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# A Comparison of Natural Language Processing Methods for the Classification of Lumbar Spine Imaging Findings Related to Lower Back Pain

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# Abstract

**Rationale and Objectives:** The use of natural language processing (NLP) in radiology provides an opportunity to assist clinicians with phenotyping patients. However, the performance and generalizability of NLP across healthcare systems is uncertain. We assessed the performance within and generalizability across four healthcare systems of different NLP representational

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CODE AVAILABILITY

https://github.com/chethanjjj/LireNLPSystem

SUPPLEMENTARY MATERIALS

Supplementary material associated with this article can be found in the online version at doi:10.1016/j.acra.2021.09.005.

methods, coupled with elastic-net logistic regression to classify lower back pain-related findings from lumbar spine imaging reports.

**Materials and Methods:** We used a dataset of 871 X-ray and magnetic resonance imaging reports sampled from a prospective study across four healthcare systems between October 2013 and September 2016. We annotated each report for 26 findings potentially related to lower back pain. Our framework applied four different NLP methods to convert text into feature sets (representations). For each representation, our framework used an elastic-net logistic regression model for each finding (i.e., 26 binary or "one-vs.-rest" classification models). For performance evaluation, we split data into training (80%, 697/871) and testing (20%, 174/871). In the training set, we used cross validation to identify the optimal hyperparameter value and then retrained on the full training set. We then assessed performance based on area under the curve (AUC) for the test set. We repeated this process 25 times with each repeat using a different random train/test split of the data, so that we could estimate 95% confidence intervals, and assess significant difference in performance between representations. For generalizability evaluation, we trained models on data from three healthcare systems with cross validation and then tested on the fourth. We repeated this process for each system, then calculated mean and standard deviation (SD) of AUC across the systems.

**Results:** For individual representations, n-grams had the best average performance across all 26 findings (AUC: 0.960). For generalizability, document embeddings had the most consistent average performance across systems (SD: 0.010). Out of these 26 findings, we considered eight as potentially clinically important (*any stenosis, central stenosis, lateral stenosis, foraminal stenosis, disc extrusion, nerve root displacement compression, endplate edema, and listhesis grade 2*) since they have a relatively greater association with a history of lower back pain compared to the remaining 18 classes. We found a similar pattern for these eight in which n-grams and document embeddings had the best average performance (AUC: 0.954) and generalizability (SD: 0.007), respectively.

**Conclusion:** Based on performance assessment, we found that n-grams is the preferred method if classifier development and deployment occur at the same system. However, for deployment at multiple systems outside of the development system, or potentially if physician behavior changes within a system, one should consider document embeddings since embeddings appear to have the most consistent performance across systems.

## Keywords

Natural language processing; Lower back pain; Document embeddings; Evaluation; Lumbar spine diagnostic imaging

# INTRODUCTION

Lower back pain (LBP) is a common condition, in which patients typically exhibit heterogeneous anatomic phenotypes and undergo a variety of treatments (1–3). LBP patients frequently receive spinal imaging, and findings identified in the resulting radiology reports are expected to help with phenotyping and decision-making (3). However, the association between many findings and LBP is uncertain, because findings can be present in both symptomatic and asymptomatic patients (4). As a result, patients with common aging-related

findings may be recommended for LBP-related treatments that are not etiologically linked to their pain. To address this uncertainty, cohort studies and pragmatic trials have investigated patterns of care among patients with LBP and sought to explore subgroups of patients based on the presence of potentially clinically important findings (4–6). To address the interpretation of radiology findings, the Lumbar Imaging with Reporting of Epidemiology (LIRE) study assessed the effectiveness of including benchmark prevalences in the asymptomatic population for findings found in radiology reports for patients who received a diagnostic imaging test of the lumbar spine to reduce subsequent spine-related interventions at four healthcare systems: Kaiser Permanente of Washington, Kaiser Permanente of Northern California, Henry Ford Health System, and Mayo Clinic Health System (7). To further assist research investigating the relationship between findings and LBP, the accurate extraction of findings from large patient groups is needed. However, manual annotation is time-consuming. Natural Language Processing (NLP)-based classification pipelines offer an automated alternative to identify key findings in radiology reports (3).

An NLP-based classification pipeline is composed of two parts: NLP methods that extract features from free-text data and convert them to a structured format (or representation) and the machine learning (ML) model that uses these representations for classification. Text conversion or feature generation can be performed using methods that range from relatively simple domain-dependent and highly manual, to sophisticated data-driven scalable strategies (8–11). Taskspecific rule-based methods identify terms in the free-text that are typically defined by domain experts for a specific outcome of interest. Word or phrase counting methods (n-grams) convert free-text to grouped consecutive words (10). Controlled vocabulary methods convert free-text into a standardized language, using resources such as the Unified Medical Language System (UMLS)'s Metathesaurus, a large vocabulary database that contains information about biomedical and health related concepts, their various names, and the relationships among them (9,12,13). Document embedding methods use neural networks to represent the semantics of documents as vectors of continuous numerical values (11). Each of these methods produces different representations that can influence ML performance. Previous studies have investigated the classification performance of these types of NLP methods (14-16), however to the best of our knowledge only one study assessed generalizability as well (17). With this study, they investigated the performance of their embeddings on a single external dataset, however without extensive validation on multiple external datasets, there is a risk of overestimating both NLP strategies and ML models' performance (8,17,18).

We hypothesize that NLP methods will have more heterogeneous performance characteristics on external data compared with internal data. The LIRE data provide an opportunity to conduct a systematic evaluation of the utility of different representational methods for identification of image findings in radiology reports drawn from multiple healthcare systems. To assess the reproducibility and reliability of NLP methods, we need to test our methods on multiple external datasets. The purpose of our study is to assess the (1) performance within and (2) generalizability across the four LIRE healthcare systems of different NLP-based feature extraction methods: rules, n-grams, controlled vocabulary, and document embeddings, coupled with elastic-net logistic regression (i.e., the ML model) for classifying radiology reports for LBP-related findings.

# MATERIALS AND METHODS

### Annotated Dataset

This was a retrospective study that utilized the same annotated dataset from a previous study that showed that ML-based models were superior to rule-based classification (3). Our work is an extension of this as we expanded our NLP methods to include controlled vocabulary and document embeddings and explicitly assessed generalizability across healthcare systems. All participating IRBs agreed that the LIRE study was minimal risk and granted waivers of both consent and Health Insurance Portability and Accountability Act authorization (IRB approval number is 476829). We used limited dataset consisting of a subsample of the LIRE cohort which consisted of approximately 250,000 patients from four healthcare systems who received a thoracic or lumbar spine plain X-ray, magnetic resonance imaging (MRI), or computed tomography (CT) between October 1, 2013 and September 30, 2016 (7). The LIRE study was a multicenter intervention study that investigated whether inserting text about finding prevalence into lumbar spine imaging reports reduced subsequent spine-related treatments (7). Once in the study, patients were followed for two years and their data was regularly collected. We randomly sampled 871 index radiology reports, the first radiology report for each patient, and stratified by system and image modality (3). The sample size was determined based on prior NLP classification tasks (19). Each report was annotated for the presence of 26 LBP-related findings (Table 1) by a team of clinical experts composed of two neuroradiologists, a physiatrist, and a physical therapist. A single report can be annotated for multiple findings. See Supplement Appendix E3 for more details on this process. These findings were based on prior research consisting of a review (20), prospective cohort study (4), and randomized control trial (21) that characterized LBP based on its causes and treatments. Out of these 26, eight were considered to be potentially clinically important: any stenosis, central stenosis, lateral stenosis, foraminal stenosis, disc extrusion, nerve root displacement compression, endplate edema, listhesis grade 2 (4,5,7). Further details of this sampling and annotation process are presented in a previous study (3).

# **Classification Pipeline Overview**

Our classification pipeline analyzed the 871 LIRE radiology reports with the goal of learning patterns that are predictive of each of the 26 findings (Fig 1). The pipeline can be separated into three steps: preprocessing, featurization, and ML.

## **Preprocessing and Featurization**

For preprocessing, we developed regular expressions to help isolate the finding and impression sections of the 871 radiology reports. For featurization, rules, n-grams, controlled vocabulary mapping, and document embedding methods were used to extract features from the finding and impression sections. Rules require domain experts to identify terms that are related to an outcome of interest. This method is timeconsuming, but since the rules were developed by clinician experts, they can be considered a proxy for clinicians' judgement for annotations. In our implementation, we developed regular expressions based on the terms our team of clinical experts identified for each finding during the annotation process. For each report, we split the text into sentences. For each sentence, we identified the

presence of a finding using the regular expression and checked for negation (22). However, the presence of findings may be uncertain as radiology reports can have terms such as "suggesting" and "not definite." We minimized this uncertainty by coding these and other similar terms as indicating the presence of a finding. We used Java (v4.6.0), using Apache Lucene (v6.1.0), Porter Stemmer, and NegEx (23,24).

N-grams is a simple, but powerful method in NLP that converts free-text across the radiology reports to n-grams (phrases of different lengths) and indicates their presence and absence in each report (25). In our implementation, we used R (v3.6.1) package Quanteda (v2.0.1) to convert the text into un-, bi-, and trigrams.

Controlled vocabulary is a filtered version of the n-grams approach that leverages only clinically-related features from a standardized medical terminology mapped from the text (9). In our implementation, we first split the text into its constituent sentences using the maximum entropy model in the Apache OpenNLP toolkit to infer the end of a sentence (26). We then applied MetaMap Lite and an assertion classifier developed by Bejan et al., to each patient's radiology text report to obtain standard UMLS concepts and assertions of whether they were present, absent, conditional, possible or associated with someone else (27,28). We used MetaMap Lite because previous literature demonstrated MetaMap Lite's performance was comparable to or exceeded MetaMap and other similar methods (28). In addition, we also implemented a version of the controlled vocabulary method (*controlled vocabulary filter only*) that outputs recognized concepts as raw text, instead of the mapped UMLS terms to assess how a "many-to-one" mapping affects classification performance (29).

Document embeddings is a sophisticated approach that uses a neural network to convert the semantics of text into a continuous numerical vector (11). For the document embedding method, we used the Python (v3.7.3) package Gensim (v3.7.1) to implement the Distributed Bag of Words (DBOW) (11,30). We set the vector length to 600, number of epochs to 500, and allowed the model to initially learn word embeddings using the skip-gram architecture prior to learning document embeddings. We pretrained our DBOW architecture on the full text using two data sets: 522,283 radiology reports from the third version of Medical Information Mart for Intensive Care (MIMIC-III) (31), and the finding and impression sections from 255,094 unannotated reports from the LIRE study. We refer to these as *document MIMIC* and *document LIRE*, respectively. The former reflects a typical pretraining scenario and the latter assesses how pretraining on a corpus similar to our actual train/test corpus of 871 reports affects classification performance. We used these pretrained models to derive numerical vector representations of each of the 871 radiology reports. See Supplemental Appendix E1 for additional details. At the end of the featurization step, the textual data from radiology reports are represented as rules, n-grams, controlled vocabulary, controlled vocabulary filter only, document MIMIC, and document LIRE.

# **Rule- and Machine Learning-Based Model**

For the *rules*, we used a rule-based model to classify a report as "positive" for a finding if at least one mention in a report was non-negated and "negative" if there was no mention or all mentions of the finding were negated. We then used the trapezoidal rule approximation to calculate the area under the curve (AUC) (32). For each non-*rule* representation (i.e., feature

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set) and the finding labels from the annotation process, we developed an elastic-net logistic regression model to predict the presence of each finding (i.e., 26 binary or "one-vs.-rest" classification models) using the R (v3.5.1) package caret (v6.0–80). Within the training set, ten-fold cross validation was used to adjust the value of our regularization parameter (lambda) to perform feature selection on our predictors by shrinking our nonimportant predictors' coefficients towards 0. For each resulting finding-specific model, we identified the optimal threshold based on the training set's receiver operator characteristic (ROC) plot; the threshold is the point closest to the true positive rate of 1 and false positive rate of 0 (i.e., the point closest to the top left corner of the curve) using Euclidean distance (33).

# Performance and Generalizability Assessment

We used R (v3.5.1) to evaluate each representation. We used AUC of an ROC plot as the primary evaluation metric. This is because we envision our pipeline as an efficient "first-pass" screening tool intended to favor the identification of more true positives. For performance assessment, we randomly split our full dataset into 80% (697/871) for training and 20% (174/871) for evaluating our model for each finding. We assessed group-level performance by averaging the evaluation AUC across all finding-specific models and across all potentially clinically important finding-specific models, separately. We repeated this process 25 times with each independent repeat using a different random train/test split of the data, so that we could estimate 95% confidence intervals. For each finding/group, a t-test was used to assess significant performance comparing the 25 repeats of the best representation to the next best representation. We used Bonferroni correction to correct for multiple hypothesis testing; for the two groups, we considered p-value 0.025 (0.05/2 groups) to be significant, and for the 26 findings, we considered p-value 0.0019 (0.05/26 findings) to be significant. To assess generalizability across healthcare systems, we trained our model on reports from three systems and evaluated on the fourth, iteratively, for each finding. For each finding, we calculated the mean and standard deviation of the AUC across the four systems. We calculated group generalizability by averaging the AUC across all findings, and across all potentially clinically important findings for each system and then calculated the mean and standard deviation. We chose mean and standard deviation to quantify generalizability, because the former measures the quality, while the latter measures the consistency of performance across systems. For the generalizability assessment, we included all representations except for *rules* because they were developed using reports from all four systems, eliminating the possibility of evaluation using data unavailable at the point of algorithm development.

Additional details for these and other secondary analyses are provided in the Supplementary Materials.

# RESULTS

# **Data Summary**

In our dataset (n = 871), we sampled reports with similar proportions of image type (i.e., X-ray and MRI) and patients' age and gender across our healthcare systems (Table 2). For performance assessment, we've shown that our training set is representative of our test set

for 23/26 findings by using a t-test to assess the significant difference in the prevalence between sets across the 25 repeats for each finding (Fig 2). For generalizability assessment, we found that each healthcare system was comparable since the finding label prevalence across healthcare systems is overall similar with *any degeneration* having the highest label prevalence (0.896) and *listhesis grade 2* having the lowest label prevalence (0.028) (Fig 3).

# Comparing the Group and Finding Level Predictive Performance of Individual Representations

To assess the best performing representation, we trained and tested 26 finding-specific models for each representation and calculated finding-level and group-level AUC. On average across all findings, we found that the models generally performed well, with average AUC values above 0.87. *N-grams* and *controlled vocabulary* had the best (AUC = 0.960) and worst average (AUC = 0.879) performance, respectively (Table 3). At the finding level, *n-grams* had better performance than the corresponding second-best representation (which differed from finding to finding) for 22 findings, 11 of which were statistically significant. These results suggest that on average, the relatively simple methodology of *n-grams* is sufficient to classify our 26 findings.

In addition to assessing the performance of *n-grams*, we were also interested in characterizing the performance relative to *rules*, a representation requiring domain-expertise, and *document LIRE*, an advanced data-driven representation. This comparison is of interest as each of these three representations reflect different disciplines in featurizing textual data that range from domain-expertise to advanced domain-agnostic implementations. On average across all findings, *rules* (AUC = 0.897) performed worse than *n-grams* and *document LIRE*. At the finding level, *rules* was out performed by *n-grams* for eight out of twelve rare findings (prevalence < 20%). Additionally, while *document LIRE* had better overall performance than *rules*, it was not the best representation for any of the findings. This may be due to the fact that *document LIRE* may not have been the best representation but had stable performance across findings (min AUC = 0.799, max AUC = 0.979) compared to *rules* (min AUC = 0.649, max AUC = 0.999). These results also suggest when considering only rare findings, *n-grams* still perform better than other representations.

# Comparing the Group and Finding Level Generalizability Performance of Individual Representation Across Healthcare Systems

To assess the best generalizable representation, we trained 26 finding-specific models for each representation on data from three systems and tested on the fourth system, iteratively. For each finding/group, we calculated the mean and standard deviation of the test AUC across the four systems. At the group level, *n-grams* had the best average performance across all findings (mean AUC = 0.902) and at the finding level, it was the best performing representation for 10 findings (Table 4). The next best representation was *document LIRE* at the group level (mean AUC = 0.879) and it was the best method for 10 findings as well (Table 4). Interestingly, when considering standard deviation at the group level, we found that *document LIRE* and *n-grams* were the most (standard deviation = 0.010) and least consistent (standard deviation = 0.051) representations across all findings, respectively (Table 5). We found *n-grams* could not generalize well to system two, particularly resulting

in a lower sensitivity and higher specificity compared to other representations (Fig 4); we verified this result through complementary analyses (Supplemental Appendix E3). At the finding level, *document LIRE* was the most consistent representation for 11 findings. These results suggest that while *n-grams* had relatively the best performance, it had the worst consistency across systems. Instead, document embeddings pretrained on study-specific data (*document LIRE*) had relatively the most consistent classification performance on average across our systems.

# Assessing Performance and Generalizability of Potentially Clinically Important Findings

In our previous sections, we focused on all 26 findings, however we consider eight of these findings to be potentially clinically important. As a result, we believe it's important to present results for this important subset of findings. For performance assessment, *n-grams* had the best performance (AUC = 0.954), and it was significantly better than that of *document LIRE*, the second-best representation (AUC = 0.910) (Table 3). At the finding level, *n-grams* also had the best performance for all eight potentially clinically important findings, six of which were statistically significant. For generalizability assessment, *n-grams* had better performance (mean AUC = 0.898) than *document LIRE* (mean AUC = 0.890) (Table 4). At the finding level, *n-grams* and *document LIRE* had the best performance for seven of these findings. For consistency, *document LIRE* was the most consistent representation with standard deviation of 0.007 compared to *n-grams*' 0.076 (Table 5). At the finding level, *document LIRE* and *MIMIC* had the most consistent performance for six and two potentially clinically important findings, respectively, with one tie *endplate edema* (Table 5). These results indicate for this subset of findings, we still observe the same trend where *n-grams* has the best performance, but *document LIRE* has the best consistency.

# DISCUSSION

Manual extraction of information from radiology reports can be burdensome, making automated NLP methods attractive for such tasks. However, correctly estimating these methods' performance across multiple healthcare systems requires an understanding of their generalizability on external datasets. In this study, we compared and contrasted the performance of different NLP methods coupled with elastic-net logistic regression to classify 26 findings related to LBP through performance and generalizability assessment. Our study suggests that if classifier development and deployment occur at the same system, then *n-grams* may be preferable. However, for deployment at multiple systems outside of the system of development, one should consider n-grams with the caveat that it's consistency can vary across systems, while document embeddings pretrained on study-specific data (*document LIRE*) or a publicly available dataset (*document MIMIC*) had the most consistent performance.

Overall, based on performance assessment, n-grams, a relatively simple, data-driven, domain-agnostic method, is superior to more sophisticated methods (document embeddings and controlled vocabulary) in extracting known findings from text. These results are in line with prior studies (14,34). Additionally, for rare findings (prevalence < 20%), n-grams had the highest AUCs, which is consistent with a prior study evaluating n-grams coupled

with LASSO logistic regression to classify acute LBP (prevalence of 22%) (35). However, n-grams did not generalize well across the four healthcare systems when compared to document embeddings. This performance-generalizability duality can be explained as follows: the n-grams method is dependent on the precise phrasing in the training text. When we considered performance assessment, we split the full dataset into 80% (697/871 reports) for the train set and 20% (174/871 reports) for the test set; both sets contained the four healthcare systems, and their text were representative of each other. However, when considering each system independently for the generalizability assessment, n-grams were more susceptible to overfitting, i.e., they may have contained predictors more relevant to the training systems than the test system. When comparing summary statistics of the raw text among systems, we found that system two was indeed different from the other systems (Supplemental Appendix E3) and changing classification thresholds for the models tested on this system did not affect performance. In comparison, document embeddings better captured differently worded but synonymous concepts by transforming the raw text into abstract numerical representations that reflect semantics, leading to less deviation in performance across systems but slightly worse performance overall.

Document embeddings are of particular interest because they represent a sophisticated method of featurization that are pretrained on large-scale corpora to learn more generalizable representations of text. Here, document embeddings were pretrained on two different data sources, unannotated LIRE reports (smaller but more relevant to LBP) and MIMIC-III (larger but less specific). While *document LIRE* overall performed better than *document MIMIC*, the difference was modest, suggesting that a lack of task-specific corpus is not a barrier for using document embeddings in clinical tasks. This observation is consistent with other studies (16,36,37).

*Controlled vocabulary* and *controlled vocabulary filter only* are two representations that can be considered filtered versions of the n-grams approach that leverages only clinically related features. These representations differ from each other in that the former maps the clinically relevant raw terms to standardized terms to then use as features, while the latter does not map and instead uses the clinically relevant raw terms as features. As a result, *controlled vocabulary* performs a "many-to-one" mapping that can affect performance. When comparing these two representations, we found that *controlled vocabulary* marginally outperformed *controlled vocabulary filter only* in both performance and generalizability. Our study indicates that this "many-to-one" mapping is not detrimental to performance, but does not provide a substantial improvement relative to using only the clinically relevant raw terms as features.

Beyond performance and generalizability, scalability and interpretability are important factors to consider when choosing a NLP-based feature extraction method. Rules are the most interpretable method, because they solely rely on domain experts to provide the synonyms to search for in text. However, this method cannot scale well as expanding the synonyms for a more complete identification of findings and larger number of findings will require more time and domain experts. In contrast, n-grams, controlled vocabulary, and document embeddings are domain-agnostic computational methods, and as a result they can scale well to a large number of radiology reports and findings. These methods differ in their

interpretability. Controlled vocabulary and n-grams are the most interpretable methods as the former provides an ML model clinically relevant terms as features, while the latter provides the raw text as features. It is relatively easy for a researcher to examine a model's features and coefficients based on either of these two methods and understand what aspects of a radiology report are predictive of the outcome of interest. Document embeddings is the least interpretable method as it uses a neural network to represent a document's semantics as a vector of continuous values. These values are no longer interpretable as they are a result of the interactions of different word embeddings from the radiology reports in the neural network. When considering subsequent implementation, it is important to consider factors such as scalability and interpretability in addition to performance and generalizability.

There are several limitations to this study. First, our pipeline required binary annotations for findings, however the presence of findings may be uncertain as radiology reports can have terms such as "suggesting" and "not definite." We minimized this uncertainty by coding these and other similar terms as indicating the presence of a finding. Second, while our sample size was in line with recommendations for classification tasks, larger training and testing sets could have led to less variable performances across our different NLP methods (19). Third, we evaluated the algorithms but not the entire pipeline involving the querying and transfer of data; there may be discrepancies in our performance estimates when compared to those at actual deployment. Fourth, we could not assess our rules' generalizability, since the search terms were developed from reports from all four systems. Finally, in the case of document embeddings, because of our limited computational resources, we had to sequentially adjust hyperparameter values in the pretraining step, rather than conducting a grid search. With a more extensive hyperparameter search, we may have been able to improve performance.

Diagnostic imaging is often an early step for LBP patients that eventually leads to interventions, however the association between findings and LBP is uncertain (4). Jarvik et al. investigated this association and identified eight (potentially clinically important) findings that may be associated with a history of LBP and of these eight, *nerve root contact, disc extrusion*, and *central stenosis* may be associated with a new onset of pain (4,5,7). We've shown that our pipeline can automate classifying reports for these potentially clinically important findings using n-grams, and can generalize across healthcare systems using document embeddings. Our automated pipeline can assist similar studies by developing large cohorts quickly and inexpensively to investigate the association between findings and a clinical outcome within and across healthcare systems using text-based data.

# Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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# Abbreviations:

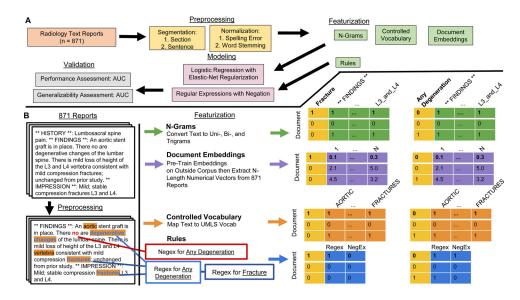
LBP	Lower Back Pain
NLP	Natural Language Processing
ML	Machine Learning
UMLS	Unified Medical Language System
LIRE	Lumbar Imaging with Reporting of Epidemiology
MRI	Magnetic Resonance Imaging
СТ	Computed Tomography
DBOW	Distributed Bag of Words
MIMIC-III	Medical Information Mart for Intensive Care
AUC	Area Under the Curve
ROC	Receiver Operator Characteristics

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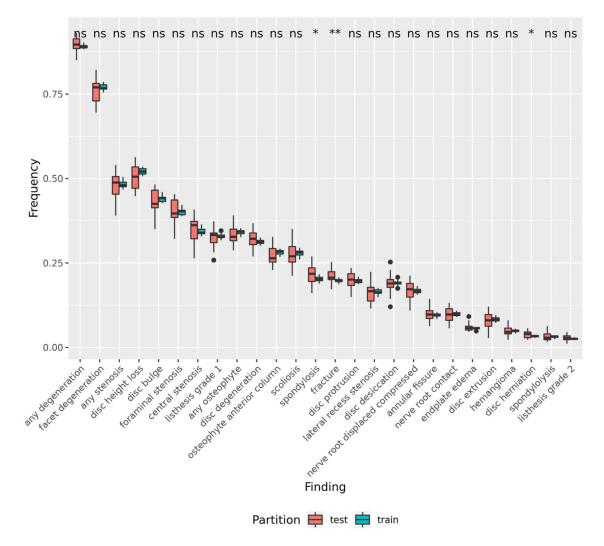
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### Figure 1.

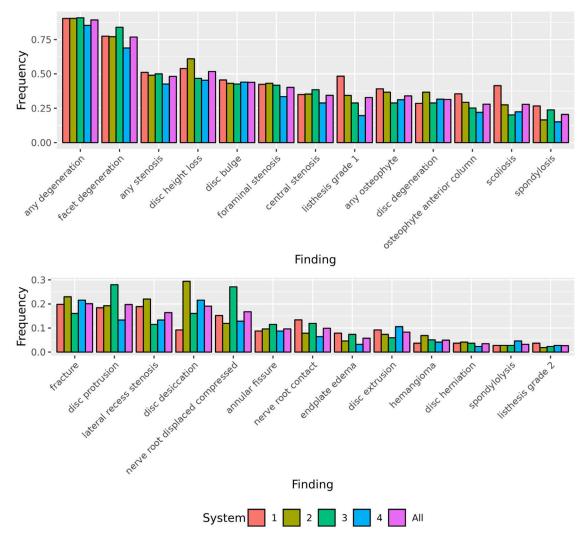
Overview of the Pipeline. (A) This visualization shows the different steps of our pipeline where we collect 871 radiology reports from our four systems, perform preprocessing to clean the text data, perform feature extraction using our four different methods. We load the *n-grams, controlled vocabulary*, and *document embeddings* feature matrices into a logistic regression model to predict the presence of these findings. For *rules*, we instead use a rule-based model that classifies a report as "positive" if at least one mention was non-negated and "negative" if there was no mention or all mentions of the finding were negated. We perform two types of assessments: generalizability and performance based on AUC. (B) A visual representation using the four different NLP methods to featurize the text for two example findings: *fracture and any degeneration*. The resulting finding-specific feature matrices are then used for the machine learning model, which uses the first column as the labels and remaining columns as features to predict the presence of these findings. AUC, Area Under the Curve; UMLS, Unified Medical Language System.



# Figure 2.

Comparison of the Finding Label Prevalence Between the Training and Test Set. We compared the finding label prevalence between the train and test sets across the 25 repeats. To assess a significant difference, we performed a t-test between the two sets for each finding. An asterisk indicates a significant difference, while "ns" indicates no significant difference.

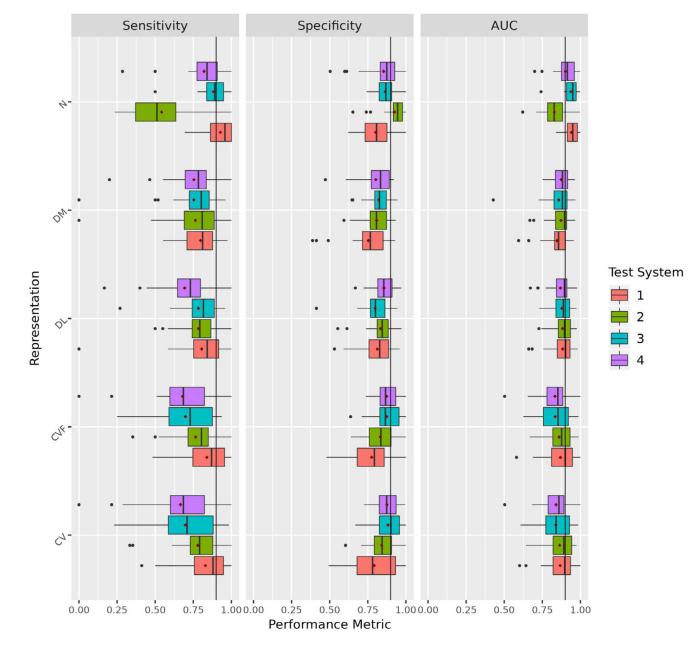
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# Figure 3.

Comparison of the Finding Label Prevalence Across the Healthcare Systems. We compared the finding label prevalence across the four healthcare systems. 1 = Kaiser Permanente of Washington, 2 = Kaiser Permanente of Northern California, 3 = Henry Ford Health System, 4 = Mayo Clinic Health System, All = all four systems.

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# Figure 4.

Assessing Generalizability of Individual Representations. We compared the generalizability of each of our representations and assessed performance using sensitivity, specificity, and AUC. For each representation, we plotted a boxplot to represent the distribution of the 26 findings for each test performance metric across healthcare systems. CV, Controlled Vocabulary; CVF, Controlled Vocabulary Filter Only; DM, Document MIMIC; DL, Document LIRE; N, N-grams. 1 = Kaiser Permanente of Washington, 2 = Kaiser Permanente of Northern California, 3 = Henry Ford Health System, and 4 = Mayo Clinic Health System.

# TABLE 1.

# The 26 imaging findings of our study

Type of Finding	Imaging Finding
Deformities	Listhesis-Grade 1
	Listhesis-Grade 2 or higher*
	Scoliosis
Fracture	Fracture
	Spondylosis
Anterior Column Degeneration	Annular Fissure
	Disc Bulge
	Disc Degeneration
	Disc Desiccation
	Disc Extrusion*
	Disc Height Loss
	Disc Herniation
	Disc Protrusion
	Endplate Edema or Type 1 Modic*
	Osteophyte-anterior column
Posterior Column Degeneration	Any Stenosis*
	Facet Degeneration
Associated with Leg Pain	Central Stenosis*
	Foraminal Stenosis*
	Nerve Root Contact
	Nerve Root Displaced/Compressed*
	Lateral Recess Stenosis*
Nonspecific Findings and Other	Any Degeneration
	Hemangioma
	Spondylolysis
	Any Osteophyte

Any stenosis refers to any of central, foraminal, lateral recess, or not otherwise specified. Any degeneration refers to any of disc degeneration, facet degeneration, or degeneration not otherwise specified. \* indicates the potentially clinically important findings.

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# TABLE 2.

Summary of the 871 Reports Based on Image Type, Average Text Length, and Patients' Age and Gender Across the Four Healthcare Systems

System	Image Type		IN III DARASEL AVELAGE TEAL LEIIGUL AVELAGE AGE	Avelage Age	remaie 70
Kaiser Permanente of Washington	X-Ray	102	$132 \pm 34$	$70.35\pm13.84$	0.60
	MR	115	$267 \pm 106$	$58.90 \pm 14.35$	0.49
	Total	217	$203 \pm 105$	$64.28 \pm 15.20$	0.54
Kaiser Permanente of Northern California	X-Ray	104	$143 \pm 38$	$67.51 \pm 16.82$	0.61
	MR	114	$270 \pm 95$	$57.1 \pm 14.96$	0.53
	Total	218	$210 \pm 97$	$62.06 \pm 16.68$	0.56
Henry Ford Health System	X-Ray	103	$121 \pm 57$	$67.15 \pm 16.04$	0.72
	MR	115	$268\pm152$	$58.95 \pm 15.77$	0.5
	Total	218	$199 \pm 137$	$62.96 \pm 16.44$	0.61
Mayo Clinic Health System	X-Ray	103	$141 \pm 39$	$69.35 \pm 16.15$	0.61
	MR	115	$222 \pm 104$	$55.13 \pm 15.44$	0.58
	Total	218	$184 \pm 90$	$61.85 \pm 17.28$	0.60
All	X-Ray	413	$134 \pm 44$	$68.58 \pm 15.76$	0.63
	MR	458	$257 \pm 118$	$57.52 \pm 15.17$	0.52
	Total	871	$199 \pm 109$	$62.79 \pm 16.42$	0.58

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report. For average text length, we calculated the average text length for both the finding and impression sections, since these sections were required for our pipeline. For both average text length and age, we included standard deviation.

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TABLE 3.

Best performing individual representations based on group and finding level AUC

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Finding	Prop.	P-Value	N-Grams	Document LIRE	Rules	Document MIMIC	Controlled Vocabulary	Controlled Vocabulary Filter Only
All Findings		1.06E-24 <sup>†</sup>	0.960 (0.949, 0.972)+	$0.910~(0.892, 0.929)^{\diamond}$	0.897 (0.882, 0.911)	0.894 (0.872, 0.916)	0.882 (0.862, 0.901)	0.879 (0.857, 0.902)
Potentially Clinically Important Findings		2.06E-13 <sup>†</sup>	0.954 (0.925, 0.983)+	$0.910(0.878,0.942)^{\diamond}$	0.821 (0.789, 0.852)	0.888 (0.856, 0.920)	0.857 (0.821, 0.894)	0.854 (0.813, 0.895)
any degeneration	0.896	0.03289	0.947 (0.906, 0.989)+	0.896 (0.820, 0.972)	0.850 (0.770, 0.931)	$0.874\ (0.789,\ 0.958)$	$0.936(0.911,0.961)^{\diamond}$	0.913 (0.882, 0.943)
facet degeneration	0.762	0.13484	0.970 (0.940, 0.999)+	$0.963(0.935,0.991)^{\diamond}$	0.873 (0.832, 0.914)	0.949 (0.919, 0.979)	0.923 (0.888, 0.959)	0.922 (0.883, 0.961)
disc height loss	0.507	5.46E-10 <sup>†</sup>	$0.931 (0.891, 0.970)^{+}$	0.833 (0.791, 0.875)	0.877 (0.829, 0.925)	0.830 (0.774, 0.886)	0.874 (0.812, 0.935)	$0.878~(0.826,0.930)^{\diamond}$
any stenosis*	0.480	0.09604	0.972 (0.950, 0.994)+	0.957 (0.930, 0.983)	0.893 (0.856, 0.930)	$0.967~(0.945,0.988)^{\diamond}$	0.955 (0.926, 0.984)	0.961 (0.935, 0.987)
disc bulge	0.435	0.00276	0.986 (0.972, 1.000)+	$0.978~(0.956,1.000)^{\diamond}$	0.976 (0.953, 1.000)	0.967 (0.939, 0.994)	0.955 (0.923, 0.987)	0.952 (0.920, 0.984)
foraminal stenosis*	0.400	$0.00153^{\dagger}$	0.950 (0.922, 0.978)+	0.935 (0.899, 0.971)	0.914 (0.876, 0.952)	$0.936(0.906,0.965)^{\diamond}$	0.932 (0.898, 0.966)	0.930 (0.896, 0.964)
central stenosis*	0.351	6.37E-10 <sup>†</sup>	0.950 (0.919, 0.981)+	0.903 (0.876, 0.930)	0.773 (0.731, 0.815)	$0.915~(0.884, 0.947)^{\diamond}$	0.907 (0.861, 0.953)	0.901 (0.852, 0.950)
any osteophyte	0.332	3.44E-07 <sup>†</sup>	0.955 (0.916, 0.994)+	0.888 (0.845, 0.932)	$0.925~(0.894, 0.955)^{\diamond}$	0.886 (0.835, 0.937)	0.875 (0.818, 0.933)	0.880 (0.819, 0.94)
listhesis grade 1	0.324	0.03888	0.967 (0.941, 0.994) <b>+</b>	0.927 (0.880, 0.974)	$0.958 (0.928, 0.989)^{\diamond}$	0.910 (0.858, 0.961)	0.945 (0.903, 0.987)	0.947 (0.904, 0.989)
disc degeneration	0.322	0.00203	0.935 (0.898, 0.973)+	0.830 (0.792, 0.869)	0.904 (0.865, 0.944)	0.784 (0.726, 0.842)	$0.909~(0.842, 0.976)^{\diamond}$	$0.909~(0.863, 0.954)^{\diamond}$
scoliosis	0.274	0.03726	0.969 (0.943, 0.994)+	0.924 (0.888, 0.961)	$0.959~(0.924, 0.994)^{\diamond}$	0.929 (0.887, 0.971)	0.900 (0.850, 0.949)	0.901 (0.854, 0.948)
osteophyte anterior column	0.271	5.13E-10 <sup>†</sup>	0.953 (0.930, 0.976) <sup>+</sup>	0.867 (0.820, 0.914)	$0.913 (0.872, 0.954)^{\diamond}$	0.846 (0.785, 0.907)	0.882 (0.831, 0.933)	0.874 (0.825, 0.923)
spondylosis	0.217	0.39974	0.992 (0.977, 1.010)+	0.936 (0.881, 0.99)	$0.990~(0.974, 1.010)^{\diamond}$	0.901 (0.846, 0.955)	0.900 (0.836, 0.964)	0.883 (0.817, 0.949)
fracture	0.212	0.62647	0.949 (0.912, 0.987)+	$0.947~(0.921, 0.973)^{\diamond}$	0.883 (0.816, 0.95)	0.896 (0.839, 0.953)	0.914 (0.868, 0.960)	0.925 (0.878, 0.972)
disc protrusion	0.197	0.69348	0.977 (0.953, 1.000)	0.940 (0.906, 0.973)	0.927 (0.879, 0.974)	$0.910\ (0.866,\ 0.954)$	0.982 (0.952, 1.010)+	$0.980~(0.948,1.010)^{\diamond}$

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	0.957 (0.923, 0.990) $0.913 (0.854, 0.972)^{\diamond}$ 0.941 (0.909, 0.973)	0.958 (0.921, 0.994)* 0.705 (0.000 0.073)	$0.923\ (0.884,\ 0.962)$	0.817 (0.745, 0.888)	0 802 (0 747 0 808)
	*	0.705 (0.500 0.873)			(0/0.0.0) 770.00
		0.782 (0.098, 0.872)	0.907 (0.848, 0.966)	0.870 (0.819, 0.921)	0.864 (0.817, 0.911)
		0.649 (0.567, 0.731)	$0.948(0.918,0.978)^{\diamond}$	0.843 (0.771, 0.914)	0.844 (0.773, 0.915)
	0.944 (0.886, 1.000)	0.957 (0.905, 1.010)+	0.922 (0.848, 0.996)	0.763 (0.667, 0.860)	0.755 (0.650, 0.860)
+(966.0	$0.921~(0.859,0.982)^{\diamond}$	0.910 (0.827, 0.993)	0.914 (0.856, 0.973)	0.797 (0.716, 0.877)	0.818 (0.741, 0.894)
0.994 (0.963, 0 1.020) <sup>+</sup>		0.886 (0.775, 0.997)	0.948~(0.901, 0.995)	0.969 (0.914, 1.020)	0.962 (0.897, 1.030)
0.916 (0.831, 0 1.000) <sup>+</sup>		0.854 (0.732, 0.976)	0.838 (0.713, 0.963)	0.789 (0.683, 0.896)	0.778 (0.626, 0.930)
$0.991 (0.950, 0.1.030)^{\diamond}$		$0.999 (0.995, 1.000)^{+}$	0.875 (0.705, 1.050)	0.963 (0.867, 1.060)	0.951 (0.845, 1.060)
$0.956 (0.878, 0.1.003)^{\diamond}$		0.975 (0.927, 1.020)+	0.912 (0.740, 1.080)	0.760 (0.526, 0.994)	0.786 (0.520, 1.050)
0.981 (0.940, 0 1.020) <sup>+</sup>		$0.936~(0.795, 1.080)^{\diamond}$	0.866 (0.747, 0.986)	0.754 (0.507, 1.000)	0.785 (0.561, 1.010)
0.905 (0.741, 0 1.070) <sup>+</sup>		0.767 (0.580, 0.955)	$0.683\ (0.486,\ 0.88)$	0.735 (0.559, 0.91)	0.706 (0.510, 0.902)
For each representation, we trained and tested 26 models (one for each findin, potentially clinically important findings for each representation. We repeated	g) on 80% and 20% of t this process 25 times wi	he dataset, respectively. ith different splits of the	For group level, we aver data to calculate 95% co	aged the AUC across all fin onfidence intervals. Table sh	dings and across all ows the best performing
	(0.963, 0 (0.831, 0 (0.950, 0 (0.950, 0 (0.940, 0	<ul> <li>(0.963, 0.979 (0.954, 1.000)*</li> <li>(0.831, 0.868 (0.765, 0.970)*</li> <li>(0.950, 0.899 (0.807, 0.991)</li> <li>(0.978, 0.899 (0.807, 0.991)</li> <li>(0.940, 0.890 (0.715, 1.060)</li> <li>(0.940, 0.832 (0.654, 1.010)</li> <li>(0.741, 0.799 (0.604, 0.994)*</li> </ul>	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	$\begin{array}{r llllllllllllllllllllllllllllllllllll$

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representation ordered left to right based on the All Findings row (1<sup>st</sup> row). The first column indicates the finding/group. For the findings, the second column indicates prevalence in the test set represented

the parentheses. Finally, for each finding and group, we performed a t-test comparing the best representation's distribution of AUC values for the 25 repeats to the second-best representation.  $\dot{\tau}$  indicates significant comparisons with Bonferroni correction (p-value significance for the groups: 0.025 = 0.05/2 groups, p-value significance for the findings: 0.0019 = 0.05/26 findings). Finally \* indicates the as a proportion. For each row, + indicates best performing individual representation based on the average AUC and \* indicates the second-best representation. We show the 95% confidence interval in

potentially clinically important findings. AUC = Area Under the Curve, Prop. = Proportion.

# TABLE 4.

Best performing individual representations based on group and finding level mean of AUC across systems

Finding	Proportion	N-Grams	Document LIRE	Document MIMIC	<b>Controlled Vocabulary</b>	Controlled Vocabulary Filter Only
All Findings		0.902+	0.879	0.868	0.857	0.853
Potentially Clinically Important Findings	ı	0.898	0.890	0.887	0.834	0.833
any degeneration	0.896	0.905	0.881	0.848	0.927+	0.874
facet degeneration	0.762	0.919	0.952+	0.927	0.914	0.898
disc height loss	0.507	0.845*	0.754	0.748	0.845+	0.837
any stenosis*	0.480	0.907	0.957+	0.955	0.946	0.943
disc bulge	0.435	0.954	0.963+	0.953	0.930	0.929
foraminal stenosis*	0.400	0.885	0.922+	0.913	0.904	0.888
central stenosis*	0.351	$0.916^{+}$	0.881	0.897	0.891	0.878
any osteophyte	0.332	0.874*	0.874+	0.860	0.831	0.833
listhesis grade 1	0.324	0.905	0.899	0.891	0.930	0.936+
disc degeneration	0.322	0.873	0.710	0.716	0.861	+806.0
scoliosis	0.274	0.911	0.892	0.891	0.865	0.870
osteophyte anterior column	0.271	0.832	0.825	0.818	0.845+	0.825
spondylosis	0.217	0.985*	0.801	0.764	0.823	0.781
fracture	0.212	0.910	0.926+	0.889	0.910	0.906
disc protrusion	0.197	0.948	0.935	0.904	0.977	0.978+
disc desiccation	0.189	$0.929^{+}$	0.909	0.850	0.796	0.797
nerve root displaced/compressed*	0.169	$0.918^{+}$	0.906	0.894	0.845	0.855
lateral recess stenosis*	0.163	0.915	0.932	0.934+	0.831	0.841
annular fissure	0.099	0.908	0.921+	0.886	0.766	0.763
nerve root contact	0.097	0.917	0.885	0.854	0.736	0.746
disc extrusion*	0.079	0.947	0.972+	0.925	0.968	0.959
endplate edema*	0.059	0.844	0.865+	0.841	0.725	0.762
hemangioma	0.049	0.917	0.869	0.908	0.963+	0.963+
disc herniation	0.038	0.820	0.826+	0.798	0.725	0.700

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spondylolysis	0.032	0.938* 0.814	0.814	0.849	0.775	0.804
listhesis grade 2*	0.028	0.838+	0.740	0.681	0.657	0.614
For each representation, we trained our model	on re	s from three sys	ports from three systems and evaluated on the fourt	ed on the fourth, iterati	ively, for each finding. Fo	r each finding, we calculated the mean of the AUC across

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the four systems. We calculated group-level performance by averaging the AUC across all findings, and across all potentially clinically important findings for each system and then calculated the mean across the systems. Table shows the best performing representation ordered left to right based on the All Findings row (1 st row). The first column indicates the finding/group. For the findings, the second column indicates prevalence in the test set represented as a proportion. + indicates the best performing representation for that finding and group. Finally \* indicates the findings that were potentially clinically important. AUC = Area Under the Curve.

# TABLE 5.

Most consistent individual representations based on group and finding level standard deviation of AUC across systems

Finding	Proportion	Document LIRE	<b>Controlled Vocabulary</b>	Document MIMIC	Controlled Vocabulary Filter Only	N-Grams
All Findings	ı	0.010+	0.012	0.013	0.014	0.051
Potentially Clinically Important Findings		0.007+	0.035	0.024	0.031	0.076
any degeneration	0.896	0.021	0.031	0.024	0.041	0.020+
facet degeneration	0.762	0.018+	0.039	0.022	0.038	0.064
disc height loss	0.507	0.033	0.050	0.010+	0.057	0.103
any stenosis*	0.480	0.023	0.032	0.013+	0.035	0.051
disc bulge	0.435	0.019	0.022	+600.0	0.015	0.040
foraminal stenosis*	0.400	0.016+	0.058	0.035	0.077	0.042
central stenosis*	0.351	0.030+	0.046	0.038	0.056	0.043
any osteophyte	0.332	0.034	0.099	0.031+	0.094	0.108
listhesis grade 1	0.324	0.032	0.014	+600.0	0.014	0.076
disc degeneration	0.322	0.033	0.052	0.092	0.049	0.026*
scoliosis	0.274	0.017+	0.051	0.025	0.043	0.068
osteophyte anterior column	0.271	0.025	0.084	0.020+	0.083	0.088
spondylosis	0.217	0.040	0.156	0.036	0.164	0.015+
fracture	0.212	0.019+	0.028	0.029	0.039	0.029
disc protrusion	0.197	0.021	0.011+	0.018	0.016	0.053
disc desiccation	0.189	0.024	0.052	0.020+	0.051	060.0
nerve root displaced/compressed*	0.169	0.020	0.018	0.015	0.011+	0.021
lateral recess stenosis*	0.163	0.017+	0.038	0.019	0.034	0.063
annular fissure	0.099	0.015+	0.064	0.057	0.065	0.063
nerve root contact	0.097	0.071	0.035	0.044	0.014+	060.0
disc extrusion*	0.079	0.007+	0.024	0.015	0.023	0.076
endplate edema*	0.059	0.059+	0.164	0.059+	0.091	0.126
hemangioma	0.049	0.073	0.048	0.007+	0.048	0.105
disc herniation	0.038	0.046*	0.079	0.092	0.065	0.094

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AUC across the four systems. We calculated group-level consistency by averaging the AUC across all findings, and across all potentially clinically important findings for each system as a test set and For each representation, we trained our model on reports from three systems and evaluated on the fourth, iteratively, for each finding. For each finding, we calculated the standard deviation of the

then calculated the standard deviation across the systems. Table shows the most consistent representation ordered left to right based on the All Findings row (1<sup>st</sup> row). The first column indicates the finding/group. For the findings, the second column indicates prevalence in the test set represented as a proportion. + indicates the most consistent representation for that finding and group. Finally \* indicates findings that were potentially clinically important. AUC = Area Under the Curve.