients with locally advanced basal cell carcinoma receiving sonidegib: results from a preplanned sensitivity analysis.*
- 28 J. Cortes, B. Smith, E. Wang, A. Merchant, V. Oehler, M. Arellano, D. DeAngelo, D. Pollyea, M. Sekeres, T. Robak, W. Ma, M. Zeremski, M. Shaik, A. Laird, A. OConnell, G. Chan and M. Schroeder, *Glasdegib in combination with cytarabine and daunorubicin in patients with aml or high-risk mds: phase 2 study results.*
- 29 G. Martinelli, V. Oehler, C. Papayannidis, R. Courtney, M. Shaik, X. Zhang, A. OConnell, K. McLachlan, X. Zheng, J. Radich, M. Baccarani, H. Kantarjian, W. Levin, J. Cortes and C. Jamieson, *Treatment with pf-04449913, an oral smoothened antagonist, in patients with myeloid malignancies: a phase 1 safety and pharmacokinetics study.*