# **Assessing Fracture Healing** with Artificial Intelligence:

Using Transfer Learning to Predict the Radiographic Union Score for Tibial Fractures

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# **Declaration of Authorship**

I, Shen Zhou Hong, declare that this thesis titled "Assessing Fracture Healing with Artificial Intelligence: Using Transfer Learning to Predict the *Radiographic Union Score for Tibial Fractures*" and the work presented in it are my own. I confirm that:

- Where I have consulted the published work of other authors, this is always clearly attributed.
- Where I have quoted from the work of others, the source is always given. With the exception of such quotations, this thesis is entirely my own work.
- This thesis is entirely human work. No part of this thesis contains content generated by ChatGPT or other large language models.
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### **Abstract**

The use of AI models in the processing and interpretation of medical imagery is an emerging field, particularly in the domain of anomaly detection and localisation. However comparatively little work has been done with the characterisation of anomalies, such as the scoring of radiographs with standard clinical instruments. This study explores the use of transfer-learning as a method to develop an AI model that is able to score radiographs of long-bone fractures using Whelan et al's *Radiographic Union Score for Tibial Fractures*. By first training a baseline model using end-to-end training, the validity of transfer-learning as an approach is validated. Next, two different models with the same architecture, but different weights, were evaluated and compared. Finally, taking the best-performing ImageNet weights, a hyperparameter search was conducted, yielding optimal hyperparameters for a final model. This model achieved an AUROC of 0.890 on the hold-out evaluation set, demonstrating persuasive performance despite the limited data available.

Keywords: transfer-learning, medical imaging, radiography, union, machine learning, AI, computer vision.

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# Chapter 1

### Introduction

Long bone fractures are a frequent effect of high-energy trauma [1], among which tibial fractures of the lower extremities are the most common. These fractures require long-term follow-up, where after initial fixation the fracture site must be re-examined at regular intervals for callus formation<sup>1</sup>, bridging, and union [2]. Whelan et al's Radiographic Union Score for Tibial Fractures (RUST score) is a discrete 12-point scale that serves a common metric for the assessment of union from the lateral and anteroposterior<sup>2</sup> radiograph of a fracture [3]. This project proposes a means to automate the assessment of fracture healing, by using transfer-learning to develop a machine learning model to classify radiographs according to their RUST Scores.

#### 1.1 Aims and Motivation

Non-union and delayed union are significant complications in fracture healing, one which results in heightened morbidity, loss-of-function, and infection risk [4]. As a result, it is important for physicians to determine non-union events so that further treatment and corrective surgery may be taken. Although non-union may be determined through a variety of clinical assessments (e.g. palpation, weight-bearing), the RUST score is emerging as a quantitative radiography-based measure with high consistency [5], biomechanical correlativity [6], and good guidance for postoperative rehabilitation [7]. However, in order to assess a fracture using the RUST score, a orthopaedic must examine at least two radiographs (one lateral, one anteroposterior) for callus formation — a non-trivial process.

Recent advances in deep learning, coupled with the increasing availability of large radiographic datasets (e.g. CheXpert, LERA, MURA) [8, 9, 10] offer the

<sup>&</sup>lt;sup>1</sup>The development of cartilaginous material containing bone-forming cells.

<sup>&</sup>lt;sup>2</sup>i.e. front-to-back.

possibility of automating the process of fracture classification. Certain research models such as Rajpurkar et al's DNN ConvNet are able to meet, or exceed radiologist-level performance for abnormality classification in specific anatomical domains [10], and as of 2021 commercial developments are beginning to see regulatory clearance<sup>3</sup> [11].

However, much of the current available literature<sup>4</sup> is focused on the mere detection and classification of fractures (i.e. abnormality detection). Such models either perform binary classification (e.g. "Is this a *normal* radiograph?"), multi-class (e.g. "Is this a *leg*, *arm*, or *knee* fracture?"), or localisation (e.g. "Where *is* the fracture on this radiograph?"). Comparatively less work has been done on the *characterisation* of radiographs, where the properties of a fracture are described [12]. The existence of this 'research gap' can be in part attributed to the absence of large radiographic datasets with quality labelling.

This gap in the field offers opportunity for further investigation, especially as it is not the mere presence of a fracture which informs medical decision-making, but rather its severity and properties. This study aims to build a machine-learning model which infers the RUST score of a fracture from a pair of radiographs, specifically utilising the technique of *transfer-learning* in order to address the challenges of working with small datasets. By building a model that infers the RUST score of a fracture from it's associated radiograph, we hope to demonstrate the feasibility of such an approach, and open the door to further investigation,

### 1.2 Objectives and Evaluation

The objective of this project is to develop an AI model using *transfer-learning* that is able to predict the RUST score of a pair of anteroposterior and lateral radiographs. Because our primary constraint is the size of our dataset (around 3,000 radiographs, see Dataset), we will be utilising the technique of transfer-learning to demonstrate the feasibility of training AI models on datasets of this order of magnitude. We will be using the InceptionV3 model architecture [13] as the base model with a custom classifier for our transfer-learning approach.

First, we will build and evaluate model that is end-to-end trained without the use of transfer-learning (Protocol I). This will serve as a baseline benchmark for our subsequent attempts. Next, we will evaluate two different transfer learning approaches based on the InceptionV3 model architecture. The first model will use InceptionV3 trained with the general-purpose ImageNet dataset [14] (Protocol II). The second model will use the same InceptionV3 architecture, but trained with the domain-specific RadImageNet dataset [15] (Protocol III). The development and comparative evaluation

<sup>&</sup>lt;sup>3</sup>Authorisation for real-world clinical use by national health agencies like the Food and Drug Administration (FDA), Health Canada, *Conformité Européenne*, etc.

<sup>&</sup>lt;sup>4</sup>A selection of which are analysed with commentary in chapter 2.

of these two base models for transfer learning will allow us to compare and contrast the use of a model pre-trained on a general-purpose dataset (ImageNet) versus a model pre-trained on a slightly smaller, but domain-specific dataset (RadImageNet).

Afterwards, we will select the best-performing base model and find the best-performing hyperparameters. This will be done via a hyperparameter search in two steps, called Regime I and Regime II. Regime I will be a random-search on batch size and dropout, while Regime II will be a grid search on learning rate and epsilon. Finally, the model trained with the best-performing hyperparameters will be evaluated on the hold-out test set, and the overall performance will be examined.

## **Chapter 2**

# **Background Research**

The use of artificial intelligence in the analysis of radiography predates deep learning, with early approaches reliant on predefined engineered features and handcrafted algorithms (e.g. edge detection, wavelet transform) [16]. With the advent of more powerful computer hardware and the democratisation of machine learning through open source frameworks, we see deep learning techniques being applied to the field of medical imagery. These studies often had to solve unique, domain-specific challenges — such as small datasets, the need for *evidence-based* labeling 1, and difficult image-preprocessing requirements. This project will face similar challenges. Hence, by taking a survey of existing literature, we may be better informed in overcoming these challenges.

### 2.1 Early Period: Small Datasets, Feature-Engineering

The first period of AI-based based fracture detection was characterised by the limitations of small datasets <sup>2</sup>, and innovative approaches aimed to overcome said limits. One example of work in this period was Cao et al's use of feature fusion in randomforests, which allowed multiple categories of features to be considered by the model [17]. Likewise, Dimililer's *Intelligent Bone Fracture Detection system* used meticulous image pre-processing, with Haar wavelet transforms and Sub-Variant Feature Transform (SIFT) <sup>3</sup> in order to extract invariances from the radiography image [18]. Both authors relied on a combination of domain-specific image pre-processing and feature-engineering, in order to compensate for limited datasets that they had. The limitations of their data further manifested in difficulties with model evaluation. In

<sup>&</sup>lt;sup>1</sup>Labelling that is done by a clinical professional who is empowered to issue diagnoses.

<sup>&</sup>lt;sup>2</sup>Often working with an individual hospital or medical center, these studies these studies generally had hand-annotated datasets of up to a hundred images, and seldom more than two hundred.

<sup>&</sup>lt;sup>3</sup>Both functions are algorithms from the domain of signal processing, designed to compress spatiotemporal information in a manner that preserves invariances.

[18], only 100 labelled radiographs were available, making the use of a distinct hold-out validation set impossible. The lack of a separate validation set makes it difficult to judge whether or not the model performs without overfitting, weakening the study's conclusion. One possible mitigation that the author of [18] could have considered was k-fold cross-validation, which [17] does implement. In [17] Cao et al. implements 10-fold cross-validation over a data set of 145 radiographs, hereby yielding a more rigorous assessment of the model's performance. However, [17] did not use any data augmentation strategy, which may have been useful in light of the limited dataset available.

What lessons can we draw from these two studies? Radiographic imagery is highly hetereogenous, with individual x-rays mostly consisting of dark and light regions with sharp transitions in-between. Pre-processing steps such as scaling or down-sampling must capture these discontinuities faithfully, in order to prevent information-loss in the input. We can take particular inspiration from [18], who uses Haar wavelet transform in order to downsample inputs without loss of detail in the transition boundaries. Likewise, we must use k-fold cross validation, and consider a data augmentation strategy for our own model. Overall, these two examples are representative of early, exploratory work in AI-based fracture detection, and they are useful to illustrate both solutions to working with small datasets (pre-processing, feature-engineering), as well as challenges (difficulties with validation). The context of the above studies help us better understand later work, which began with the advent of large, publicly-available datasets: the most important of them being the Stanford Musculoskeletal Radiography Dataset (MURA).

<sup>&</sup>lt;sup>4</sup>According to Dimililer: "Unlike the discrete cosine transform, the wavelet transforms are not fourier based and therefore discontinuities in image data can be handled with better results using wavelets." [18]

### 2.2 MURA: Large Musculoskeletal Radiography Datasets

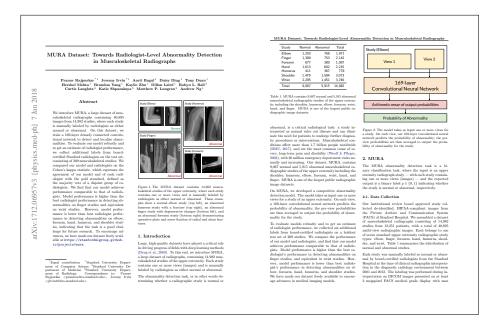


Figure 2.1: Thumbnail of article by Rajpurkar et al. [10]

MURA is one of the first large radiographic datasets focused exclusively on musculoskeletal imagery, as well as the name of a study conducted alongside said data by the Stanford Center for AI in Medical Imaging [10]. The MURA study marks a distinct landmark in the field of AI-assisted fracture classification, because it was the first to apply a deep-learning model on a large (40,561 radiographs), publicly available dataset. As a result, an examination of MURA allows us to contextualise our study in important ways: first, the success of MURA establishes the *possibility* of using deep learning to robustly analyse radiography. Rajpurkar et al. demonstrates near-radiologist levels of performance, validating the feasibility of our project in general. Secondly, the fact that the MURA model takes multi-view input imagery will help inform our own architectural design for processing both lateral and antero-posterior data. Finally, the limitations of MURA being a pure anomaly-detection system allows us to understand the need for a model which goes beyond binary classification, i.e. what our project is trying to accomplish.

To begin, it will be useful to look at MURA's architecture The MURA model is a 169-layer convolutional neural network which takes one or more views as input<sup>5</sup>, and delivers a *probability of anomaly*. The final probability is the arithmetic mean of output probabilities from every view [10]. Every type of radiograph may have

<sup>&</sup>lt;sup>5</sup>Different views are radiographs of the same subject taken at different standard perspectives.

multiple standard views, and the model architect has a choice of either training a model on a single view, or combining information from views in some ensemble stage. For anomaly detection, MURA chose a fairly conservative approach of assessing a separate probability of anomaly for each view, and then finding their average. This approach will be similar to the one that our project must take: which is to look at both the lateral and anteroposterior view of the tibia.

Another aspect of MURA that is worth examining, is their evaluation process. The model was accessed in two different ways: first, the model's precision versus recall plotted out in a Receiver Operating Characteristic (ROC) plot, and then the Area-Under-Curve of the ROC (AUROC) was quantified. The AUROC is a common metric to assess diagnostic ability since it serves to quantify the precision-recall curve of the model. The MURA model had an AUROC of 0.929. Second, a panel of three radiologists were assembled to evaluate a set of radiographs, and their performance was compared against the model. This kind of competitive evaluation allowed the study to compare the model performance against human clinicians, yielding an informative baseline for the AUROC. With this information, we can contextualise the earlier AUROC value of 0.929, and see that the model performs slightly worse than humans.

The use of AUROC as an evaluative metric, as well as competitive evaluation with human clinicians, present an advancement in model evaluation compared with the two earlier studies. However, in practice any AI model in radiography will not aim to replace human radiologists entirely, but serve as an additional tool which augments a human clinician's own diagnostic ability. Thus, the MURA study's evaluation is not representative of what real-world deployment would look like. This is why we must turn to Lindsey et al's *Deep neural network improves fracture detection by clinicians*, for a more holistic example of model evaluation.

### 2.3 Lindsey et al: Refinements to Model Evaluation



Figure 2.2: Thumbnail of article by Lindsey et al. [19]

In this study, a deep convolutional neural network is trained on a dataset of 31,490 wrist radiographs [19]. Like MURA, the study presents an AI-based fracture detection model, outputting both labels as well as a location heatmap. We include this study in the background research, for it's more holistic approach to evaluation, which simulates a real-world use-case. "Radiographic interpretation often takes place in environments without qualified colleagues available for second opinions" [19], the paper acknowledges, before proposing a model which aims to serve a possible 'second opinion.' After an initial evaluation which demonstrated an AUROC of 0.967 on certain datasets<sup>6</sup>, Lindsey et al. proceeds to conduct a second experiment, where a group of clinicians were shown radiographs from the same test set, and tasked to evaluate the radiograph both with and without the model's assistance:

<sup>&</sup>lt;sup>6</sup>For model testing, the study utilised two distinct categories of radiographs divided into 'Test Set 1' and 'Test Set 2' (a slight improvement over MURA), with the former containing a collection of singleton wrist radiographs, and the latter of wrist radiographs with anterio-posterior and lateral views.

For each radiograph shown, the clinicians were asked whether or not a fracture was present. After a clinician made a response, the model's semantic segmentation prediction [i.e. heatmap] was shown overlaid on the radiograph; the model's clinical determination was shown as text, and the clinician was asked the same question again. [19]

This experiment simulates the workflow of a clinician when using a fracture-detection model as a part of a CAD workflow. By doing so, we evaluate the effectiveness of the model as an useful tool within a broader clinical practice. The study found that the "... sensitivity and specificity of the emergency medicine MDs were significantly improved with the assistance of the deep learning model." [19]

The key advantage of Lindsey et al's paper lies in its evaluative design, which our own project must take inspiration from. It may be impractical to setup a similar trial involving radiologists or clinical professionals, but one feature that we do aim to replicate is the output of a heatmap which highlights the location of features that the model detects. For the Lindsey et al. and the MURA study, these heatmaps highlighted fracture-sights, whereas for our project they will highlight the sites of callus formation and bridging. By having this as a output, we will make the model's behaviour much more interpretable, and allow for integration into real clinical workflows.

<sup>&</sup>lt;sup>7</sup>Such an addition to this project is, however, within the realm of possibility, in collaboration with the medical faculty at METRC.

#### 2.4 Kim & MacKinnon: Cross-Domain Transfer Learning



Figure 2.3: Thumbnail of article by Kim and MacKinnon. [20]

Studies [10, 19] both rely on large, labelled radiographic datasets. While such datasets exist for fracture *detection* and fracture *classification*, the classification of fracture healing by their RUST scores is without precedence in literature, and to our knowledge there are no openly available datasets with the aforementioned labels. For our project, we will be using radiographic data internally collected by the Major Extremity Trauma Research Consortium (METRC), a research unit at the Johns Hopkins Bloomberg School of Public Health. Although this dataset has the RUST-scores we require, the number of radiographs we have access to is limited. Hence our approach is to utilise *transfer learning*, a process where the model is first trained on a larger, general-purpose dataset, before being fine-tuned on the study data. Thus, we turn to Kim and MacKinnon's *Artificial intelligence in fracture detection* for their innovative use of transfer learning. [20]

In Kim and MacKinnon's study, a dataset of 1,389 radiographs was available for a binary classification task (i.e. labels were "fracture" or "no fracture"). Instead of training a model *ab initio*, a CNN trained on a general purpose, non-radiographic dataset was re-applied on the radiography data. The authors used the Inception v3 network, an object-detection and image classification network [21] with the topmost layer fine-tuned on the radiographic data. [20]. By using a pre-existing model and

data augmentation<sup>8</sup>, Kim and MacKinnon were able to achieve an AUROC of 0.954, a value that is within the same standard deviation of [10, 19], studies which both had much larger datasets for their models.

Our inclusion of Kim and MacKinnon's study serves as an important validation for the aims of our project. The METRC dataset will consist of around two to three thousand radiographs sourced from a handful of METRC studies. If Kim and MacKinnon's model is already able to achieve a robust performance with only 1,389 samples, than it is quite possible for our own project to achieve it's aims. The theoretical reasoning behind why transfer learning can achieve high levels of model performance is because the first layers of a CNN are primarily responsible for high-level feature extraction, e.g. edge detection, pattern recognition, with only the latter layers responsible for task-specific classification [20]. This does, however open up a further avenue of inquiry. Would model performance improve further, if the base model that the transfer learning is conducted from, was itself originally trained on a domain-specific dataset, like the MURA data from [10]? Our project will explore this possibility, through it's own application of transfer learning to the prediction of radiographic union with RUST scores.

<sup>&</sup>lt;sup>8</sup>The authors of [20] ultimately generated 11,112 augmented samples from their initial set of 1,389 radiographs.

### **Chapter 3**

# Methodology

The primary objective of our study is to develop an AI model using the technique of transfer-learning, to automatically predict RUST scores from an input radiograph. In this chapter, we will first discuss our model design, contextualising the choices that we made in choosing our classifier, optimizer, and pre-training datasets. Next, we will discuss the primary datasets used in the study, before proceeding to the define the three protocols that we will conduct, and the hyperparameter tuning regime. Afterwards, the evaluation and endpoints will be presented, as well as a discussion on patient privacy and ethical considerations.

### 3.1 Model Design

Transfer learning is a technique which uses a model trained upon a larger dataset first, before being applied to a smaller, task-specific dataset. Assuming our task-specific dataset (in our case, the RUST data from METRC) is fixed, the performance of our transfer-learning model is determined by the following factors:

- 1. The architecture of the original model.
- 2. The initial, large dataset that the original model is trained on.

Hence, in the context of our study design, we must first choose (or create) a model architecture, and select an initial large dataset to train our model on. Only then are we able to apply the transfer learning procedure (e.g. freeze model weights, add new classifier, fine tuning, etc) upon our task-specific dataset.

<sup>&</sup>lt;sup>1</sup>i.e. the datasets that are used to train the base model, before transfer-learning.

#### 3.1.1 Model Architecture Choices

The first decision that we must make in our experiment design is the choice of model architecture. According to a literature survey by Litjens, Kooi, Bejnordi et al., convolutional neural networks (CNNs) are the most common deep learning architecture deployed for medical image analysis, vastly outnumbering alternative methods such as Stacked Autoencoders (SAE), or Recurrent Neural Networks (RNNs) [22, p. 77]. This is expected, as CNNs exhibit strong performance with image classification tasks, and as our project is an image classification task (albeit with radiographs), we will be using a CNN.

The choice now remains to either design our own CNN from scratch, or use an existing CNN architecture. Although designing a CNN *ab initio* allows the possibility of further experimentation and potentially finer-grained control, such a *de novo* model is difficult to assess holistically: there would be no prior work in literature to serve as a basis for comparison. In contrast, if we use a well-documented, existing CNN as our transfer-learning foundation, we may compare the performance of our implementation against other instances of the same model architecture used for transfer-learning in different domains.

Thus, we will be using the InceptionV3 model, a convolutional neural network developed by Szegedy, Vanhoucke, Ioffe, et al. from Google [13]. Our choice of InceptionV3 is based upon a 2022 literature review of transfer-learning models for the domain of medical image analysis. According to Kora, Ooi, Faust et al., CNNs with a broad (as opposed to deep) network topology perform well in transfer-learning tasks [23], with models like InceptionV3 outperforming more parameterized models like AlexNet [24]. Indeed out of the 54 studies included in the review, Inception-style models both the most common (14 out of 54) and among the highest-performing. [23]

#### 3.1.2 'Top-Classifier' Choices

Following the choice of our base model, we must then build a trainable classifier on top of the frozen base model layers. It will be the role of the classifier to transform the base model outputs into the 18-length one-hot encoded vector which will serve as our model's output. Top-classifiers are generally composed of a pooling layer, followed by layers of two or more densely interconnected neurons, with dropout. For our model, we have implemented the following classifier:

```
classifier: tf.keras.Sequential = tf.keras.Sequential([
    layers.GlobalMaxPooling2D(),
    layers.Dense(1024, activation='relu'),
    layers.Dropout(dropout_rate),
    layers.Dense( 512, activation='relu'),
    layers.Dropout(dropout_rate),
    layers.Dense( 256, activation='relu'),
    layers.Dropout(dropout_rate),
    layers.Dropout(dropout_rate),
    layers.Dense( 18, activation='sigmoid')
])
```

Listing 1: Classifier Architecture (Github)

Our use of three Dense layers is fairly typical of classifiers built for transfer-learning applications. What does warrant a brief explanation, however — is the choice of a GlobalMaxPooling2D() layer as a means to flatten the two-dimensional output of the base model. Why MaxPooling, instead of AveragePooling? We use MaxPooling because it 'selects' the brightest pixels per block-size. This performs well for images that are primarily composed of light pixels on dark backgrounds, such as our radiographs.

#### 3.1.3 Model Optimizer Choices

Finally, the last architectural decision we must make in our model design, is the choice of an optimizer. We have two possible approaches: we may either select an optimizer from *a priori* principles, or consider the selection of an optimizer to be a hyperparameter, and benchmark a variety of optimizers with our model on the data-set. Because we are already evaluating two variants of InceptionV3, the additional task of iterating through different optimizers will be infeasible given the time and compute constraints of the project.

Hence, our choice of an optimizer is determined by a review of available benchmarks and literature. The study and benchmarking of deep learning optimizers is a fairly recent field, initiated by the development of robust, reproducible benchmarks. Projects like Schneider, Balles, et Hennig's *DeepOBS: A Deep Learning Optimizer Benchmark Suite* allowed researchers to evaluate optimizers against an assay of realistic optimization problems, simulating common deep learning tasks and neural network architectures. [25] The availability of reproducible benchmarks allowed the first large-scale empirical experiments to be conducted on optimizer performance, culminating in Schmidt, Schneider, et Hennig's 2021 paper in optimizer benchmarking. [26] In an evaluation of 15 popular optimizers<sup>2</sup> across a total of more than 50,000 epochs of training, significant data on optimizer performance was gathered.

Of the eight optimization problem assays in the benchmark suite, our task of

<sup>&</sup>lt;sup>2</sup>AMSBound, AMSGrad, AdaBelief, AdaBound, AdaDelta, Adam, LookaheadMomentum, Lookahead-RAdam, Momentum, NAG, NAdam, RAdam, RMSProp, and SGD, respectively. The full list of results are available in their supplementary appendix.

radiographic image classification is bears closest resemblance to the CIFAR-10 benchmark: a CNN-based image classification task. According to the latest results available on the paper's website<sup>3</sup>, the ADAM optimizer has a slight accuracy improvement over Momentum and SDG. However, the differences are so small that the author mentions:

... a practitioner with a new deep learning task can expect to do about equally well by taking almost *any method* from our benchmark and tuning it, as they would by investing the same computational resources into running a set of optimizers with their default settings and picking the winner. [26, p. 2]

Therefore, we will select the ADAM optimizer, and spend time on fine-tuning the model and hyperparameter choices, instead of devoting further resources to selecting an optimizer.

#### 3.1.4 Pre-Training Dataset Choices

Now that we decided our model architecture, the next step is to select the initial, or pre-training dataset, that is used to train the base model.

#### **ImageNet Dataset**

Originally, InceptionV3 was trained on the ImageNet dataset: a general-purpose collection of more than 14 million everyday images. [14] The overwhelming size of the ImageNet dataset serves as a robust foundation for the InceptionV3 model, which exhibits strong performance in classification tasks upon it. However, the statistical characteristics of data in ImageNet is quite different from data in a typical radiography dataset. Hence, it remains a valid research question to ask whether a InceptionV3 model trained on a smaller, but more domain-specific dataset will exhibit better performance when applied to our transfer-learning task.

#### RadImageNet Dataset

Thus, we will also investigate RadImageNet, a collection of 5 million medical images composing of radiographic (CT), MRI, and ultrasound images. [15] As an open radiologic dataset developed specifically for transfer learning applications, a base model trained on RadImageNet may exhibit better performance on our radiography data, as the original network is trained upon images in a similar domain. However, the advantages of a similar domain is moderated by the corresponding smaller dataset size (5 million versus ImageNet's 14 million).

Therefore, in this project we will evaluate the use of a InceptionV3 model trained both on the ImageNet dataset, as well as the RadImageNet dataset as the base model for transfer learning. By comparing the performance of both models, we will be able to select the best-performing one for further development and refinement.

<sup>&</sup>lt;sup>3</sup>https://deepobs.github.io/leaderboardP4.html

#### 3.2 Dataset

Our transfer-learning models will be trained on a corpus of radiographic data sourced from a set of three METRC studies. [27, 28, 29] The data consists of pairs of radiographs (the anteroposterior view and the lateral view) along with a corresponding RUST score.

Dataset	Name	Samples
[27] RETRODEFECT	Retrospective Study of the Treatment of Long Bone Defects	1389
[28] PAIN	Pain Management & Long Term Outcomes Following High Energy Orthopedic Trauma	1233
[29] PACS	Predicting Acute Compartment Syndrome using Optimized Clinical Assessment	316
Sum		2938

Table 3.1: Sources of Radiographic Data with RUST labels in REDCap

The data is downloaded from the METRC Redcap database using the PyCap API library [30] and a set of Juypter notebooks (see Github repository link in Appendix), and then processed into a Tensorflow tf.data.Dataset object.

#### 3.2.1 One-Hot Encoding for Labels

RUST Scores will be one-hot encoded as 18-length label Tensors. Recall that our model must predict a RUST score from a pair of input radiographs. RUST scores are measures of orthopaedic union (i.e. healing), and ordinarily consist of two subscores for each view (the anterior-posterior and medial-lateral views), quantifying the development of bone calluses and bridging over the fracture line. The score components are as follows:

Radiographic Feature	Score
Fracture Line, No Callus	1
Fracture Line, Visible Callus	2
No Fracture Line, Bridging Callus	3
No Fracture Line, Remodeled.	4

Table 3.2: Radiographic Union Score for Tibial Fractures (RUST) Rubric

These score components  $^4$  are then used to assess the features of a fracture from the

<sup>&</sup>lt;sup>4</sup>Note that the original Whelan et al. paper [3] does not include a value for remodelled fractures, however as the METRC dataset includes this category, we will be using the modified RUST variant specific to Johns Hopkins for this study.

anterior, posterior, medial, and lateral cortices:

Anterior Cortex
Posterior Cortex
Medial Cortex
Lateral Cortex

Table 3.3: RUST Scoring Instrument

Finally the resulting subscore components are summed in order to yield a RUST score for the fracture as a whole. For this study, our model is designed to predict every component subscore. Hence, the label will consist of a 18-value tf.Tensor with the shape (18,), consisting of 16 one-hot encoded values for the RUST subscores (four for each cortice), as well as two additional one-hot values to represent the view (anterioposterior or medial-lateral).

We explicitly choose to one-hot encode every component subscore of the RUST instrument in order to avoid bifurcating our already small dataset. Recall that it takes a pair of radiographs (one anteroposterior, one lateral) in order to score a fracture. If we chose to label each radiograph simply according to their component cortex sum (e.g. Anterior cortex score + Posterior cortex score), then we would have to develop two separate models, one for the anterioposterior views, and another for the lateral views. This would cut the effective size of our dataset in half, as there are two radiographs for every fracture.

#### 3.2.2 Data Augmentation Strategy

Because we have a small dataset, a robust data augmentation strategy is needed to combat overfitting. In order to develop our data augmentation strategy, we followed suggestions from Bejani et Ghatee's *systematic review on overfitting control in shallow and deep neural networks* [31]. We may categorise data augmentation strategies to three broad families of methods: [32]

- 1. Geometric Methods: rotation, cropping, flipping.
- 2. **Photometric Methods:** noise, color-shifting, edge modification.
- 3. 'Complex' Methods: generating artificial data using GANs, style transfer.

Of the three, 'complex' data augmentation methods such as generating artificial data points is inappropriate for our use case, as it will compromise the evidence-based labelling of our dataset. Hence, we may recourse to either geometric methods, or photometric methods. Originally, our intuition was that photometric data augmentation methods would be of limited applicability to radiographs, which are entirely greyscale. Upon further investigation, it turns out that when evaluated individually, the most effective data augmentation strategy is cropping and rotation. [33] Our intuition was that cropping may not be appropriate, given that fracture-lines may lie on the peripheries of radiographs. Hence, we restrained ourselves to using only rotation, via random-flipping, as our method of data augmentation.

#### 3.3 Evaluation

We will evaluate the performance of our models using a combination of K-fold cross-validation (for the initial evaluation and hyperparameter search), and a hold-out test set (for the final evaluation of the completed model).

#### 3.3.1 Performance Metric: AUROC

The performance metric that we will be targetting is AUROC, the Area-Under-Curve of the Receiver Operating Characteristic (ROC) graph. This is the area underneath the precision-recall plot of the model, and is a common metric used to assess diagnostic accuracy. An AUROC of 0.50 indicates a diagnostic performance of no better than chance, while an AUROC of 1.00 indicates perfect performance.

#### 3.3.2 K-Fold Cross-Validation

We will be using K-Fold Cross validation as a technique to get a more accurate assessment of our model performance. K-fold cross validation will be used during the initial evaluation of the different models (see section 3.4), as well during hyperparameter search. We will be using a k value of k = 10 during the initial evaluation, and then a smaller k value of k = 6 for each sample of the hyperparameter space during the hyperparameter search (see section 3.5)

K-Fold	Percentage (of 3000)	Samples
6	14%	420
10	8.5%	255

Table 3.4: Size of validation set, per k-fold value.

Larger k values yield a more thorough measurement of a model's performance at the cost of additional computational costs, while lower k values risk lowering the training-validation split ratio until the training set is too small for adequate training.

#### 3.3.3 Hold-Out Test Set

A separate hold-out test set containing entirely unseen data will be reserved for a evaluation of the final model performance, after hyperparameter search. This hold-out

test set will be kept strictly separate for final evaluation only, and will be used only once at the very end. With a dataset of  $\approx 3,000$  samples, we will be setting aside 15% of the initial data ( $\approx 450$  samples) for the hold-out test dataset.

#### 3.4 Protocols

Now that we have finished discussing the model design, data, augmentation, and evaluation; it is time to define the study protocols. The experimental portion of this study will consist of three protocols:

- 1. Protocol I: Baseline: InceptionV3 end-to-end trained on RUST Dataset.
- 2. Protocol II: Transfer-learning w/ InceptionV3 trained on ImageNet.
- 3. Protocol III: Transfer-learning w/ InceptionV3 trained on RadImageNet.

All protocols will utilise the same data augmentation pipeline.

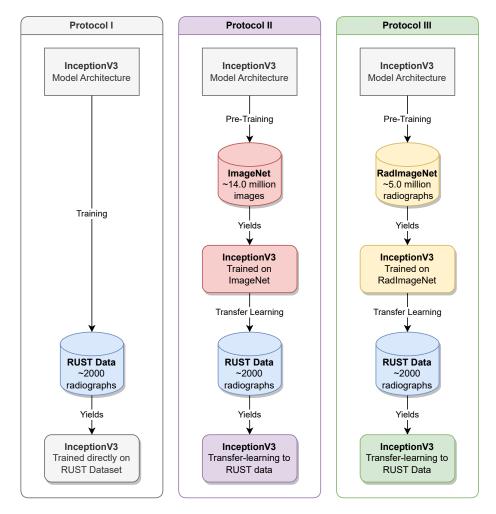


Figure 3.1: Overview illustrating the three model development protocols.

#### 3.4.1 Protocol I: InceptionV3 End-to-End Trained

Protocol I is designed to help us establish a baseline level of performance that we will use to evaluate the feasibility of using transfer learning. Hence we will be using the same model architecture (InceptionV3) and classifier (see subsection 3.1.2), but instead of using transfer-learning, we will train our model directly on the RUST dataset. We may expect a model that has only marginal performance, as the full InceptionV3 model has 23.9M trainable parameters [13], which will almost certainly not converge on a dataset of only 3000 samples. The performance of the Protocol I model will be used as a baseline for our subsequent investigations in transfer learning.

The model will be evaluated using k-Fold cross-validation, with k = 10.

#### 3.4.2 Protocol II: InceptionV3 with ImageNet

After establishing a baseline, we will move on to evaluating the use of ImageNet, or RadImageNet as our pre-training dataset for the transfer-learning model. Protocol II uses a base model (InceptionV3) with ImageNet weights. Here we use transfer-learning to train our classifier on the RUST dataset, and evaluate the AUROC of the resulting model using k-fold cross validation with k=10.

#### 3.4.3 Protocol III: InceptionV3 with RadImageNet

Finally, we instantiate a third model that uses the InceptionV3 architecture, except this time our pre-training dataset is RadImageNet. Here we use transfer-learning on the InceptionV3 model with RadImageNet weights, to train our classifier on the RUST dataset once more, and evaluate the AUROC of the resulting model using k-fold cross-validation with k=10.

Once this is complete, we will choose the better-performing model out of Protocols II and III (Protocol I's performance will almost certainly be marginal at best), and select it for further hyperparameter optimisation. This will take place according to two steps defined in the following section.

### 3.5 Hyperparameter Search

Using whichever is the better performing model, we will now attempt to optimise it's performance further by searching for better hyperparameters. We have the following hyperparameters that are available for consideration: *dropout rate*, *batch size*, *learning rate*, and *epsilon*.<sup>5</sup> What is the best method for us to systematically find the best hyperparameters? A brief survey of machine learning literature informs us that a variety of methods are possible, including sophisticated approaches such as Bayesian optimization, and genetic algorithms. [34] However, due to time and resource limitations we will be implementing only random search and grid search. We will perform a

<sup>&</sup>lt;sup>5</sup>A momentum factor similar to learning-rate decay specific to the Adam optimiser.

random search on the hyperparameter space defined by dropout and batch size, and then a grid search over a grid of different learning rates and epsilons.

This choice of performing hyperparameter search across two regimes is a practical compromise to the amount of compute resource that this project has. Although every hyperparameter influences every other hyperparameter, in practice it is rarely possible to conduct a hyperparameter search which varies over every hyperparameter under consideration at the same time. Instead, by constraining ourselves to two hyperparameters per search regime, we get to avoid the 'curse of dimensionality' Regime I is concerned with hyperparameters that are specific to the model, while Regime II is concerned with hyperparameters which are specific to the optimizer. Likewise, due to resource limitations, we will be performing k-fold cross validation for every search sample with a k = 6, instead of k = 10.

#### 3.5.1 Hyperparameter Search Regime I

Regime I uses a stochastic search process where the hyperparameter hypothesis space is randomly sampled t times. We use random search for Regime I because random search performs better than grid search over large hypothesis spaces, where the process of randomly sampling the hypothesis space will yield better hyperparameters as t increases, in a manner similar to Monte Carlo methods in statistics. [35] We begin by defining the range of batch size from 16 to 2048, and the range of possible dropout rates from 0.00 to 0.50.

Given this domain of batch size versus dropout, we will randomly sample the hypothesis space 100 times, where each sample contains a batch size and dropout rate pairing. Using this sampled pair, we will conduct a k-fold cross-validation experiment with k=6, where for every experiment the model will be trained for 20 epochs with said hyperparameters. The resulting average AUROC across the k-fold cross validation experiment will be used to assess the performance of said hyperparameters.

#### 3.5.2 Hyperparameter Search Regime II

After concluding Regime I, we will use the best-performing dropout and batch size, and proceed on to Regime II. Here, we aim to find the best learning rate and epsilon  $\epsilon$  (an Adam-specific momentum factor). Possible values for epsilon and learning rate span not merely a short interval (like those in Regime I), but across several orders of magnitude. Indeed, possible choices for learning rate can range from  $10^{-4}$  to  $10^{-1}$ , while possible choices for epsilon can range from  $10^{-7}$  to  $10^{-1}$ . Due to the sheer size of the search space, I judged random search to be infeasible. Instead, we will perform a grid search with a fixed set of learning rate and epsilon values, spanning each order of magnitude.

 $<sup>^6</sup>$ The concept in which the size of a hypothesis space grows combinatorially (n!) with the number of free parameters, i.e. degrees of freedom, which are added.

Specifically, we will investigate the following learning rates:  $10^{-4}$ ,  $10^{-3}$ ,  $10^{-2}$ ,  $10^{-1}$  against the following epsilon values:  $10^{-7}$ ,  $10^{-6}$ ,  $10^{-5}$ , ...,  $10^{-1}$ .

#### 3.6 Research Questions and Final Evaluation

Once we have completed the Hyperparameter Search Regime II, we will have obtained the optimal set of batch size, dropout, learning rate, and epsilon for our model. Now will be the occasion for us to evaluate this final model against our hold-out test set (see subsection 3.3.3). To do this, we will first train a new instance of our model with said optimal hyperparameters, observing its validation AUROC. Next, we will evaluate the model against the hold-out test set, and note the test AUROC.

Once we have done so, we will be able to answer the following research questions:

- 1. Does Transfer-Learning using the InceptionV3 model perform better than end-toend training with the InceptionV3 model?
- 2. Does the Transfer-Learning Model with the RadImageNet weights perform better than the Transfer-Learning Model with the ImageNet Weights?
- 3. Is Transfer-Learning able to achieve a persuasive level of performance, as measured by it's AUROC?

Protocol I and Protocol II will allow us to answer questions one and two. Once we have found the best-performing variant of InceptionV3, we will then begin the process of optimising model performance as measured through AUROC. The final model, as evaluated on the hold-out test set, will be used to answer question three. For the purpose of this study, we define a persuasive level of performance as a final AUROC that is greater than or equal to 0.80.

#### 3.7 Ethical Considerations

Any research project or study that involves human subjects will necessitate ethical consideration. This section is only a brief discussion of ethical risks and mitigations. The aim of this project is to validate transfer learning as a technique to infer RUST scores from a small dataset of radiographs. Although we aim to achieve robust performance in order to demonstrate the feasibility of this technique as an avenue for further research, at no time do we imply to develop a *diagnostic* tool for clinical use. The overarching spirit of the study is to explore interdisciplinary applications of AI in medical imaging, in hopes of reducing clinical caseload for medical practitioners, leading to better standards of care for all patients overall. This study has passed Goldsmiths IRB review.

#### 3.7.1 Human Subject Research, and HIPAA Compliance

This study works with radiographs collected from human subjects, that were a part of past or on-going METRC studies in high-energy trauma. This data is fully anonymised, and does not contain any personally identifiable information. As a part of Johns Hopkins Bloomberg School of Public Health's IRB requirements, all researchers working with human data, even anonymised data, must complete a Human Subject Research certification, and a Information Privacy Security certification. The author of this paper attests that they hold the aforementioned qualifications (see Appendix, Human Research Certification).

### **Chapter 4**

# Implementation and Results

We begin with the establishment of an initial baseline, by training an end-to-end (i.e. without transfer learning) variant of the InceptionV3 model on our radiography dataset. This will serve as a baseline that will allow us to validate the transfer-learning *technique* against regular end-to-end training.

Following the establishment of this baseline, we will proceed to begin an initial evaluation of two different transfer-learning base models. We will compare the performance of InceptionV3 trained with ImageNet weights [14], against InceptionV3 trained with RadImageNet [15] weights. This initial evaluation will help us explore whether a base model trained on the smaller, but domain-specific RadImageNet dataset will have any advantages over the larger, but general ImageNet dataset. We will select the better performing base model out of the two options, and proceed to optimize the model's hyperparameters.

Our model's hyperparameter search procedure consists of two steps, which we term hyperparameter search Regime I and hyperparameter search Regime II. As per our methodology, in Regime I we find the optimal batch size and dropout rate for our model. This is done using a stochastic search process where the hyperparameter space of the model is randomly sampled for t trials, where each trial consists of a k-fold cross-validation of the model with the selected hyperparameters. Once the optimal combination of batch size and dropout rate are found, we will set these hyperparameters as constant and proceed to the second hyperparameter search regime. In Regime II we find the optimal learning rate and epsilon value  $\epsilon$  for the Adam optimizer, by conducting a grid search over a selection of possible values.

#### 4.1 K-Fold Evaluation

Before we begin, we must first implement our k-fold cross-validation routine. Since model performance is sensitive to the network's random weight initialisation  $^1$  [36], our methodology requires k-fold cross-validation to be conducted on every experiment (i.e. model run). My implementation of the k-fold cross-validation process consists of two parts: a function which will divide the dataset into k folds, as well as a function that runs the k-fold cross-validation on the given model. The the k\_fold\_dataset() function is given as follows:

```
def k_fold_dataset(ds: tf.data.Dataset, k: int = 10) -> list[tuple[tf.data.Dataset,

    tf.data.Dataset]]:
    # First shard the given dataset into k individual folds.
    list_of_folds: list[tf.data.Dataset] = []
    for i in range(k):
        fold: tf.data.Dataset = ds.shard(num_shards=k, index=i)
        list_of_folds.append(fold)
    # Next, generate a list of train and validation dataset tuples
    list_of_ds_pairs: list[tuple[tf.data.Dataset, tf.data.Dataset]] = []
    for i, holdout_fold in enumerate(list_of_folds):
        ds_valid: tf.data.Dataset = holdout_fold
        # Select every fold except holdout_fold as the training folds
        training_folds: list[tf.data.Dataset] = list_of_folds[:i] +

    list_of_folds[i+1:]

        # ds_train size is \frac{k-1}{k} of the original dataset
        ds_train: tf.data.Dataset = training_folds[0]
        for fold in training_folds[1:]:
            ds_train = ds_train.concatenate(fold)
        ds_pair: tuple[tf.data.Dataset, tf.data.Dataset] = (ds_train, ds_valid)
        list_of_ds_pairs.append(ds_pair)
    return list_of_ds_pairs
```

Listing 2: Sharding dataset for K-Fold Cross Validation (Github)

One thing of note, is that our k\_fold\_dataset() function conducts all dataset-related operations using the Tensorflow's high-performance tf.data.Dataset API. This allows support for pre-fetch, caching, and other low-level optimisations. This function serves as a dependency which is called by cross\_validate(), which runs the actual K-fold cross validation experiments on the given model:

<sup>&</sup>lt;sup>1</sup>This is particularly true on small datasets with unbalanced classes like ours.

<sup>&</sup>lt;sup>2</sup>The code listings provided in this document *are for illustration only*. The actual implementation is generally longer, and contains docstrings, debugging instrumentation, file I/O logic, as well as additional function arguments. Every listing will have a link to it's corresponding implementation in the git repository.

```
def cross_validate(ModelClass: tf.keras.Model, ds: tf.data.Dataset, epochs: int = 50,
   batch_size: int = 128, k: int = 10) -> list[tf.keras.callbacks.History]:
   history_list: list[tf.keras.callbacks.History] = []
    train_valid_pairs: list[tf.data.Dataset] = k_fold_dataset(ds, k)
    for i, (ds_train, ds_valid) in enumerate(train_valid_pairs):
        # Reset tensorflow gradient tape
        tf.keras.backend.clear_session()
       model = ModelClass()
        model.compile(
            optimizer=tf.keras.optimizers.Adam(),
            loss=tf.keras.losses.BinaryCrossentropy(),
            metrics=metrics
        history = model.fit(
            ds_train,
            {\tt validation\_data=ds\_valid,}
            epochs=epochs,
            batch_size=batch_size,
       history_list.append(history.history)
   return history_list
```

Listing 3: K-Fold Cross Validation (Github)

The output of every k-fold cross-validation experiment will be a 'history list' containing k tf.keras.callbacks.History objects. This History object will contain training and validation metrics which will be used to calculate the average metric over k folds:

```
def calculate_mean_metrics(kfold_metrics: list[dict[str, float]]) -> dict[str,

    listΓfloat]:

    # Initialise aggregate metrics with appropriate keys
    aggregate_metrics: dict[str, list[float]] = {}
    for fold in kfold_metrics:
        for metric in fold.keys():
            if metric not in aggregate_metrics:
                aggregate_metrics[metric] = []
    # Calculate the average metric per epoch for every fold
    number_of_folds: int = len(kfold_metrics)
    for metric in aggregate_metrics.keys():
        number_of_epochs: int = len(kfold_metrics[0][metric])
        for epoch in range(number_of_epochs):
            # A list of every value for that given metric in this epoch across folds
            values_per_epoch: list[float] = [x[metric][epoch] for x in kfold_metrics]
            mean_per_epoch : float = sum(values_per_epoch) / number_of_folds
            aggregate_metrics[metric].append(mean_per_epoch)
    return aggregate_metrics
```

Listing 4: Calculating Mean Metrics from K-Fold Data (Github)

The above code now completes the prerequisites necessary for data gathering.

#### 4.2 Establishing Baseline Performance Target

In this section, we will establish the baseline performance targets for our transfer-learning model by training and developing an InceptionV3 model that is directly end-to-end trained on our radiography dataset. This evaluates whether the *technique* of transfer learning is appropriate for our dataset. Let us start by defining our base model:

```
class TransferLearningModel(tf.keras.Model):
   def __init__(self, dropout_rate: float, **kwargs):
        super().__init__(**kwargs)
        self.input_layer: tf.Tensor = layers.InputLayer(input_shape=(299, 299, 3))
        self.data_augmentation: tf.keras.Sequential = tf.keras.Sequential([
            layers.RandomFlip(seed=RNG_SEED),
        ٦)
        self.inceptionv3: tf.keras.Model = tf.keras.applications.InceptionV3(
            include_top=False,
            weights='imagenet'
        self.inceptionv3.trainable = False
        self.classifier: tf.keras.Sequential = tf.keras.Sequential([
            layers.GlobalMaxPooling2D(),
            layers.Dense(1024, activation='relu'),
            layers.Dropout(dropout_rate),
            layers.Dense( 512, activation='relu'),
            layers. Dropout(dropout\_rate),\\
            layers.Dense( 256, activation='relu'),
            layers.Dropout(dropout_rate),
            layers.Dense( 18, activation='sigmoid')
        self.model: tf.keras.Sequential = tf.keras.Sequential([
            self.input_layer,
            self.data_augmentation,
            self.inceptionv3,
            self.classifier
        ])
    def call(self, inputs):
        return self.model(inputs)
```

Listing 5: Model Class for InceptionV3 (Github)

We define a **class TransferLearningModel** which will be instantiated by every k-fold validation trial. Note that for this particular experiment, as we are establishing an end-to-end trained baseline, we will be setting self.inceptionv3(weights=None) and the attribute self.inceptionv3.trainable = True. Naturally in the actual implementation (Github) this is done through an argument in the class constructor, however the listing is simplified for the purpose of size and readability. So what happens now when we run the kfold experiment (Github)?

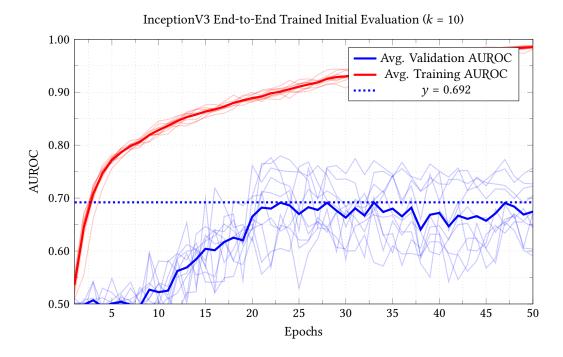


Figure 4.1: InceptionV3 Model Trained on Study Data.

The bold lines in the chart represent the *average* training (red) and validation (blue) AUROC, as measured after performing k-fold cross-validation on 10 folds (k = 10). The transparent lines indicate the observed training and validation AUROC per each individual fold: this per-fold performance has been charted in order for us to better observe the consistency of model performance per epoch. Variations in performance per fold is due to a combination of different random starting conditions (due to random weight initialisation at the start of a model's training), as well as variances in floating-point calculations.

We can observe that the end-to-end variant performs marginally: achieving an average AUROC of 0.692. However, upon a closer examination it is clear that the validation AUROC of each individual k-fold trial is highly erratic. The large spikes in validation AUROC indicates a failure to converge, as the dataset is too small for the number of tunable weights in the model. The InceptionV3 model has 189 layers, with a combined total of 23.9 million trainable weights: representing a parameter space several orders of magnitude larger than our dataset. The highly erratic validation AUROC is only a symptom of the model's inability to converge, and demonstrates clearly that regular end-to-end training is insufficient and inappropriate for our dataset.

### 4.2.1 Baseline Metrics

Having completed assessing our two baseline models, we are left with the following metrics that will help us in our own evaluation:

Baseline	Validation AUROC
Random (no better than chance)	0.50
End-to-End Model (InceptionV3)	0.69
Performance Target	0.80

Table 4.1: Baseline Benchmarks

The first level of performance that our subsequent models are expected to achieve is a validation AUROC > 0.50. As a measurement of classifier performance, an AUROC of 0.50 indicates performance no better than chance (i.e. the same as choosing by random). If we are unable to meet the minimum baseline of > 0.50, then our entire approach may be unrealistic and infeasible. Next, the baseline that we established allows us to validate the suitability of the transfer-learning technique. If our model is unable to meet an AUROC of > 0.69, then we will be better served to train our model architecture directly on our dataset.

With this information in hand, it is time for us to embark on the second part of our study: developing a transfer-learning model to infer the RUST score of radiographs of long-bone fractures. In the transfer learning technique, a base model is trained on a larger dataset, before having it's weights frozen, and then used as a component of a classifier trained on the task-specific dataset. A key decision in this process is the choice of the larger dataset that the base model will be trained on. Thus, we are lead to the second part of our study: evaluating the performance of InceptionV3 trained on ImageNet, and RadImageNet.

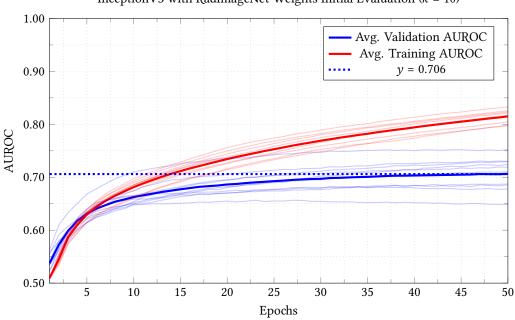
### 4.3 InceptionV3 with Transfer Learning

Let us begin by evaluating the performance of the RadImageNet weights (see subsection 3.4.3). Following the same procedure as we did earlier, we instantiate a InceptionV3 model with our classifier, & set self.inceptionv3 (weights='radimagenet.h5') and self.inceptionv3.trainable = False. This class instance now has InceptionV3 with pre-trained ImageNet weights, which we have frozen. Now when we run the training process, only our classifier will be trained. We conduct the experiment in the following Jupyter notebook (Github).

<sup>&</sup>lt;sup>3</sup>I choose to conduct Protocol III first, due to an idiosyncrasy in the process of loading Tensorflow weights. However despite this switch-up in ordering, the overall method remains the same.

### 4.3.1 Base Model Trained on RadImageNet Dataset

The model trained on RadImageNet weights achieves an average validation AUROC of 0.706. Note how unlike the version of InceptionV3 that was end-to-end trained, the per-fold validation AUROC is fairly consistent: we do not see any large spikes in validation performance. Likewise, although the model begins to exhibit overfitting after epoch 15, the degree of overfitting is relatively well controlled.



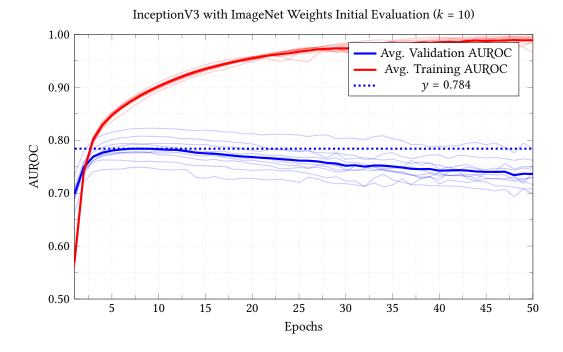
InceptionV3 with RadImageNet Weights Initial Evaluation (k = 10)

Figure 4.2: InceptionV3 with RadImageNet Weights

This preliminary information helps inform us that the technique of transfer learning is appropriate for our use case and dataset. All that follows now is for us to evaluate the ImageNet weights, and compare their performances together.

### 4.3.2 Base Model Trained on InceptionV3 Dataset

We now conduct the same experiment in a separate Jupyter notebook with ImageNet weights (Github). Recall that the difference between RadImagenet and ImageNet is that former contains approximately 4.1 million images [15], while the latter contains around 15.0 million [14]. While the RadImageNet dataset is exclusively sourced from medical imagery, including radiographs — the vastly larger ImageNet dataset has the potential to perform better, simply because the model was trained on a larger dataset. Does this assumption hold true?



### Figure 4.3: InceptionV3 with ImageNet Weights

Our data indicates that the InceptionV3 weights yield a higher validation AUROC of 0.784, in comparison to the RadImageNet model with an AUROC of 0.706. This means that although the RadImageNet dataset was more domain-specific to our needs (i.e. medical radiography classification), it appears the sheer size of the ImageNet dataset yielded weights which performed better.

### 4.3.3 Comparison between RadImageNet and ImageNet

However, despite this difference in 'naive' (i.e. untuned, without hyperparameter optimisation), the performance characteristics of both models are quite different. Observe how the overfitting profile of the RadImageNet model is less severe than that of ImageNet, despite achieving a lower validation AUROC overall. Likewise, while the validation AUROC of the ImageNet model begins to decrease past epoch 15 due to overfitting, the validation AUROC of RadImageNet is still growing by epoch 50. Although our methodology calls for us to select the better performing model out of both of them, the information shown here offers room for further investigation — which can be the subject of a future study.

At this point, having assessed the 'naive' performance of both the RadImageNet weights and the regular ImageNet weights, we will select the better-performing

ImageNet weights as the basis for our transfer learning model. Going forward, we are now ready to tackle our problem directly: and begin the process of hyperparameter search.

### 4.4 Hyperparameter Search

Recall that our methodology calls for a two-part hyperparameter search (see section 3.5), consisting of Hyperparameter Search Regime I, and Hyperparameter Search Regime II. Regime I is concerned with finding the best combination of batch size and dropout, while Regime II is for the best learning rate and epsilon  $\epsilon$  for the Adam optimizer.

### 4.4.1 Hyperparameter Search Regime I

To begin, let us construct a search function which conducts t trials, where during each trial a random portion of the hypothesis space is sampled, and then evaluated using k-fold validation with k=6. Recall that our methodology specifies the use of a slightly lower k value as a concession to the amount of compute resources available. Given 20 epochs of training per k-fold, and 6 folds per trial, this hyperparameter search regime conducts a total of 12,000 epochs of training over the course of 2 days (Github).

```
def hyperparameter_search(trials: int, kfolds: int = 6, epochs: int = 20) ->
→ list[dict[str, Union[int, float, list[tf.keras.callbacks.History]]]]:
   search_results: list[dict[str, any]] = []
   for trial in range(trials):
        # Randomly pick hyperparameter options
       rng = np.random.default_rng()
       batch_size : int = rng.integers(16, 2048, endpoint=True)
       dropout_rate: float = rng.uniform(0.0, 0.5)
       # Conduct K-Fold cross-validation with given hyperparameters
       results: list[tf.keras.callbacks.History] = cross_validate(
           TransferLearningModel.
           ds_train_and_valid,
           k=kfolds
           epochs=epochs.
           batch_size=batch_size,
           model_kwargs={"dropout_rate": dropout_rate},
       )
       search results.append({
            "batch_size" : batch_size,
            "dropout_rate": dropout_rate,
            "history_list": k_fold_results
       })
   return search_results
```

Listing 6: Hyperparameter Search Regime I (Github)

The results of this hyperparameter search are then plotted, with the maximum observed validation AUROC represented as the colour of the data point in the following scatter

plot. Note that the color-map is scaled according to the highest and lowest observed average validation AUROC.

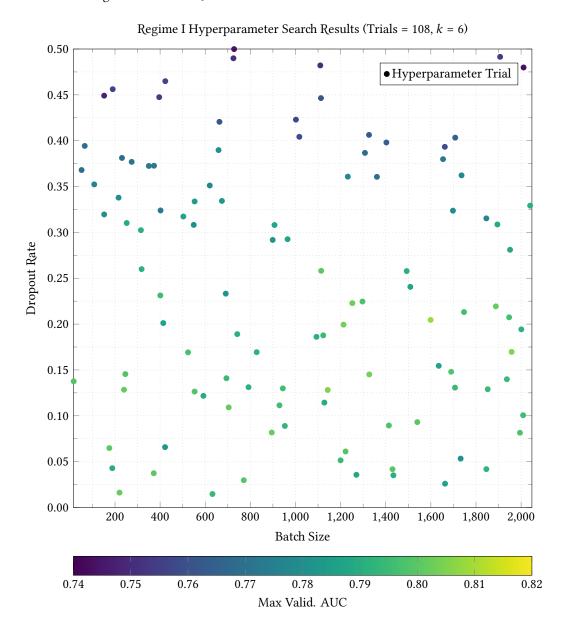


Figure 4.4: Results for the Hyperparameter Search Regime I

What were some of the learning rate and batch size combinations that we found? The following table lists out the top ten hyperparameter options.

Batch Size	Dropout Rate	Max Validation AUROC
1599	0.20450534699237183	0.8082009057203928
1958	0.16968093634911133	0.8052726884682974
1143	0.12816312484235065	0.8047231038411459
1327	0.14511673948299006	0.8034322460492452
1213	0.19936088970848825	0.8029904663562775
1252	0.22284567287235169	0.8021498719851176
1888	0.21933964138972928	0.8016549249490103
895	0.08177228898982053	0.8012685179710388
2009	0.10061578442603508	0.7949380973974863
401	0.23126040688192015	0.794610470533371

Table 4.2: Top Ten Hyperparameter Options for Regime I

As we can from a glance, a reasonable batch size seems to hover between 1500 to 2000. Likewise, a good dropout value seems to be around 0.20. This corresponds the colour gradient that we can see in the scatter plot. As an additional visualisation, we will also graph a random assortment of nine hyperparameter choices from our hypothesis space sample:

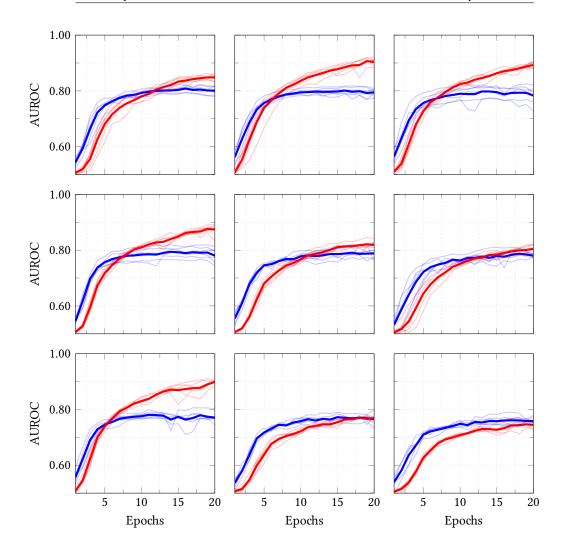


Figure 4.5: Random examples of models from hyperparameter search regime I.

The graphs above represent a selection of nine batch size and dropout combinations, sorted by best-performing to worse-performing. The first graph represents the top-performing hyperparameter combination, which we will graphs in greater detail below:

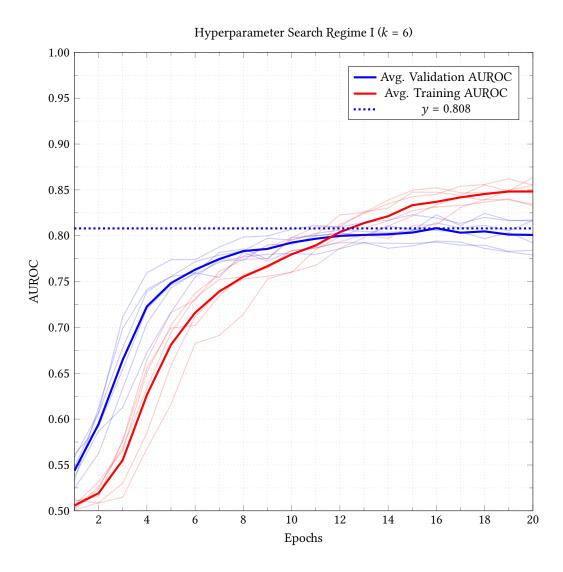


Figure 4.6: Best performing model in Regime I

Thus, we complete the first hyperparameter search regime, obtaining the hyperparameter options of 1599 for batch size and 0.205 for dropout. Going forward in all subsequent models, we will use those values, after rounding them up slightly to the nearest whole figure: batch\_size: int = 1600, dropout\_rate: float = 0.20. With this information, we may now move on to the second hyperparameter search regime, where we find the best hyperparameters for the Adam optimizer.

### 4.4.2 Hyperparameter Search Regime II

Recall that for Regime II, we must find the best learning rate and epsilon factor for the Adam optimizer. We begin by implementing a grid search function, as seen below. We sample the grid at discrete intervals for both learning rate and epsilon:

```
def learning_rate_gridsearch(kfolds: int = 6) -> list[dict[str, Union[int, float,
→ list[tf.keras.callbacks.History]]]]:
    # Grid i: 1.0 \times 10^{-1} \le learning\_rate \le 1.0 \times 10^{-4}
    learning_rates: list = [1 * np.float_power(10, -exp) for exp in range(1, 5)]
    # Grid j: 1.0 \times 10^{-1} \le epsilon_rate \le 1.0 \times 10^{-8}
    epsilon_rates : list = [1 * np.float_power(10, -exp) for exp in range(1, 9)]
    search_results: list[dict[str, Union[int, float,
      list[tf.keras.callbacks.History]]] = []
    for i, learning_rate in enumerate(learning_rates):
        for j, epsilon_rate in enumerate(epsilon_rates):
             # Conduct K-Fold Experiment
            k_fold_results: list[tf.keras.callbacks.History] = cross_validate(
                 TransferLearningModel,
                 ds_train_and_valid,
                 k=kfolds.
                 epochs=EPOCHS,
                 batch_size=BATCH_SIZE,
                 model_kwargs={"dropout_rate": DROPOUT_RATE}
                 optimizer_kwargs={"learning_rate": learning_rate, "epsilon":

    epsilon_rate},
             search_results.append({
                 "learning_rate": learning_rate,
                 "epsilon_rate" : epsilon_rate,
                 "history_list" : k_fold_results
            })
    return search_results
```

Listing 7: Hyperparameter Search Regime II (Github)

Running the above search routine in a Jupyter notebook (Github) on our hosted compute provider, we obtain the following results:

 $10^{-1}$ 0.7405 0.7963 0.5744  $10^{-2}$ 0.7903 0.5678 0.7339 0.7466  $10^{-3}$ 0.7032 0.7938 0.7624 Epsilon  $(\epsilon)$  $10^{-4}$ 0.7744 0.7907 0.7466  $10^{-5}$ 0.7847 0.7969 0.7436  $10^{-6}$ 0.7865 0.7958 0.7075  $10^{-7}$ 0.7030 0.7823 0.7969  $10^{-8}$ 0.7919 0.7817 0.7493  $10^{-4}$  $10^{-3}$  $10^{-2}$  $10^{-1}$ Learning Rate

Regime II Hyperparameter Grid Search Results

0.50 0.52 0.54 0.56 0.58 0.60 0.62 0.64 0.66 0.68 0.70 0.72 0.74 0.76 0.78 0.80 Max Valid. AUC

Figure 4.7: Results for the Hyperparameter Search Regime II

Learning Rate	Epsilon $\epsilon$	Max Validation AUROC
$1.0 \times 10^{-3}$	$1.0 \times 10^{-7}$	0.7969321310520172
$1.0 \times 10^{-3}$	$1.0 \times 10^{-5}$	0.7969251374403635
$1.0 \times 10^{-1}$	$1.0 \times 10^{-1}$	0.7962859372297922
$1.0 \times 10^{-3}$	$1.0 \times 10^{-6}$	0.7957556545734406
$1.0 \times 10^{-3}$	$1.0 \times 10^{-3}$	0.7938481668631235
$1.0 \times 10^{-4}$	$1.0 \times 10^{-8}$	0.7918900847434998
$1.0 \times 10^{-3}$	$1.0\times10^{-4}$	0.7907490432262421
$1.0 \times 10^{-2}$	$1.0 \times 10^{-2}$	0.7903381884098053
$1.0 \times 10^{-4}$	$1.0 \times 10^{-6}$	0.7865246037642161
$1.0 \times 10^{-4}$	$1.0 \times 10^{-5}$	0.7846748034159342

Table 4.3: Top Ten Hyperparameter Options for Regime II

### 4.4.3 Final Hyperparameters

Thus, we have completed our hyperparameter search. After implementing and running Regime I and Regime II (Github), we were able to find the following hyperparameter options for our final model:

Batch Size	Dropout Rate	Learning Rate	Epsilon ( $\epsilon$ )
1600	0.20	$1.0 \times 10^{-3}$	$1.0 \times 10^{-7}$

Table 4.4: Final Hyperparameters for InceptionV3 with ImageNet Weights

The implementation, analysis, and search of our model hyperparameters are now complete. Now we are ready for the final evaluation of the complete model on our hold-out test set.

## **Chapter 5**

# **Results and Discussion**

As per our methodology (see section 3.6), we train a new model using the optimal set of hyperparameters. The following graph presents the training history of our final model:

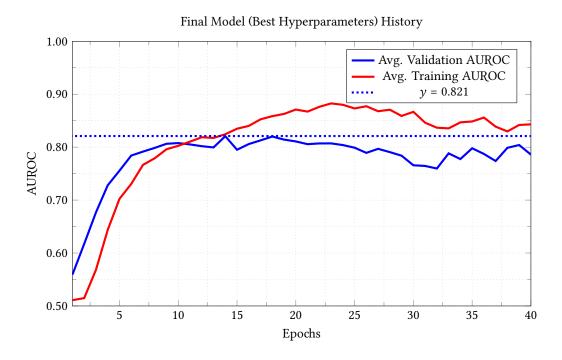


Figure 5.1: Training History of the Final Model (with Best Hyperparameters)

Using the best-performing model checkpoint at epoch 14, we evaluate the final model

on our hold-out test set. We obtain the following results:

Metric	Results
AUROC	0.890
Precision	0.880
Recall	0.409

Table 5.1: Performance characteristics of the final model

With this final piece of information, we may now turn and look back at the questions and endpoints (see section 3.6), and evaluate how our project turned out.

### 5.1 Does Transfer-Learning Perform Better than End-to-End Training?

The first question that we defined in our evaluation, was whether transfer-learning as a method is able to perform better than end-to-end training. Machine learning problems in medical imagery is often constrained by the challenge of small datasets, and ours is no exception. With only 3,000 radiographs bearing RUST labels, the very reason that we implement transfer learning is an attempt to mitigate issues that the end-to-end training of a larger model may face. Is our assumption validated? We can answer this question by examining the performance of our end-to-end model in subsection 4.2.1 versus the performance of our final model here. The end-to-end model achieved an AUROC of only 0.692, versus the AUROC our final transfer-learning model achieved of 0.890. Indeed, even our interim models built without hyperparameter optimisation achieved performances that were higher than the end-to-end baseline.

By examining the training and validation AUROC graphs of the end-to-end training implementation, we can see that the validation AUROC is highly erratic, both per k-fold cross-validation runs, as well as within the epochs of an individual training cycle. Furthermore, the end-to-end trained model overfits severely from the very beginning. These observations are strong indications that that end-to-end training is inappropriate for datasets of our order of magnitude. In contrast, if we look at the training and validation AUROC graphs of our final model, we can see a much smoother curve, and minimal overfitting. Hence, we may conclude that transfer-learning, as a method does perform better than end-to-end training, especially in the application of medical image processing where datasets are relatively small.

# 5.2 Do RadImageNet Weights Perform Better than ImageNet Weights?

The next question that we can answer, is whether or not RadImageNet weights perform better than ImageNet weights. We must keep in mind that whatever conclusions we can draw are only applicable specifically to our use case of inferring RUST scores from radiographs. In our case, it appeared that the ImageNet weights exhibited a higher level of performance, compared to the RadImageNet weights. This may be due to how the ImageNet dataset is fundamentally larger than the RadImageNet dataset (13 million images versus 5 million). The larger dataset size may have allowed the weights to be better placed to serve as general-purpose feature extractors. However, although the RadImageNet model achieved a lower level of performance overall, it did exhibit significantly less overfitting than the un-tuned ImageNet model. This keeps RadImageNet as an open avenue for additional exploration — and indeed, if there was more time, it would have been better to perform a hyperparameter search on the RadImageNet model as well, for a more robust comparison.

# 5.3 Is Transfer-Learning Able to Perform Persuasively, in the Task of Predicting RUST Scores?

Finally, we must ask ourselves: were we able to achieve the overarching goal of developing an AI model that is able to predict the RUST score of a fracture from a pair of radiographs? In our methodology, we explicitly defined a performance target of an AUROC of 0.80 or above. Our model, when evaluated on the hold-out test set, was able to achieve an AUROC of 0.890. Does this mean we achieved our goal?

One one hand, the AUROC of 0.890 is a strong result, approaching the same percentile as the other studies that we reviewed in our background research. Recall that the 2017 [10] study, which utilised a significantly larger dataset, was able to achieve an AUROC of 0.921. Likewise, the Lindsey et al. study utilising 31,490 radiographs achieved a AUROC of 0.967. [19] For a project utilising a data set containing only around 3,000 radiographs, an AUROC of 0.890 is certainly persuasive — at least as a proof of method.

However, the AUROC value is not the only metric that we must consider, when evaluating a model's performance. The model's precision and recall must also be evaluated, and although we achieved a precision of 0.880, our recall is only 0.409. This significantly weakens our results, insofar that it indicates that the true positive rate of our model is much lower than the AUROC would lead one to expect. Why is the recall so low, when compared to the Precision? One reason why this might be the case, may be due to the highly imbalanced nature of our dataset's class distribution.

As a *multi-class*, *multi-label* classification problem, our model must essentially tag a radiograph with the right set of cortex scores. However, the distribution of scores cannot be assumed as even — indeed, the dataset contains many more instances of non-union (i.e. non-healing fractures), than remodelled fractures. A low recall can be a symptom which indicates poor performance on a highly class-imbalanced dataset.

Hence, when considering both the precision and recall of our model, the ultimate performance still offers further room for improvement. This is not a bad conclusion to draw — since ultimately, the goal of this project is to validate the use of transfer-learning on a small dataset, demonstrate the feasibility of achieving results even with a difficult multi-class, multi-label problem, and ultimately offer the possibility for further inquiry — into the goal of building more advanced AI models that can characterise fractures using established clinical instruments.

### 5.4 Limitations and Further Works

Ultimately, it is important to stress that despite achieving an AUROC of 0.890, the model that we built in this study is still very far from a diagnostic or even research tool — particularly due to it's low recall. But instead, the results that we achieved should offer a path towards further questions and avenues of additional exploration. Some of these further works may include:

Is it possible to change the performance metric of the model from AUROC, to another metric that will allow us to maximise both precision *and* recall? One possible method that the author considered, but did not ultimately pursue, was the use of the F1 harmonic mean as the performance metric. The F1 harmonic mean is derived from the precision and the recall, and may potentially allow the training of a more robust model. Ultimately, the reason that this method was not attempted in the study, was because the F1 harmonic mean is not a differentiable function, hence gradient descent via backpropagation does not work for it. However, there exists alternative functions such as the F1SoftMax() which implement differentiable approximations to the F1 harmonic. This remains an intriguing direction for further inquiry.

Likewise, one of the decisions that we made was to develop one model for both anterioposterior and lateral views. This allows us to avoid bifurcating our limited dataset, but in retrospect it may be a decision that deserves further examination. Would an ensemble model, composed of two different neural networks, potentially avoid some of the issues the current project experienced with low recall? How would the RUST labels be encoded, in the context of an ensemble model? These questions remain to be answered.

Finally, further work can certainly be done in the domain of hyperparameter optimisation. In this study, we implemented a basic random search, followed by a grid

search. But given the provisioning of more compute resources, alternative methods like genetic algorithms may prove vastly superior to our current hyperparameter regimes. Likewise, the provisioning of further compute could allow us to optimise hyperparameters on both the RadImageNet model as well as the ImageNet model, potentially resulting in a fairer evaluation for both.

### Chapter 6

### Conclusion

In conclusion, this study explores the process and validity of building an AI model to automatically characterise fracture union, using the *radiographic union score for tibial fractures*. By taking pairs of radiographs as inputs, and inferring a RUST score as output, we hope to advance the intersection of medical imaging and machine learning. Although a great deal of literature exists in the use of AI and machine learning models for the automatic detection of fractures, and other anomaly-detection tasks, comparatively little work has been done to date in the characterisation of fractures — especially via an existing clinical instrument like RUST. Likewise, due to the comparatively smaller nature of datasets in the field of medical imaging, we faced challenges specific to small-datasets, which we hoped to overcome with the method of transfer learning.

We began by validating transfer-learning as a method, by training and evaluating our base model directly on the RUST dataset. Next, we evaluated two different set of weights (i.e. pre-training sources) for our application, being RadImageNet and ImageNet. After selecting the better-performing ImageNet weights, we conducted a hyperparameter search regime, ultimately finding the best dropout, batch size, learning rate, and epsilon. With the best-performing hyperparameters in hand, we proceeded to train a final model, which we then evaluated on our hold-out test set, yielding an AUROC of 0.890.

Using the data that we gathered in our experiments, we were able to examine them critically and contextualise our results in the basis of prior work in the literature. We acknowledge that although an AUROC of 0.890 approaches the same percentile as previous works, our model suffers from a low recall, despite the high precision and AUROC. However, the fact that we were able to achieve such a level of performance

even in light of a small dataset, indicates that this direction of inquiry is not without merit, and indeed better model design, the use of alternative performance metrics, and further data — could offer more significant performance gains.

Ultimately as radiography plays an increasing role not just in fixation, but also in rehabilitation, the use of AI can become a powerful tool in alleviating workloads for clinicians, and reducing healthcare cost for patients. By developing AI models which can accomplish routine tasks like scoring radiographs, we hope to empower clinicians in their practice, ultimately helping to fulfil the promise of medicine.

### Appendix A

### **Additional Materials**

### A.1 Project Code and Github Repository

All of the implementation details, model architecture, and data are made available in this project's Git repository (Github). Every code listing contains a link to the specific implementation, and different experiments also contain links to their corresponding Jupyter notebooks where the code was originally run. As a part of this project's commitment to reproducibility, all Jupyter notebooks are documented, and readers are encouraged to follow along and run the experiments for themselves. For further information, please see the repository README.md.

https://github.com/ShenZhouHong/radiography-ai-project/

### A.1.1 Initial Evaluation Models

Jupyter notebooks used to run the initial evaluations of LeNet 1998, InceptionV3 with end-to-end training, and initial transfer learning models:

https://github.com/ShenZhouHong/radiography-ai-project/tree/master/python/initial-evaluation

### A.1.2 Hyperparameter Search Code

Jupyter notebooks used to perform the hyperparameter search regime.

https://github.com/ShenZhouHong/radiography-ai-project/tree/master/python/hyperparam-search

### A.1.3 Analysis Notebooks

Jupyter notebooks used to analyse the raw data, process for insights and visualisations, and output CSV files:

https://github.com/ShenZhouHong/radiography-ai-project/tree/master/python/analysis

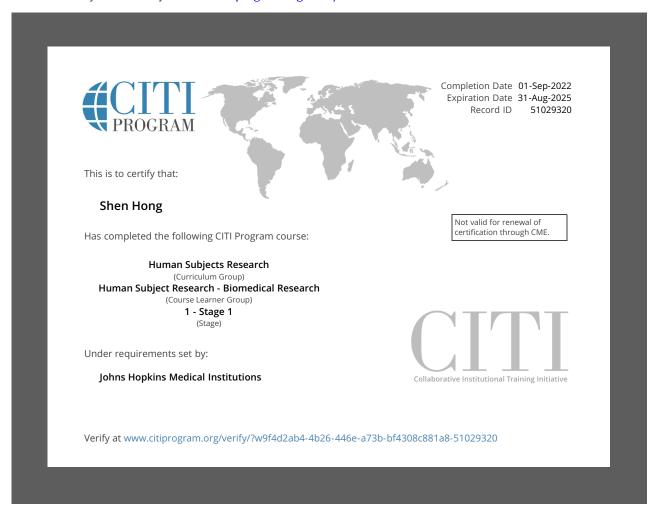
### A.1.4 Final Model Weights and Examples

Weights and instructions for the final model trained on the best set of hyperparameters, as well as example notebooks:

https://github.com/ShenZhouHong/radiography-ai-project/tree/master/python/final-model and the project of the

### A.2 Human Research Certification

Collaborative Institutional Training Initiative (CITI Program) certificate demonstrating compliance with human subject research. Verify authenticity at: <a href="https://www.citiprogram.org/verify/?w9f4d2ab4-4b26-446e-a73b-bf4308c881a8-51029320">www.citiprogram.org/verify/?w9f4d2ab4-4b26-446e-a73b-bf4308c881a8-51029320</a>



### A.3 Project Proposal Presentation

Project proposal presentation given on 2022-01-11, at the Johns Hopkins University, East Baltimore Campus.

# Evaluating Fracture Healing with Artificial Intelligence

Using Transfer Learning to Predict RUST Scores from Radiographs.
A Major Extremity Trauma Research Consortium Project.

Shen Zhou Hong <shong@jhu.edu>









### **Introduction and Table of Contents**

- Table of Contents:
  - §1. Background Information
  - §2. Project Design, Methodology, & Endpoints
  - §3. Current Progress, Roadmap, & Challenges
  - Q&A Session.

### **Background Information**

2022-01-11 METRC, Johns Hopkins 2 2022-01-11 METRC, Johns Hopkins

### Using AI to Evaluate Fracture Healing

- We want to develop an AI that can infer RUST scores from radiographs.
- RUST: Radiographic Union Score for Tibial Fractures. Also used in other long bone fractures.
- RUST measures the progression of fracture healing via callus formation and bridging.



Image source: D. B. Whelan, M. Bhandari, D. Stephen, et al. 2010

22-01-11 METRC, Johns Hopkins 4 2022-01-11 METRC, Johns Hopkins

2022-01-11

2022-01-11

### Why is this project useful?

- The AI model can automate the analysis of archived study data.
- RUST scores have good biomechanical correlation, can serve as guidance for rehab.
- Project is novel: prior work in AI-radiography mainly focused on fracture detection.

022-01-11 METRC, Johns Hopkins

### Prior Research in AI-Radiography

- MURA: Anomaly detection. 40.5k radiographs, achieved AUROC of 0.929
- Lindsey et al: Anomaly detection & localization.
   31.0k radiographs, achieved AUROC of 0.967
- Kim et MacKinnon: 1.3k radiographs, achieved AUROC of 0.954

2022-01-11 METRC, Johns Hopkin:

### Why has nobody done this before?

- Most individual institutions do not have access to datasets large enough to perform training.
- Most large radiography datasets do not have specialized, *evidence-based*, *adjudicated* labels.
- Most AI research organizations do not have access to medical research organizations.

METRC, Johns Hopkins

METRC is uniquely positioned to conduct this research.



2022-01-11 METRC, Johns Hopkins

### Design, Methodology, & Endpoints

### AI Models versus Statistical Models

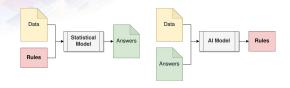


Image source: METRC 2022

METRC, Johns Hopkins

### METRC Radiography Datasets

METRC. Johns Honkins

RetroDEFECT: 741 radiographs

• OUTLET: 707 radiographs

• PAIN: 370 radiographs

• PACS: 195 radiographs

• Total: ~2,013 radiographs

• Our study architecture is data-constrained.

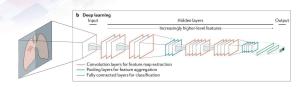
### Model Architecture

- *Transfer learning*: train an AI model on a large, general-purpose dataset. Then *fine-tune* on the smaller, task-specific dataset.
- Data Augmentation & K-Fold Validation
- We will use ImageNet and MURA as base models for transfer-learning.

122-01-11 METRC, Johns Hopkins 12 2022-01-11 METRC, Johns Hopkins 1

2022-01-11

### Anatomy of a Deep Neural Network



age source: Hosny, Parmar, Quackenbush et al. 2018

METRC, Johns Hopkins

### Endpoints

- Model performance measured via AUROC: Area Under Curve (of) Receiver Operating Characteristic.
- Endpoint 1: Model AUROC > 0.50
- Endpoint 2: Model AUROC > "naive" ConvNet
- Endpoint 3: Model AUROC > 0.75

METRC, Johns Hopkins

### Progress, Roadmap, & Challenges

### Roadmap and Current Progress

- 2022-11-15: Complete background research
- 2022-12-01: Study and DNN architecture design
- · 2022-12-12: Initial exploration of METRC datasets
- 2023-01-02: Preprocessing of METRC radiographs. (We are here)
- 2023-01-31: Initial model development
- 2023-02-27: Fine-tuning, hyperparameter search.
- 2023-03-31: Deadline: Impl. & Analysis
- 2023-05-02: Deadline: Poster Presentation at UoL Goldsmiths
- 2023-05-12: Deadline: Submission to UoL Goldsmiths

2022-01-11 METRC, Johns Hopkins 2022-01-11 METRC, Johns Hopkins

### Challenges

- · Dataset validation and "cleaning."
- · Combating model overfitting.
- Programmatically parsing data from REDCap.

### REDCap Libraries and Tools





- REDCap Branch Parser: https://github.com/metrc /redcap-branch-parser
- REDCap Schema: https://github.com/metrc /redcapschema

2022-01-11 METRC. Johns Hopkins 2022-01-11 METRC, Johns Hopkins

### **Future Pathways**

- Heat-maps and "Explainable AI"
- Further assessment of model performance with human orthopedic specialists/surgeons
- Collect more RUST data. Train a more robust model, conduct serious evaluations of using AI as a diagnostic tool.

The End. Thank you for your attention!

2022-01-11 METRC. Johns Hopkins 2022-01-11 METRC. Johns Hopkins

### A.4 Poster Presentation

Final project poster presentation, given on 2023-05-11 at Goldsmiths College, University of London.

— Avg. Validation AUROC
— Avg. Training AUROC
v = 0.821

to find that ImageNet weights had slightly better performance on our eters, we were able to yield a Final Model with a k-fold cross-validated validation AUC of 0.821 (see graph above). This best-performing model was then evaluated upon our hold-out test set, yielding a AUC of 0.891. The performance achieved by this model is promising, especially given

By evaluating both ImageNet weights and RadImageNet, we were able baseline model. Using ImageNet and the best-performing hyperparam

JOHNS HOPKINS Goldsmiths

ng Transfer-Learning to

# Shen Zhou Hong. Supervisor: Georgios Mastorakis, Renan Castillo

Final Model (Best Hyperparameters) History

# Predict RUST Scores in X-Rays

data. To mitigate this issue, we use the technique of transfer learning to train an AI model first on a large, general-purpose dataset, before set. Small models with few parameters have limited ability to generalise. while large models with many parameters struggle to converge on limited In partnership with the Major Extremity Trauma Research Consortium challenge in this project is dealing with the limitations of a small data transfer-learning said model on our smaller radiography dataset.

Methodology & Datasets

0.80 0.70

> of the precision/recall graph. We use the InceptionV3 CNN architecture as our base model with the Adam optimizer. We evaluate two different size, dropout rate, learning rate, and epsilon. This hyperparameter search is RUST labels are one-hot encoded into 18-value sparse vectors, and model performance is assessed via the AUC (Area Under Curve) pre-training datasets for our base model: Stanford's ImageNet dataset (14.0 million images), and Cornell's RadImageNet dataset (5.0 million el, we proceed to search for optimal hyperparameters, iterating on batch images). RadImageNet is a smaller dataset but offers potential advantag es by being domain-specific to our task. With the better-performing mod

- Regime I: Random Search on Batch Size & Dropout (see bottom fig.)
- with k = 6. After finding the best set of hyperparameters, an evaluation of the final model is performed using k-fold cross-validation, and then the best-performing fold is evaluated on the hold-out test set. Each trial in the search space is evaluated using k-fold cross-validation Regime II: Grid Search on Learning Rate & Epsilon

practicality of using

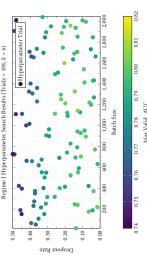
loads. Is it possible to train an AI model to automatically infer RUST transfer-learning as a technique to build an AI model on a limited dataset. to automatically predict RUST scores from radiographs of fractures.

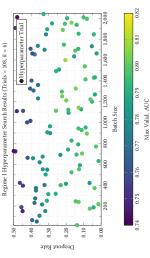
scores from radiographs? This study

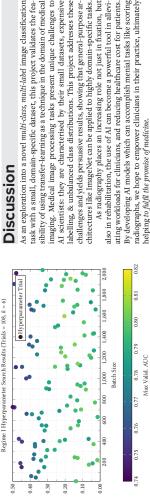
The precision achieved over the one-hot encoded RUST labels on the hold-out test set is 0.880. However, the recall is only 0.409, high-lighting difficulties with the highly imbalanced nature of our dataset. This

the small size of the dataset that it is trained on.

# maging. Medical image processing tasks present unique challenges to As an exploration into a novel multi-class, multi-label image classification sibility of using transfer-learning as a technique in the domain of medical AI scientists: they are characterised by their small datasets, expensive labelling, & unbalanced class distributions. This project addresses these challenges and yields persuasive results, showing that general-purpose artask with a small, domain-specific dataset, this project validates the fea may indicate that the model performance degrades with edge cases Discussion eter Search Results (Trials = 108, k = 6) Regime I Hyperpa







Project Report: https://s.hong.io/project.pdf

As radiography places an increasing role not just in fixation, but

# Repository & Data URL: https://github.com/ShenZhouHong/radiography-ai-project

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mation, bridging, & union. Whelan et al's Radiographic Union Score for Long-bone fractures are serious injuries which require longterm care and

Tibial Fractures (RUST) is a 12-point clinical instrument

rehabilitation. After fixation, a fracture must be monitored for callus-for-

union from the antereoposterior & lateral radiographs of a fracture. However scoring radiographs for RUST is a tricky process, requiring the attention of a trained orthopaedic clinician, potentially increasing work-

used to assess

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