

# Contents

<b>1</b>	<b>Abstract</b>	<b>1</b>
<b>2</b>	<b>Introduction</b>	<b>2</b>
<b>3</b>	<b>Fundamentals on paleoecology</b>	<b>3</b>
3.1	Basics on ecology . . . . .	3
3.2	Paleoenvironmental reconstruction . . . . .	3
3.3	Composition of mammal teeth . . . . .	4
<b>4</b>	<b>Deep Neural Networks for Optical Character Recognition</b>	<b>5</b>
4.1	Deep Neural Networks . . . . .	5
4.2	Training neural networks . . . . .	5
4.2.1	Loss functions . . . . .	6
4.2.2	Evaluating model performance . . . . .	6
4.3	Architectures . . . . .	7
4.3.1	Convolutional layers . . . . .	7
4.3.2	Transformers . . . . .	7
4.3.3	Autoencoders . . . . .	7
4.4	Techniques and heuristics for improving performance . . . . .	7
4.5	Transfer learning . . . . .	7
4.5.1	Foundation models . . . . .	7
<b>5</b>	<b>Related work</b>	<b>8</b>
5.1	Approaches to digitization of handwritten archival data . . . . .	8
5.2	Approaches to character recognition with small target domain datasets . . . . .	8
<b>6</b>	<b>Experimental setup</b>	<b>8</b>
6.1	Data description . . . . .	8
6.1.1	Notes on creating the dataset . . . . .	8
6.1.2	Unicode characters used for data labeling . . . . .	9
6.2	Data preprocessing . . . . .	9
6.3	Methods: base models and transfer learning techniques . . . . .	10
6.3.1	Problem formulation . . . . .	10
6.3.2	Base model selection . . . . .	11
6.3.3	Transfer learning method selection . . . . .	11
<b>7</b>	<b>Results and discussion</b>	<b>11</b>
<b>8</b>	<b>Conclusions</b>	<b>11</b>

## 1 Abstract

Keywords: Optical character recognition, Few-shot transfer learning, Vision transformers, Paleontological databases

## 2 Introduction

broad research area: paleontology: data analysis on fossil finds dig fossil from the ground. identify: which bone, species, time. write this on a field slip, a slice of baking sheet like paper analysis: take set of fossils, use methods for deducing eg climate, habitat, vegetation why this is interesting reactions of ecosystems to climate change what ancient worlds were like how ancient humans lived mass extinction events

now we have a stack of baking sheets in a room in kenya. paleontologists from all over the world want to solve climate change, among other problems big data methods would explode what paleoecology can do so we need to put the baking sheets on the computer to do analysis on big data what is the topic of my work? the baking sheets contain weird characters that a normal reader cannot read. my topic is to read them

given scanned images and data with bounding boxes of sentences and words where tooth denoting words are badly read, how can tooth element recognition results be improved? Goal is to have both this is what the element / nature of specimen column says (eg example here) and what/which teeth are found in this specimen in standardized format (eg example here)

limitations: compute (no HPC utilized), small amount of labeled data, scope is masters thesis so very advanced techniques are not feasible.

relevance of this work: KNM is able to have way more precise dental element markings to other catalogues: previous project + this a complete solution to digitizing the handwritten data relevance of this work: any field that does: - ocr on unconventional characters - ocr where each character has a multivariate output (eg. this is an a. it is underlined could be letter and underlined /not underlined)

The direct impact of this work is an improved precision of the tooth element entries in the digitized fossil catalogues of the National Museum of Kenya, but the results are applicable to a wider domain of problems. Intuitively, the results are directly applicable to other fossil archives using similar notation: only a fine-tuning of the models to the new archive data is necessary. For other handwritten archives, the results presented can be used to improve recognition accuracy, especially in cases where the data contains characters other than latin letters or arabic numerals. Additionally, this work presents a potential solution for when the target character set can be expressed with multivariate output data. This could, for instance, be handwriting with occasional underlinings, where the bivariate output could be the letter and a boolean variable for whether the character was underlined.

The rest of this thesis is organized as follows. First, the necessary background theory is presented. For deep neural networks the following concepts are introduced: the basic network structure, how training is conducted, basic building blocks of character-recognizing network architectures, performance-improving heuristics, and transfer learning. For paleoecology, the background covers foundational ecological laws followed by a brief introduction to methods used in paleoenvironmental reconstruction, especially focusing on inferences from tooth data. As the last background section, the composition of mammal teeth is presented. Second, related work is presented, both on handwritten archive digitization and transfer learning with character-recognizer models. Next, the experimental setup is introduced, covering dataset creation, labeling and data preprocessing, followed by base model and transfer learning method selection. After this, results of the experiments are presented and discussed. Finally, the work is concluded.

### 3 Fundamentals on paleoecology

Nature is highly complicated -> models, approximate models and assumptions enable drawing conclusions from known distributions of species.

each assumption / model can be questioned but they hold as a rule. only models briefly presented here, all statements here can be questioned to some extent

idea: what fossil/dental data can be used for. how fossil/dental data is used

to highlight why accurate, fine-resolution (ie specific) and large magnitude of fossil, esp dental fossil data is genuinely useful.

chapter overview: review ecology and assumptions the analysis is based on. then, short overview of main techniques for paleoenvironmental reconstruction, the main application area of fossil data. last, mammal teeth row is presented to introduce terminology present in the data.

#### 3.1 Basics on ecology

basic laws: theory that the data analysis relies on

Tolerances and niches (fundamental + realized): basis for environmental reconstruction [3] ch 2

tolerance = range of an environmental variable that is hospitable for the species, eg. imaginary small mammal (come up with some imaginary name) can live when avg temperature is +10-+15. Niche = set of tolerances the species has. fundamental niche = possible environments for the species, realized niche = where it lives. center of tolerances is better than the edge (ch2)

main assumption uniformitarianism (the fact that tolerances constraint things has not changed)

niche conservatism: Assume that nearest living relative has same tolerances now -> get past environment (ch3, lyman 2017),

this presents a mapping from taxa to environmental variables -> basis of analysis of past environments.

modern alternative to this: transfer functions: mappings from taxa data to environment learned using machine learning / statistical models (ch9, birks 1995)

benefit instead of tolerances/niches: they have subjective interpretation problems (book ch 9)

esp. teeth: dental ecometrics = inference of transfer functions given dental data (ch9 liu et al 2012)

next, turn to how to solve the problem of information to environment given the taxonomic information to environmental indicators mapping

#### 3.2 Paleoenvironmental reconstruction

why: get information of what is to come with climate change (faith ch2)

definition of the problem: what ancient habitats were like and what changes they underwent at which times (ch2)

overview main techniques: presence/absence, abundance, taxon free, diversity based, size clines

presence absence (ch5): dataset is list of taxa that are present. absence is also indicator but worse since might be that the species just was not preserved / found. two approaches: fix location or fix set of species. fix species: find where this set of species lives now (climate maps & areas of sympatry), this indicates that historical locality had this climate. place-fixing: analyze how which species show up in this place changes over time, reduce species showing up data to lower dimension (like pca), eg how many warm-climate vs cold-climate species show up, this gives ideas on changes in climate over time

abundance get relative NISP (number of species in sample): percent of species in sample is of this taxon (grayson 1984b, ch6) do like presence absence but weigh signal according to abundance and assume abundant species lives more toward center of tolerance grayson 1981: fossil accumulation affects datasets -> use with caution

taxon free cornelis van der klaauw 1948 (p 160) relation of traits of animals and livelihood = ecomorphology: eg what the animal eats also eg how diverse the place is. for environment mostly diet -> what plants grew and habitat -> climate etc helps with not having to assume that closest relative tolerances were the same. (ch7) eg usage of teeth: dental microwear. calandra and merceron 2016: analyze miniscratches on the surface to get diet to get vegetation and climate oma: to conduct this you need many samples of the same bone need to know exactly which bone it is since different teeth used for different things

diversity and size clines as environmental indicators andrews et al 1979: how diverse also has a mapping to environment eg tropical is more diverse than arctic coarse indicators of the environment lyman 2008: strong sample size effects: bigger sample is more diverse so to do this you need equal size samples from all time periods analyzed size clines basic idea: coarse indicators for size of bone -> environment eg larger libs warmer climate (mayr 1970) also dental measures correlate with environment, eg mandibles (faith et al 2016). for that you need sufficient tooth samples and correct identification which tooth it is

end lesson: sample sizes and data precision are important. therefore it is superimportant to get more data / improve accuracy, which is the main goal of the digitization effort. scale sense: faith lyman ch 4 said 1000 is solid 10 000 whopping sample size dataset in question in this work has 90,000 so the value is pretty clear

### 3.3 Composition of mammal teeth

Fossils occur when animal / plant remains are deposited in a sediment in a way that preserves some part of its original form. Since teeth are the hardest material in animals, large fraction of found parts are teeth. Fossil finding is followed by identification to most specific taxon possible largely a technical skill (ch5), teeth are identified down to type and number, how manyeth the teeth are, counting from center to edge or other way round?? specimen can be either one tooth or fragments of the jaw bone where there are multiple teeth (markings like M1-3)

from [4] what teeth are composed of

the jaw bones lower jaw bones: mandibles permanent and deciduous (D), nonpermanent "milk" teeth (laita vaan jos löytyy d-hampaita)

right and left sides are always symmetrical, denoted simply L or R or Lt or Rt or left or right. left is left looking from the animal, not the observers perspective Identity also causes that sometimes tooth fossils are misidentified to the wrong side and corrected (ei lähteestä vaan nähty datasta koska l ja r on sutattu aika monta kertaa ja vaihettu

four classes, front to back: three incisors (I), one canine (C), four premolars (P), three molars (M). top bottom left right. top/bottom noting upper jaw as superscript lower jaw as lower script, purpose: incisor -> catching, canine -> stabbing / killing prey, molars are for chewing. premolars are bit like canines bit like molars, function varies lot between taxa including holding, cutting and chewing. also form and number of each present changes between taxa. sometimes lower jaw as line on top and upper jaw as line on bottom, sometimes both are used: upper script number with line on bottom. Line is "the other jaw" if there are less of a type of teeth eg two premolars, they might be no 1 and 2 or no 3 and 4

## 4 Deep Neural Networks for Optical Character Recognition

separate problems: character classification (easy, kNN, SVM), reading variable-length text (harder)  
[7] introduce the problem of ocr, example: fossil catalogue

### 4.1 Deep Neural Networks

- neurons and activation functions. maybe examples of activation functions: relu, sigmoid, softmax
- what is a neural net weights in layers: floating point numbers, grouped in groups activations: connections between weights, nonlinear scalar to scalar functions

- feed forward network computes output from input with the feed forward. you have an input, bunch of numbers then, you compute a linear combination and pass that through an activation function

$$h_d = a \left[ \theta_{d0} + \sum_{i=1}^{D_i} \theta_{di} h_{d-1} \right] \quad (1)$$

hd, hd-1 are hidden unit values. d is layer index. thetas are weights D is width of previous layer  
[9]

a is any nonlinear function, simplest is the rectified linear unit relu,  $x = x$  if  $x \geq 0$ ,  $x = 0$  if  $x < 0$ . hidden units are taken as inputs to next layer and next and next. thetas = weights largest, deepest: hundreds of layers, hundreds of millions of weights then last hidden unit output is the output of the network. tadaa.

- universal function approximator the theory of the universal approximation capacity: this algebraic construct, given correct weights, activation functions and structure, could approximate any mapping from input to output. note: input/output dimension can be anything

examples from ocr relevant in this case input is always image ie 2d matrix if grayscale or 3d tensor if rgb image.

classification problem, softmax activation image of tooth sample (letter+number) to four classes, M P I or C. input: image, output: four probabilities that sum to 1, probability this is a m p i or c. sum to 1 is achieved with softmax. output is largest probability.

multilabel classification problem: image of tooth sample. output: first MIPC, second if it is upper or lower jaw output can be two arrays, one like before, other a 0-1 probability for upper jaw. upper if this  $\geq 0.5$

present sequence to sequence learning: more complex case: image of sentence output: text on image, variable length. here output layer is a more complex structure of probabilities for each position in the result sequence.

insert image example: inputs and outputs of different learnable functions with neural networks

### 4.2 Training neural networks

u have encoded the structure, starting weights. data, input/output pairs. training = process that adjusts weights so that network becomes a good approximator

1. take part of data as training data 3 divide to batches, common batch sizes are exponents of 2, 2-32 usually 4. pass a batch through the network. get output 5. pass output and correct to loss function: map from output, correct to scalar, 0 is good, large value bad. 6. Compute loss function

gradient with respect to network weights using automatic differentiation. algorithm to get this in code is called backpropagation because it moves backward in the network 7. gradient informs how to adjust weights so that loss should decrease. adjust weights in this direction by preset amount called learning rate. Optimizers are algorithms that determine the specifics of "moving in the direction of decreasing loss" most common stochastic gradient descent (inserts randomness to movement) and adam (uses previous iteration movements known as momentum in process to make movements more smooth) 8. run batches until out of data = epoch. run many epochs, stop according to stopping condition that is known to be a state when the network weights are good. goal: reach global minimum of loss function, difficult! would be perfect approximation 8. test on unseen test data to see generalization performance

lots of details on this process, rest is about that

#### 4.2.1 Loss functions

more specific: how do you map output and label to scalar value describing how good the output was? ocr point of view

Loss function is a function from model predictions and ground truth labels that describes with a single scalar value how good the match was, low number describing a good match [9]. These functions are constructed to be equivalent with maximum likelihood solution, think the model would output a conditional distribution of outputs,  $p(y|x)$ . each ground truth label in the training set should have a high probability in this distribution. Product of all these probabilities is called likelihood. Find parameters that maximize the likelihood of the training data set. Loss functions are derived so that parameters bringing loss to zero is equivalent to the parameters with maximum likelihood. Derivations are out of scope.

- cross-entropy loss kullback-leibler divergence of correct conditional probability and conditional probability parametrized by current model parameters. (show formula, 5.27), correct is not dependent on parameters so is omitted. show 5.29, what is left from that (until here from [9])

then: how cross-entropy loss is computed used for classification problems. eg. is this letter in this image an 'a' or a 'b'. correct probabilities eg .1 and .9 for a or b. model says .2 and .8. discretized cross entropy computes it as  $-.2 \log .1 - .8 \log .9$ , log in base 2.

word detection models: have a predefined vocabulary, layer for probability of each word. loss is cross entropy for these probabilities compared to target probability distribution, where correct word has probability 1 and all other have probability 0

- CTC loss

maybe

#### 4.2.2 Evaluating model performance

common in character recognition: accuracy, f1 sometimes also used explain these and differences

accuracy: correct out of those classified

why one uses precision and recall instead of accuracy sometimes simple accuracy is not a good measure, eg cancer screenings: 1 in 500 has cancer. then a dummy saying everyone is healthy would be correct 499/500 of the time, accuracy 99.8%. measurements anyhow relevant would be 99.8-100, inconvenient range.

precision: fraction of positives identified. dummy would have precision 0 recall: out of those noted as positives, fraction of correct ones. dummy would have 0 too f1 is an average to lump these together in one number

going forward: only consider simple accuracy scores as none of the classes are very rare in the fossil case.

## 4.3 Architectures

### 4.3.1 Convolutional layers

convolution (cross-correlation) max pooling / average pooling operations

### 4.3.2 Transformers

- self-attention - multi-head self-attention - tokenizing and the cls token

### 4.3.3 Autoencoders

- encoder/decoder  
sequence to sequence [11]

## 4.4 Techniques and heuristics for improving performance

data augmentation

## 4.5 Transfer learning

this sect from [8]. they initially formalized the problem and uniformized the terminology, before that multiple terms (ref? price/goodfellow?)

basics: what it is

why: save compute and data labeling (usual constraints in ml model building [5])

task (inductive) transfer: tasks differ and domain (transductive) transfer (only data differs)

task transfer relatedness + the more related the better results, negative transfer ie things made worse if things differ too much.

initialize weights to those that suit a related task, it is assumed that the starting point is already very good catastrophic forgetting = forgetting the previously learned after finetuning

broad methods for inductive transfer: feature representation transfer, parameter transfer; priors or hyperparameters of model are assumed to be shared, instance transfer: reuse source data (not considered), relational-knowledge-transfer: when data is not i.i.d (not considered here) ont considered because does not apply here

feature representation transfer seeing model as model doing the feature engineering: all but last layer are like feature extraction that allows a linear model to be fit from features to output. assumption for last layer transfer: reusing the feature extractor should bring good results with little computational cost

basically nowadays it is usually insensible to ever train from scratch (NNDL lectures)

### 4.5.1 Foundation models

- generalist models - large unsupervised training data sets

Idea: since transfer learning source task performance is not important, target task is the main point [8], you can come up with some nonsensical task such as map image to the same image to get

unsupervised task, train massively on massive compute, and use the model for varieties of tasks.  
idea: do as little as possible in the target domain

## 5 Related work

Search strategy: few seed papers and snowball search. Related conferences: Frontiers in Handwriting Recognition

Notes on choices: there is math OCR and music OCR, but they use large datasets and no transfer learning, they don't suffer from the limited target data problem. Also, the problem domain is too different from my problem to be informative.

### 5.1 Approaches to digitization of handwritten archival data

obvious solution: sit down and type

### 5.2 Approaches to character recognition with small target domain datasets

method: snowball search. for a paper to be accepted for use here, i set up these requirements: lists best accuracy clearly (percent, which dataset, number of classes) specifies the base model used (architecture and source domain dataset) specifies how transfer learning was conducted (not just used transfer learning) then reputable venue and acceptable quality of text and presentation, super subpar text was a frequent reason to skip a paper, they also often lacked relevant details

## 6 Experimental setup

### 6.1 Data description

has been done by different annotators, no logs on who logged what, everyone had a bit different style of notating. also no clearly defined standard for notating specimens. so might be that actual data used will have characters or words not present in any data, causing errors.

Identity also causes that sometimes tooth fossils are misidentified to the wrong side, seen in data with smudged over l's and r's

Catalogues have lines between entries. challenge for the model to not mark these as underlined in these cases the line is long and spans the entire image, so it should be distinguishable from a single underlined character.

#### 6.1.1 Notes on creating the dataset

Hand-labeled

Data was extracted from scans by getting bounding boxes from Azure Vision API, finding the correct column (nature of specimen or element), and cropping the image according to bounding boxes.

Non-tooth samples were not discarded since they contain bone fossil related words and good samples of the handwriting style of this dataset.

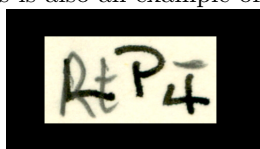


smudged-over "L" was labeled as "R", and other way around: it seems that later someone found it was the opposite side after all. Hope of this is that the model would learn to map "messy L" as "R". smudged "left" or "right" was not noted as the opposite as there were too few such samples.

Superscript seems much more rare than lower script

Data was labeled not by individual characters but as full tooth descriptions to preserve context where tooth special characters are more likely to occur

Some have been corrected by writing on top and thus are very hard even for humans to read, this is also an example of smudged-over correction:



### 6.1.2 Unicode characters used for data labeling

explain: unicode has graphemes with code points. eg a is one grapheme one code point, à is one grapheme two code points (dot on top and the letter). the top thing -like characters will be called "modifiers".

markings contain letters and numbers with no line, line on top or line at the bottom. Each character can be lower- or upper script. The modifiers used are: macron with lower (Ā) and upper variant.

Unicode [12] has characters that are for example upper script, but these were not used for two reasons:

- lower and upper script character set is incomplete for this purpose (eg 3 with upper macron and lower script needed)

- from the model perspective 3 and 3 are no more similar than A and B, however, three combined with lower script modifier and 3 with upper script modifier all contain the same unicode character 3 with only the modifier changing. The problem here is that there is no lower or upper case modifiers in unicode. Therefore, the caron (Ď) was chosen as the lower script modifier, and the circumflex accent (Â) as upper script. These were chosen since the arrow-like modifier pointing up or down is maybe the most logical placeholder for the missing modifier. More traditional workarounds of missing upper or lower script, the underscore "\_" and separate caret character "^" were not used to keep one unicode grapheme represent one character on the page. Also on the other hand using one modifier for all lowercase characters allows the model to understand that there is a similarity between all lowercase characters. The intention is that one idea about a character is encoded as one code point, so that the model can learn the mapping from the image of the character to the code point combination (until now already in thesis text) —

also: some annotators used / instead of line. left to / -i upper, right -i lower. unknown/unsure noted with x with macrons on top/bottom annotate with up/down macron

if model is toothroot classifier + tooth reader -i remove fractions notes

## 6.2 Data preprocessing

grayscale, flipping to back background white foreground

I should especially do random cropping since that occurs in real dataset!

I could do some border contrast decrease center contrast increase because border noise is present and usually less relevant

all these: test first without, then add preprocessing step, then re-test

i am able to get labeled MPIC data by getting azure outputs where the letter is mpi or c

## 6.3 Methods: base models and transfer learning techniques

### 6.3.1 Problem formulation

different types of defining inputs and outputs, discussion on what to choose

the sequence or character per character recognition question: sequence is a mapping from image to variable-length phrase character per character approach, inspired by [14]: train a classifier: is this word a tooth or not? then give non-tooth to the untuned trocr, which works very well. Then few shot transfer learn a classifier from an image with only a tooth marking (letter and one number) to tooth. Possibly extend classifier to be able to recognize multiple teeth (eg M1-3). Target could be multivariate: first would be tooth (which i1-3,c,p1-4,m1-3), second would be jaw (upper, lower, unknown), third side (left, right, unknown). Separating 'l/r/lt' from the letter+number tooth notation is fairly trivial: noncursive handwriting can be separated by finding a vertical line where there is no black. Image processing: convert to black and white, then find x coordinate with no black and split there.

sequence benefits: adaptable to many kinds and lengths of input, possible to get good inferences for surprising marking styles sequence bad sides: finetuning just one layer on 80 training images for two epochs took about 15 minutes -; all hyperparameter optimization etc is out of question with this heavy training. Also: not domain adaptation (adapting same task to different dataset), but task adaptation ie. the target set of characters has changed. Encoding this to the large encoder decoder transformers would require rewriting parts of the preprocessor and model which is too complex given the level of this work.

character by character benefits: feasible given available data and computing resources possible to encode the classes (ie, teeth) classifying characters has been essentially solved, easy problem also classifying to tooth or not tooth should be easy -; focus on model ensemble with tooth or not classifier + trocr + tooth classifier

multilabel class or classifier chains: multilabel is difficult because the basic case has just set of labels and you can have any number of any of them. we have 4 types of classes with dependencies (no 4th canine) so chains make sense, as constrained multilabel classification with class groups is a complex, nonstandard construct [13]. so, mini model approach

1. tooth or not: azure output text matches regexp letter and number or 'c' or 'C' -; 1/0 train using MPIC images 2. MPI classifier: tooth type (regexp already says c is c), upper lower classifier: 1/0 3. get number (1-4) from azure output, is generally correct by visual inspection

not inspired from any paper but my own idea, felt like a sensible way to formulate the problem, so we have for all models a well-solved well-defined basic classification problem.

reason for this choice: it encodes most prior knowledge in the output structure, use as much from azure as possible

priors: there is also the azure output for the tooth. it is likely to be very good but not constrained to the MPIC case. include the azure label as one input variable why we went for the simplest problem out there because the part of data that can be digitized with the tightest constraints and the simplest models are most accurate. Do that first, deal with the remaining later

### 6.3.2 Base model selection

initial sequence learning attempts done with [6], proof of concept so no real exhaustive comparison literature review

Chosen public dataset was EMNIST [2] because it is the closest to my problem: a large dataset of labeled letters and numbers.

According to [1], this model was the best: [10], was chosen as the base model. they trained on emnist-balanced, so no difference between upper -and lower case letters. not a problem for us since in the handwritten catalogues, upper or lowercase was not used to distinguish upper and lower jaw

### 6.3.3 Transfer learning method selection

Priors. base model already knows the output should be a word, eg "jdaskjfkds" is a highly unlikely correct answer. Bone notation has a very small subset of possible english words, eg. the word "beach ball" cannot ever be a correct answer for a reading -i these do not apply if we do character classification

## 7 Results and discussion

table of results could contain: rows: base models tested columns: transfer learning method

transfer learning methods could include

just training further (ie baseline, dont do anything special)

train the last layer (inspo from [14], test hypothesis of reusing feature extractor) what else? find from literature!

train unsupervised autoencoder on the target data (reduces domain transfer distance), transfer to supervised classification task

## 8 Conclusions

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