Martin Pettersen

Segmentation of MR Images Using CNN

Ålesund - 01.10.2019



Abstract

The need for digital segmentation of human body parts is growing and having a digital twin of the body part with an injury, or a planned surgery can help people understand that body part better and be prepared for the tasks to come. To segment bodyparts are a helpful tool to both understand and examen the human body. You can add a segmented body part in a 3-dimensional world, look at it from different angles and discover something you could not have seen from a 2-dimensional view.

During the work of this thesis, there was created a Python-script of a Convolutional Neural Network (CNN) using Keras with TensorFlow as backend. This CNN is a U-Net inspired network created to classify each pixel in an MRI scan of a knee joint. It classifies bones, PCL, ACL which all are tissues located in the knee joint. The results are promising an there where several discoveries in the result of this thesis. The weighted loss function is a necessary function to classify lower weighted tissues such as the ligaments (PCL and ACL). And it also showed that the commonly used inputs not necessary are the best inputs for this network. The single input image T1 performed slightly better than the known method by using all of the images as an input.

Sammendarag

Anvendbarheten av digital segmentering av kroppsdeler er en økende trend, og det å ha en digital tvilling av skadede legemer, eller en planlagt operasjon kan bedre forståelsen av den kroppsdelen bedre, og forberede partene involvert. Segmentering av kroppsdeler er et nyttig verktøy for å bedre forstå og undersøke menneskekroppen. Segmenterte legemer kan vises som 3D-modeller, og manipuleres slik at den kan ses fra forskjellige vinkler, og muligens oppdage noe som ikke var like synlig i det originale bildesettet.

I løpet av masterprosjektet ble det utviklet et Python-script av et Convolutional Neural Network (CNN) ved hjelp av Keras med TensorFlow som backend. Dette CNN-nettverket er inspirert av U-Net og lagd for å klassifisere hver enkelt piksel i MRI-skanninger av kneledd. Den klassifiserer ben, fremre-, og bakre korsbånd, som alle er vev som ligger i kneleddet. Resultatene er lovende, og det ble avdekket flere gode resultater. En vektet tapsfunksjon er nødvendig for å klassifisere vev som opptrer sjeldnere, slik som korsbåndene. I tillegg viste det at den mest brukte inndataen ikke nødvendigvis var det beste. Et enkelt type bilde (T1) gjenga noe bedre resultater enn en etablert metode der alle bildetyper blir brukt.

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Chapter 1

Introduction

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1.1 Problem and Motivation

A well functioning knee joint is essential to mobility and a important part of daily activities as standing, walking and running. Knee injuries, for example injuries on ligaments, meniscus, cartilage or tendons is common in sport among professional and amateur athletes. These injuries may damage their career or life quality in a significant.

Joint diseases and load over a longer time span may also also affect the knee joint mobility, and may also affect the life quality.

A lot of this injuries is a non life-threatening and the consequences from the injuries are not reducing life-quality enough. This makes the queues long and you usually have to wait a long time to get the diagnosis you need to continue the treatment of the injury.

The Magnetic resonance imaging (MRI) is a widely used technique to image and diagnose injuries like knee-injuries and gives the doctors the ability to look inside tissues and organs of the body. The data produced by MRI is not segmented. They need some kind of segmentation tool to make the segmentation of the images. These tools use a lot of time and often end up with errors in the segmentation. This is limiting the quantitative uses of the MRI images and humans has to correct the errors.

Successful treatment of a knee with injuries depends on well trained doctors and a good knowledge about the injury and the rest of the anatomy in the knee. In this master thesis we will look into how Convolutional Neural Networks (CNN) can automatically segment MRI images form knees and further findings on how to optimize the CNN.

1.2 Scope

This thesis will study the use of deep learning and the method named Convolutional Neural Network within computer science. MRI images are a tool to map the inside tissues of the body, and are widely used in medicine to diagnose injuries in the knee joint.

The scope of this thesis is therefore:

- Computer Science Artificial Intelligence / Machine Learning Deep Learning,
 CNN
- Medicine MRI MRI segmentation

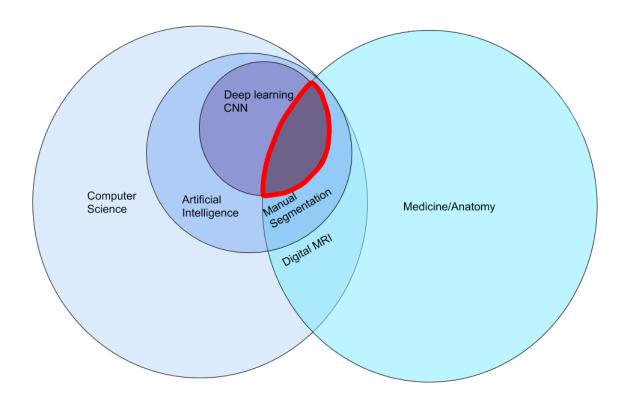


Figure 1.1: Scope

1.3 Research Question

Based on what has been mentioned in this document, the following research questions are formulated:

- Does the proposed Convolutional Neural Network (CNN) perform a good segmentation and make it more automatic then today's solution?
- How does the CNN perform in processing speed compared to older segmentation methods?
- What can be done to optimize the segmentation results?

1.4 Organization of This Report

The following document consist of six chapters that will cover this master thesis. Here is a overview of the chapters presented in this document.

1.4.1 Chapter One

The introduction is explaining the problem and gives a overview about the study and describe why this is a wanted tool.

1.4.2 Chapter Two

This chapter covers the literature review and the previous work related to this topic. It covers relevant information for understanding this thesis, and describes the scope more detailed.

1.4.3 Chapter Three

This chapter includes methods and descriptions of the CNN and processes used to solve the problem.

1.4.4 Chapter Four

The results are presented in this chapter with a description.

1.4.5 Chapter Five

Discussion of the results discovered in chapter four.

1.4.6 Chapter Six

Conclusions from this thesis.

Chapter 2

Theory and Literature Review

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2.1 Knee Anatomy

It is important to understand the parts of the knee joint, to know how to evaluate the results of the CNN. The structures of the knee can be divided into several categories such as shown in the following list and figure [2]:

- Bones
- Joints
- Ligaments
- Tendons
- Muscles
- Nerves
- Blood Vessels

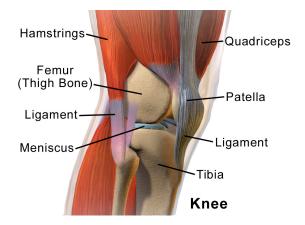


Figure 2.1: Knee anatomy: $https://upload.wikimedia.org/wikipedia/commons/b/bc/Blausen_0597_KneeAnatomy_Side.p$

2.1.1 Bones

The knee joint is where the two bones femur (top) and tibia (bottom) meet. Other bones close related to the knee joint are the patella (knee cap) and fibula. At the end of each bone, in the joint where the bone meets another bone, you have the articular cartilage. The articular cartilage has a slippery surface that allows the two surfaces at the end of each bone to slide against each other without damaging one another. The cartilage's main function is to absorb shock and give a slippery surface which helps the motion in the joint.

2.1.2 Ligaments

Ligaments are strong bands of tissue that connect the bones. You can find four ligaments in the knee joint. Two of them are located on each side of the knee joint. The inside ligament is named Medial Collateral Ligament (MCL) and the outside ligament is named Lateral Collateral Ligament (LCL).

The two other ligaments are inside the knee. The Anterior Cruciate Ligament (ACL) stretches from the front of the tibia to the back of the femur, and the Posterior Cruciate Ligament (PCL) stretches from the back of the tibia to the back of the femur. The LCL and MCL prevent movement in the side direction and ACL and PCL prevents movement too far in the front and back direction. The ligaments are the most important tissue to control the stability of the knee.

2.1.3 Meniscus

The meniscus is a fibrocartilage located between the femur and the tibia. The meniscus is important for two reasons:

- They work like a gasket to spread the force of the weight of the body over a larger area.
- They help the ligaments with the stabilization of the knee joint.

The meniscus distributes the weight from the femur over a larger area on the tibia and works like a pillow in between the two bones. This protects the bones from getting to much force and prevents the bones from taking damage. The meniscus is also thicker

on the edges and this helps the femur stay in place and not roll on the tibia.

The meniscus and ligaments are the most important part of the knee when it comes to stabilizing it. Without strong and tight ligaments to connect the two bones in the knee joint, you will end up with a loose knee joint and this can damage the knee.

2.1.4 Tendon

Tendons are similar to ligaments but connect muscles to bones. One of these is the quadriceps tendon that connects the large muscles in the tie named the quadriceps to the patella (knee cap). This tendon continues over the patella and connects to the patellar tendon which connects the patella to the tibia. The hamstring muscles in the back of the tie also have tendons that connect the hamstring with different places at the tibia.

The extensor mechanism in the knee is the motor that allows movements in the knee joint. It sits in the front of the knee joint and exist of the patella, patellar tendon, quadriceps tendon, and the quadriceps muscles. When the quadriceps muscles contract it straightens the knee joint like getting up from a squatting position.

2.1.5 Nerve

The most important nerves in the knee is the tibial nerve and the common peroneal nerve. Those nerves are positioned on the back of the knee. These two nerves travel to the lower leg and foot to give sensation and muscle control in the lower leg and foot. The sciatic nerve splits above the knee joint into the tibial nerve and the common peroneal nerve. Both the tibial nerve and the common peroneal nerve can be damaged by injuries in the knee joint.

2.1.6 Blood Vessels

The major blood vessels in the leg travels with the tibial nerve on the back of the knee. The popliteal artery and popliteal vein is the largest blood supplies to the leg. The artery carries blood out to the foot and the vein carries the blood back to the heart. If big damages happen to the popliteal artery and there are no possibilities for repair, there is most likely not possible to save the leg.

2.1.7 Injuries

The knee has an unstable design and has to support the body's full weight while standing and much more than that when walking, running or jumping. This is one of the reasons why knee problems are a common complaint among people of all ages.

All ligaments in the knee can be injured. It can be stretched, partially torn or completely torn. Among these injuries, the most common injury is completely torn. Symptoms of this are pain, popping sound when the accident happens, instability of the knee joint and joint swelling. In half of the cases where one ligament is torn, the surrounding ligaments, cartilage or meniscus are also damaged.

This includes all of the following ligaments:

- Meniscus tear
- ACL strain or tear
- PCL strain or tear
- MCL strain or tear
- LCL strain or tear

In case of injuries, an experienced hand can be accurate and tell if the ligaments are torn or not. But to confirm the injury it is often used MRI to provide images of the soft tissues like ligaments and cartilage in the knee.

2.2 Magnetic Resonance Imaging

MRI uses a magnetic field and radio frequencies to take pictures of the body instead of ionizing radiation as used in x-ray and CT scans. The magnetic field in an MRI machine is measured in Tesla which represents the magnetic flux density of the magnet. MRI machines are usually from 1.5T to 3.0T. This produces a very strong magnetic field, compared with the magnetic field of the earth that is 0.00003T. The strength of this magnet is strong enough to pick up a car.

Our body is composed of 70% water or H2O and the MRI relies on the magnetic properties of the hydrogen atom to produce MRI images. The hydrogen atom has a single proton in the center of the atom. The atom can be charged with a spinning momentum that produces a magnetic field also named a magnetic moment. Normally the protons are oriented randomly in all directions when there is no magnetic field presence. When the hydrogen atom is in a strong constant magnetic field their magnetic moment line up parallel or anti-parallel to the field. This is also referred to as a longitudinal magnetization. This can be done by a superconducting magnet such as the magnetic field produced by the primary magnet in an MRI machine.

This property of hydrogen in a magnetic field is what makes the MRI possible and is, therefore, an important part of MRI imaging.

2.2.1 MRI Components

The MRI has several components to be able to map the body in a 3-dimensional image. The components are as follows:

- Primary magnet
- Gradient Magnets
- Radio frequency (RF) coils
- Computer system

Service connections MRI Scanner cross section (including emergency cryostat vent pipe) Thermal insulation . Scanner bore Cryostat filled with liquid helium Superconducting primary electromagnet coil Magnetic gradient coils RF transmit coil Longitudinal section Scanner table Body part specific receive coil

This chapter will describe the different parts and their properties.

Figure 2.2: MRI components: https://en.wikipedia.org/wiki/File:Mri_scanner_schematic_labelled.svg

Primary Magnet

This is the magnet which makes the strong constant magnetic field in the MRI machine. As mentioned earlier this magnetic field has a strength between 1.5T and 3.0T and covers the entire machine.

Gradient Magnets

The gradient coils generate a second magnetic field overt the primary field. There are three gradient coils in different directions. The directions are in z, x, y-axis and represent the directions the MRI can take pictures. The gradient coils produce a gradient magnetic field that helps the localization of pixels in the image.

Radio Frequency(RF) Coils

The RF coils are responsible for transmitting the radio frequency or RF pulse and helps you receive images in MRI. The RF coils come in several designs to suit the part of the body that is going to be imaged. The RF pulse is responsible for flipping the protons into a high energy state and decreasing the longitudinal magnetization. It also synchronizes the protons. This turns the magnetization vector of the proton towards the transverse plane also named a transverse magnetization. The proton does not stay in this state for a long time and will end up in its normal state within a small amount of time. This change induces an electrical signal in the RF receptions and the transverse plane picks up this signal and stores it in the computer system.

Computer System

The computer system receives the signal from the RF and converts it from analog to a digital signal. This data is then applied to a Fourier Transformation and this produces the result image.

2.3 MRI Images File Format

The results produced from an MRI is an output file named DICOM. This section is about the output file.

The image format that an MRI scan is returning is a Digital Imaging and Communications in Medicine (DICOM) file. This is the standard for handling, saving, printing and transferring medical images and other information regarding this image/patient. The DICOM-file consists of a header and a data set containing the data that you want to save of the patient. The header is the part of the filer where the information about the patient is stored. Example for information that can be stored in the header is:

- Patients name, age, sex, weight and height.
- Acquisitions made from the doctor about what he thought of the image
- Image dimensions
- Matrix size
- Colour space
- Or other...

The reason why the header and the image data is in the same file is because it should be hard to separate the two of them. You do not want to lose the patient info connected to the given image. When converting the file from DICOM to another type of file you will lose the patient data. This is in our case necessary because the patient data is classified and unauthorized people are not allowed to work with this type of data. The DICOM format stores any kind of data and images are stored as pixel data from an MRI session. Since the DICOM file only stores one image each file, you need to store as many files as there are layers in the z-direction of the image. All the images are stored with the file type "*.dmc". If there is an MRI scan producing a 255*255*255 pixel image, you end up with 255 "DCM" files containing one image each file.

2.4 Machine Learning

Machine learning is all about understanding and extracting knowledge from data. It is a researching field within Artificial Intelligence and has a lot of branches branching out of it. The use of applications helped by a machine learning algorithm is to find in all our lives. Everything from picking recommended music, videos, movies, and other entertainment, to unlocking our phones and in general makes tasks in our daily lives easier. The idea of machine learning is not a new field of research. AI was first discussed by Alan Turing in the 1950s and this raised the question: Could a computer go beyond "what we know how to order it to perform" and learn on its own how to perform a specified task? The use of machine learning has a big influence on how data-driven research is done today. It helps science understand problems that not necessarily are recognized by humans such as finding particles, analyzing DNA and recognizing cancer.

2.5 Neural Networks

Neural networks are inspired by our human brain. The human brain contains roughly 86 billion neurons and the connections between these neurons are what make our mind so powerful. Our mind controls everything from your body, thoughts, memories and more. The concept of using a neural network-model in computing is not a new field of research [XXX]. It was first presented for over 60 years ago, but the technology at that time had no possibilities to apply such a model. In this section, we are going to look closer into how neural networks function and particularly how the convolution neural networks function (CNN). CNN is the computer equal to human eyes and is copying how shapes, color, and shades are processed in our brains. There are many kinds of neural networks and some of the easiest to imagine is a feed-forward neural network. The neural network is containing the following parts:

2.5.1 Input Layer

The input-layer works as an entrance to the hidden layers. This entrance is tailored for the data to fit into the neural network. The data often needs to be processed to fit

into an input layer and when the data is in the same size and dimensions as the input layer, the data is ready to be processed in the neural network.

2.5.2 Hidden Layers

The hidden layers are what makes up the neural network, and the "magic" it does. In the hidden layer, we are talking about numbers of hidden layers. This is the number of layers with a decided amount of neurons in the depth of the network. In between the layers are connections/weights which separates the neurons from the previous layer to the next layer.

2.5.3 Neurons

Neurons are connected to the previous layer and receives values from the previous neurons which are weight and then pushed through an activation function. The value of the current neuron can be updated by changing the value of the weights coming into the neuron.

2.5.4 Activation Function

As mentioned in the previous section, there is a function used to calculate the new value of a neuron. This is used to normalize data and keep the neurons under control. A neuron can often make conclusions by looking at noise (data that are misleading or confusing) and keep evolving around this misunderstanding. There are therefore a lot of different types of activation functions depending on what type of problem you want to solve. E.g. the activation function named ReLu is converting all negative values to 0 and all positive values are still the same.

2.5.5 Output layer

The output layer is similar to the input layer. It is the exit of the neural network and this is where you get your results. Therefore the output layer has to be in the same dimensions and size as the expected results. This means that the labeled known training results should have the same size and dimensions as the output layer. So if you are classifying an image to be either a "Dog" or a "Cat" the output should be an approximation of what the network thinks it is e.g. [0.95, 0.02]. This is showing that this neural network thinks it is a "Dog".

2.5.6 Data Set

Picking better data

To solve the problem, it is useful to pick the data that suits the problem best. This includes removing data that are not useful or generating/add more data that we want and supports our conclusion.

Pre-Processing and Scaling

In neural networks and convolutional neural networks, the algorithms are sensitive to the scale of the data. Therefore, it is common practice to scale and pre-process the data set before giving it to the network. The data should be adjusted in such a way that it suits and optimizes the performance of the network. In a lot of cases, the data is too detailed, the contents can be irrelevant for the problem that you are trying to solve, or just not fit for the network to function optimally. In this case scale the data to fit the network by e.g. reducing dimensions, changing the data type, or fitting it into another shape that suits better. A very important thing to remember when you are changing the training data is that the test data should be treated in the same way. If it is necessary to keep the original data types and scale for the results, this can be achieved by undoing the processes done before the neural network.

2.5.7 Multi-Class Classification

In a lot of cases, it is enough to calculate single/binary values like is it a dog in this image or not. But in cases where there are more than two classes to classify a problem appear. The problem is an instance of multi-class classification. When this is the case and every single point should be classified into only one of the categories, meaning that one of the categories is 1 and the rest is 0, you have an instance of single-label, multi-class classification.

2.5.8 One-Hot Encoding

When classifying multiple classes or solving categorical variables (single-label, multiclass classification) such as the one mentioned in the above chapter. The most common way to represent the data is using the one-hot encoding also known as categorical encoding. The reason behind this is to simplify the variables into true-false, or 0 -1 values to represent what class it is representing. E.g it is possible to classify an image and tell if there is a cat or a dog in it.

This can also be done per pixel in the entire image.

2.6 Convolutional Neural Network

The human vision is an advanced sensor. Within fractions of a second, we can identify all objects in our line of sight without even thinking about it. We can name the object, tell how far away it is, calculate its movement and predict where the object is going to be in the near future. Our eyes take raw pixels of colors and transform them into more primitive shapes like shadows, lines, curves, and other shapes. And then it gets process into one object and classified. The human vision is the motivation for the convolutional neural network and the difference between how an ordinary neural network and the CNN processes data.

Convolutional Neural Network (CNN) is a deep learning, feed-forward artificial neural network. CNN is good at recognizing objects in images and that is why CNN is a good solution for recognizing different tissues in an MRI image. The input to a CNN can be a 2D image, 3D image, sound or any data where the columns and rows are closely related to each other. This is because of the unique properties of the CNN algorithm.

CNN is a multiple layer neural network. This means that it uses several layers in depth to produce a result. The following sections will describe the layers in detail.

2.6.1 Convolution Layer

The convolutional layers perform a 2D convolution that compares a squared filter over the entire images. The filter could be an edge, line, dark spot or light spot, that scores the parts of the image to how similar that part of the image is to the filter. After applying the filters the convolutional layer gives the result to the next layer.

The first convolutional layers look at simple filters, but the filters get more complex in the following layers and could represent entire objects instead of lines or spots. In the following figure you can see how the different layers have different complexity in the filters and can recognize objects like faces and cars in the last layer.

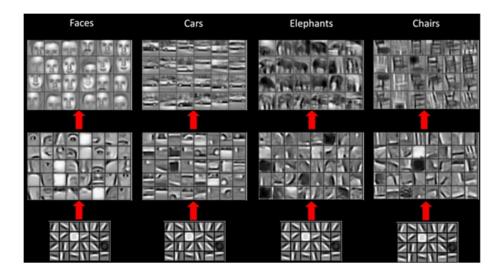


Figure 2.3: CNN Convolution Layers

2.6.2 Pooling Layer

The convolutional networks often include local or global pooling layers. This down-sampled the output and uses, for example, max pooling or average pooling to define the down-sampled value. The pooling layer also makes the filter less sensitive to the position. This is good because you do not want to focus on an exact position but what is appearing in the image.

2.6.3 Fully Connected Layer

The final layer in a CNN is named Fully Connected Layer. This is where every value that has gone through all the filters earlier gets a vote to find what the answer is going to be. We take all the values and list them up in a single fully-connected table where it weights against all the objects. The one object with the strongest average is what the CNN will return as its result.

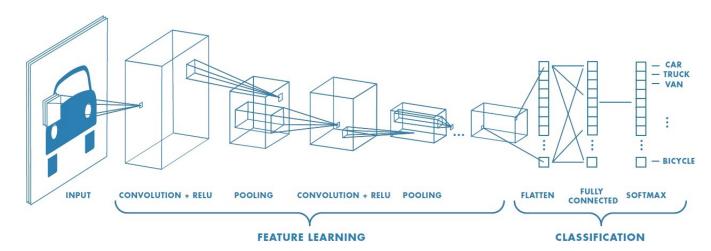


Figure 2.4: Full CNN network

Chapter 3

Methodology

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3.1 Software

The code for this was built using Python programming language, the Keras library with TensorFlow as backend.

3.1.1 Keras and TensorFlow

Keras is a widely used high-level neural network API. The Keras library is a user-friendly, modular and extensible library that allows fast prototyping. It is an excellent library for developing CNNs and runs seamlessly on both CPU and GPU. It is an open-source library that is written in python which makes it easy to understand and debug if any problems should occur.

Keras is using TensorFlow as a backend. TensorFlow is an open-source platform used for machine learning.

The U-Net model is programmed such that it is running on both of the graphic-cards on the computer.

3.1.2 Computer Specifications

The computer used to perform these tasks is a computer owned by NTNU. This computer has good hardware and is a good environment to perform a task such as this. The hardware specifications is as follows:

- Intel(R) Core(TM) i9-9960X CPU 3.10GHz (Cores)
- 64GB Ram
- SSD 1024 GB
- 2 x NVIDIA GeForce RTX 2080 with 8GB of VRAM

3.2 Data-set

The data-set is from 20 different knees, but one is not used because it occurred problems when reading the file. There are used several different imaging techniques used to image the knee. This is done by the MRI scanner and uses the different properties of the tissue to get different values from the individual tissues. In this data-set, we are using three different imaging techniques. This makes it more likely to distinguish the tissues from one another. The different imaging types that are used are T1, PD, and FS. The images are anonymous and delivered by MR-Klinikken Møre for research on this topic. The images are from the right knees from 20 individual subjects with an estimated healthy knee. The subjects are all adults in the age between 27 and 64. And it is a 50% distribution between males and females. The average age is 46.

All images are inside a preferable quality. This means that the images are in between the good conditions where the image is sharp and with good contrast and some images that did not qualify within these parameters were thrown away.

The labels or also known as the ground through is segmented by qualified persons in MR-Knlinkken Møre with help of tools that are widely used to segment such images.

3.2.1 Raw Data

There are 20 test images delivered with the raw data images from T1, PD, and FS as well as the ground truth/label saved as Nifty files.

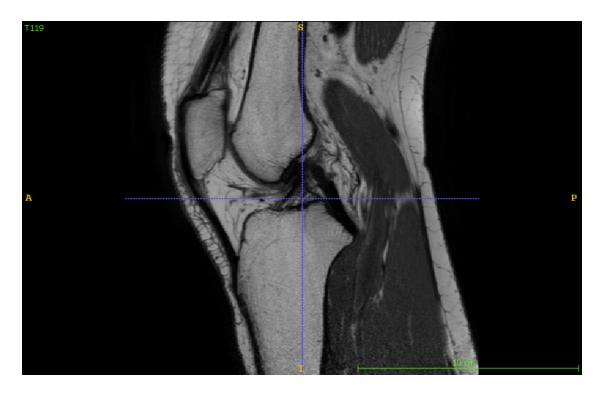


Figure 3.1: T1 raw image

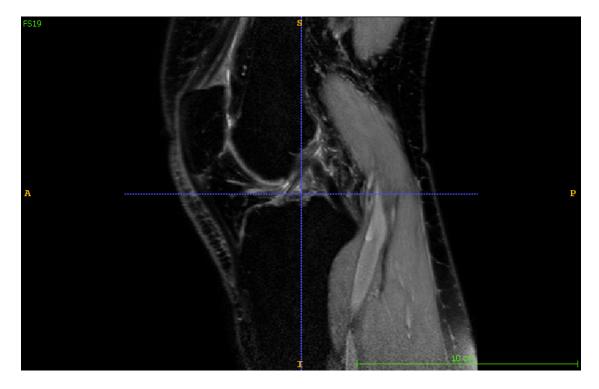


Figure 3.2: FS raw image

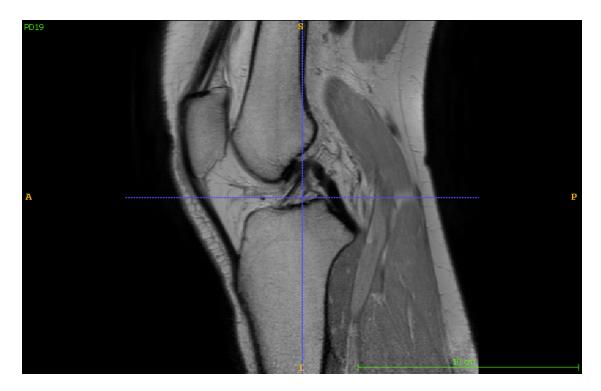


Figure 3.3: PD raw image

3.2.2 Mask/Label Annotation

The ground truth also named mask or label is a 3-dimensional image in the same shape/size as the raw image data. The difference is that it only contains values of the pixels that are classified as some of the tissues that are in the knee. This means that if a pixel is outside the knee joint or an unspecified tissue in the knee, it has the value 0. Depending on how many tissues are labeled in the image, the range of the pixels is integers reaching from 0 to the number of tissues classified in the image.

Since the labeled image contains an integer with the range from 0 to the number of tissues that are classified, the labeled image needs to be converted into one-hot encoding to fit the input values of the model [as discussed in the section about one hot encoding]. This is done while loading the labels and all the ground truth (labels) are stored as this while the program is processing the data. The labels used in this data-set is the following:

- Empty/not classified Value 0 Black
- Bone Value 1 Dark gray
- PCL Value 2 Light gray
- ACL Value 3 White

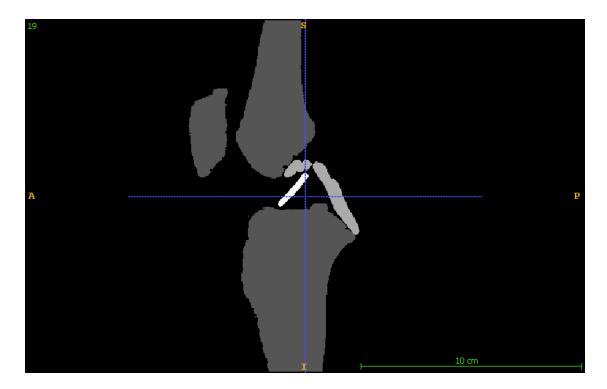


Figure 3.4: Label Image

3.3 Program

The following figure shows the flow of the python script and how it manipulates the data to get the predicted result. You can find the source code as an appendix XX.



Figure 3.5: Program Flow

The data we are using are 3-dimensional images of human knee joints. The original size

of the images is 400x400x275 where each pixel represents a space of 0.XX cm in real life. The original image is scaled down to make the images fit into the convolutional neural network. This is done by resizing the image to the size 256x256x256. The resizing of the image also applies to the labeled image, to make sure that the input and output pixels are the same amount. The input format is then given a new dimension and then the dimensions look like one of the following: 256x256x256x1, 256x256x256x2, 256x256x256x3. This depends on how many of the raw data image types we want the CNN to process.

Since the labeled data set is labeled with 3 labels. That is not included no label. This means that the output data has 4 different values as a possible value. In the labels, there are classified bones, and the ACL and PCL (see. Medicinal section). The same one-hot encoding is applied on the labeled data-set to fit it into the CNN.

The following table show how the size and dimensions are changed throughout the programming script.

Table 3.1: Dimension Size

Image Type	Raw image	Input	Output	Saved
	dimensions	dimensions	dimensions	classified
T1	1 x 400x400x275	256x256x256x1	256x256x256x4	400x400x275x4
PD	1 x 400x400x275	256x256x256x1	256x256x256x4	400x400x275x4
FS	1 x 400x400x275	256x256x256x1	256x256x256x4	400x400x275x4
PD, T1	2 x 400x400x275	256x256x256x2	256x256x256x4	400x400x275x4
PD, FS	2 x 400x400x275	256x256x256x2	256x256x256x4	400x400x275x4
FS, T1	$2 \times 400 \times 400 \times 275$	256x256x256x2	256x256x256x4	400x400x275x4
PD, T1, FS	3 x 400x400x275	256x256x256x3	256x256x256x4	400x400x275x4

3.3.1 UNet

The Convolutional neural network is a U-Net [8] inspired network and it is built using the Keras library. The figure [xxx] shows how the network structure is. It shows 5 levels of layers where the first layer is the input layer. After the input layer, there is a convolutional layer that is discussed in section 2.6.1. All the convolutional layers are followed up by a batch normalization layers. After two convolutional layers with a corresponding batch normalization layer, there is a max-pooling layer. The next level starts with a dropout layer where the first dropout layer is halved to prevent the model from forgetting any good finds in the first layer. Then it repeats the same pattern as described above. This repeats in all 5 levels and then the 5 decoding levels starts, where the goal is to reach the original image size with the labels instead of pixel values. The first level is started with an up transposed and followed by a merging with the concatenation and followed by a new dropout layer. Then it is new layers with two times of convolution and batch normalization. Next, there is a new transposed layer to double the size of the image and the layer is done. It is then repeated until it is scaled up to the same size as the input and then the output is generated. All max-pooling layers and transpose layers contain a size of 2x2, so it gets half the size each time the max-pooling is applied and double size when the transpose layer is applied.

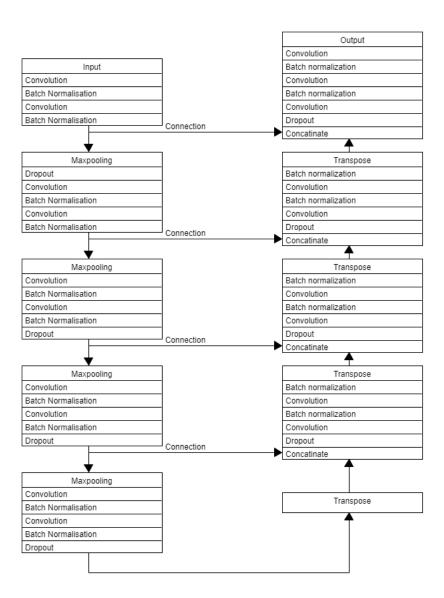


Figure 3.6: CNN structure. See detailed summary in Apendix B CNN Summary.

3.3.2 Activation Function

The convolutional layers are using ReLu as an activation function and the padding is the same to keep the same size of the image.

The last activation function is a sigmoid function. It could also have been a softmax function but this is solved later by applying an argmax function (A function that gives the class with highest probability a new value of 1 and the other classes a value of 0.)

to the output and then get the highest scored output. Sigmoid and softmax are pretty similar but softmax also applies the argmax. Because it was used sigmoid in this last layer the argmax function has to be applied after the output has been generated.

3.3.3 Loss Function

Categorical Cross-entropy

To calculate the loss with categorical cross-entropy you take the true value and multiplies this with the log value of the prediction. Say that we are looking at only one class and one pixel where the true values is 1. And for example, the prediction is right. This gives us the following loss: -1*log(1) = 0. Say that the prediction is only 50% probability on the right class, the loss will be as follows: -1*log(0.5) = 0.693 The loss is added together for all the classes in all the pixels of the image and this is how you get the total loss in the image.

Weighed Categorical Cross-entropy

For an unbalanced data-set such as this one, the original categorical cross-entropy usually updates the weights so that the most common classes are prioritized and the lower weighed classes often get ignored. This is because a class with lesser pixels in the ground truth does not add together to a big enough loss to make a difference.

To solve this there is added a weight to the calculated loss. If there is a class that has the true value of 1 and it is a class that needs to be weighed up with a value of 2 and the predicted value is 0.5. This gives us the following loss: -1*log(0.5) * 2 = 1.386

To predict the given problem and the unbalanced classes in our data-set, it is calculated the original categorical cross-entropy, and two versions of weighted categorical cross-entropy. The following values are the weights used on the loss function, where the first value represents value and class 0 and continuing left up to value and class 3:

$$WCC = [1.0, 1.05, 6.9, 11.8]$$

$$WCC2 = [1.0, 1.1, 12.0, 19.7]$$

3.4 Evaluation

Sometimes when evaluation the results, it is not enough to look at the accuracy. This especially true when you are looking at a multi-class classification problem. Like other problems such as binary classification, it is possible to evaluate the results in multiple ways and then calculate the average of all classes. This chapter will tell more about the evaluation methods used in this master theses

3.4.1 Confusion Matrix

Confusion matrices are often used for the evaluation of a model. When solving a binary classification (see chapter about binary classification) this could be illustrated pretty easily. In the next table you can see how binary classification results can be distributed four different categories. This is True Positive, True Negative, False Positive and False Negative, also known as TP, TN, FP, and FN. True positive and true negative is when you are predicting the correct value of that given point either if it is a positive value or a negative value. When you hit a false positive the model gives a false alarm a predicts this point to be something that it is not. And a false negative is when it is predicting noting but missing on a positive value.

Table 3.2: Confusion Matrix

True	Predicted	Predicted
Predictions	True	False
True	TP	FN
False	FP	TN

Metrics for Multi-Class Classification

In short steps, the multi-class classification is similar to the binary classification, but you take the average over all the classes given in the problem. When classes are imbalanced as it often is. The accuracy is not any good type of evaluating the network. An example of an imbalanced data-set where the model has classified 90 % to be class A, 6 % class B, and 4 class C. In this case, the accuracy does not mean anything at all. You could score 85 % accuracy, but this does not tell if you are correctly predicting anything from class B or C.

Multi-Class Confusion Matrix

The confusion matrix is, in this case, the only way to control that the model is correctly predicting the separate classes. Considering the same three classes (A, B, and C) as the model example above. You can see in the following table that the model is predicting the classes with a bad rate. Although the accuracy is tolerably high.

Table 3.3: Confusion Matrix with values.

True	Predicted	Predicted	Predicted
Values	\mathbf{A}	В	\mathbf{C}
A	180	0	0
В	2	6	2
С	0	3	7

This confusion matrix shows in detail how the predictions are made by the model. Known classes is representing the label data that is known to be the correct answer to this problem. The Predicted Class is what the result from the predictions the model has made. The diagonal elements in this matrices are the correctly predicted classes and the other values show how the model misses and tells you a lot about what issues this model is having with the problem it is trying to solve.

True Predicted Predicted Predicted \mathbf{C} Values \mathbf{A} В Α TP_a e_{ab} e_{ac} В TP_b e_{ba} e_{bc} \mathbf{C} TP_c e_{ca} e_{cb}

Table 3.4: Confusion Matrix Notation

The notations used for values taken from the confusion matrix when evaluation multi-class classification is as the table shown above.

3.4.2 Measure performance

There are several methods used to calculate the performance of a model. The reason behind having several methods is to understand and measure the performance of the model in more detail.

Accuracy

Accuracy is the simplest way to measure performance and often used to evaluate the model while training. It is total predictions that are true divided by total data point in the entire data set.

$$Accuracy = (TP)/(total_data_predicted)$$

Precision

Precision also named positive predictive value is measuring the accuracy within a given class. This means that if the model has predicted this class, what is the accuracy for tat this is a correct prediction.

Binary classification

$$Precision = tp/(tp + fp)$$

Multi-class classification

$$Precision(a) = TP_a/(TP_a + e_b a + e_c a) = 180/180 + 2 + 0 = 0.989$$

To calculate the total precision for the entire model, you calculate the average precision of each class.

Recall

Given that this is a value that should be positive, what is the accuracy for the model to find and predict the true value.

$$Recall = sensitivity = tp/(tp + fn)$$

$$Recall(a) = TP_a/(TP_a + e_ab + e_ac) = 180/(180 + 0 + 0) = 1.0$$

F1-score

Say that you have two different models or algorithms scoring differently on accuracy, precision, and recall. Where one is better in precision and the other is better on recall. f1-score helps you classify which one of these models/algorithms is the better one. The mathematical formula for f1-score is:

$$f1 - score = 2 * (precision * recall) / (precision + recall)$$

When scoring a multi-class classification there are two possible ways to calculate the f1-score, which is macro-f1 and weighted-f1.

Macro f1-score

Macro f1-score is the simplest way to calculate the f1-score for multi-class classification. This takes the average of the f1-scores for each class. And the mathematical formula is as follows [7]:

$$Macro - F1 = (F1_a + F1_b + F1_c)/3$$

The macro f1-score does not consider that the classes are differently weighted. this tells us a lot about how it scores on each class but does not consider the total image.

Weighted f1-score

The weighted f1-score takes to a count that the classes may not be equally weighted. This means that the F1-score gets weighted by how many percentage of this class is represented in the data-set.

The mathematical formula for this is as follows:

$$weighted - f1 = (F1_a * true_a + F1_b * true_b + F1_c * true_c) / (total_p redictions)$$

Chapter 4

Results

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4.1 Loss Function

As discussed earlier there are used 3 different loss functions used to calculate the loss on CNN. See section 3.3.3. Early findings in the project show that the original categorical cross-entropy was not able to score any of the lower weighted classes.

As showing in the upcoming tables, there is a significant difference in the performance between the functions used. Especially when training the model for lesser iterations, you will see that the categorical cross-entropy has ignored the lower weighted classes and does not hit any pixels in those classes.

4.1.1 Categorical Cross-entropy

The evaluations of the categorical cross-entropy with one T1 input and 10 iterations and all three inputs with 1000 iterations are shown in the tables below. The categorical cross-entropy has greater problems when scoring the lower weighted classes. But the scores in the first two classes seems to be a good representation of the ground truth. The macro F1-score shows that the average over the F1-scores is pretty bad. Although the model that used 1000 iterations of training scores way better, and are evening out the gap from the other two loss-functions. It still scores the worst on the smallest class value 3.

Table 4.1: CC (T1 10)

Class	True	Recall	Precision	F-Score
	Positive			
0	39847412	0.997962	0.9975	0.998
1	3969431	0.98535043	0.9799	0.983
2	0	0	Nan	Nan
3	0	0	Nan	Nan

Note: Macro F1-score: 0.495077424, Weighted F1-Score: 0.995356337.

Table 4.2: CC (PDFST1 1000)

Class	True	Recall	Precision	F-Score
	Positive			
0	39826899	0.997448257	0.998432467	0.997940119
1	3974149	0.986521602	0.979427857	0.982961931
2	31011	0.82651919	0.763216184	0.793607329
3	2361	0.449971412	0.191670726	0.26883006

Note: Macro F1-score: 0.76083486, Weighted F1-Score: 0.996307596.

4.1.2 Weighted Categorical Cross-entropy

The evaluations of the weighted categorical cross-entropy is shown in the tables below. The tables are showing one T1 input with 10 iterations of training and all three inputs with 1000 iterations of training. As the macro f1-score shows that the weight helps the training find the lower weighed classes early in the training.

Table 4.3: WCC (T1 10)

Class	True	Recall	Precision	F-Score
	Positive			
0	39789051	0.9965	0.999	0.9977
1	3987212	0.9898	0.972	0.981
2	34902	0.9302	0.649	0.7649
3	3904	0.744	0.313	0.4402

Note: Macro F1-score: 0.795939725, Weighted F1-Score: 0.995899592.

Table 4.4: WCC (PDFST1 1000)

Class	True	Recall	Precision	F-Score
	Positive			
0	39821311	0.997308308	0.998482868	0.997895242
1	3974473	0.98660203	0.978368421	0.982467975
2	32523	0.866817697	0.701349953	0.775354027
3	2467	0.470173432	0.260699567	0.335418083

Note: Macro F1-score: 0.772783832, Weighted F1-Score: 0.996214022.

4.1.3 Weighted Categorical Cross-entropy 2nd Version

The evaluations of the weighted categorical cross-entropy version two is shown in the tables below. The tables are showing one T1 input with 10 iterations of training and all three inputs with 1000 iterations of training.

Table 4.5: WCC2 (T1 10)

Class	True	Recall	Precision	F-Score
	Positive			
0	39820557	0.99729	0.9982	0.99773
1	3956394	0.98211	0.9834	0.98277
2	35418	0.94398	0.5659	0.70763
3	4858	0.92586	0.2354	0.37535

Note: Macro F1-score: 0.765870872, Weighted F1-Score: 0.996036625.

Table 4.6: WCC2 (PDFST1 1000)

Class	True	Recall	Precision	F-Score
	Positive			
0	39819616	0.997265857	0.998468482	0.997866807
1	3974554	0.986622137	0.977598792	0.982089738
2	31909	0.850453092	0.687560602	0.760380798
3	2311	0.440442157	0.31796918	0.36931682

Note: Macro F1-score: 0.777413541, Weighted F1-Score: 0.996144863.

4.2 Compared results of different input

In this section, we will show the results of the performance of the different imaging types. As discussed earlier in section 3.2 there are 3 different imaging techniques delivered in this data-set and we will look deeper into how these images performed. And maybe discover if one of them are containing more information or a better choice when you want to segment the classes.

4.2.1 One Input

In this section, we will look at how the different image techniques performers when training the model. Comparing the three different imaging types FS, PD, and T1 to see which one of these images performs the best. We will look into how it scores in the evaluation methods discussed in section 3.4.2. There seems to be a big gap between how the different imaging types scores in the early stages. As shown in the tables below there are some imaging types that are not able to detect some of the classes. Although FS and T1 seem to do pretty decent there are some big differences with the PD images.

PD

The figure shows how the PD image type has evolved the weights in the network with the weighed categorical cross-entropy after just 10 iterations of training. You can see that this input has problems detecting the lower weighted classes such as the ACL and PCL values 2 and 3. In the table, you can see the evaluation of the entire 3-dimensional image. The calculated average F-scores is bad because it does not score the third class at all. The average F1-score and weighted F1-score is as follows:

Average = 0.629306106



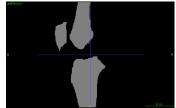




Figure 4.1: Raw PD, Predicted output, Ground truth

Table 4.7: PD 10 iterations

Class	True	Recall	Precision	F-Score
	Positive			
0	39777731	0.996216865	0.997099323	0.996657899
1	3935749	0.97698939	0.965525878	0.971223809
2	18446	0.49163113	0.609261461	0.544161897
3	0	0	Nan	Nan

FS

The following figure shows how the FS image type has evolved the weights in the network with the weighed categorical cross-entropy after just 10 iterations of training. It manages to find all the classes but still has some problems classification the lowest weighted classes.

Average = 0.714017027



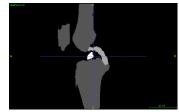




Figure 4.2: Raw FS, Predicted output, Ground truth

Table 4.8: FS 10 iterations

Class	True	Recall	Precision	F-Score
	Positive			
0	39669131	0.993497023	0.994866632	0.994181355
1	3825669	0.949663716	0.944144318	0.946895974
2	31276	0.83358209	0.611851243	0.705709805
3	2963	0.56470364	0.128440765	0.209280972

T1

The following figure shows how the T1 image type has evolved the weights in the network with the weighed categorical cross-entropy after just 10 iterations of training. The average Macro F1-score is the best score in all of the results with only 10 iterations.

Average = 0.795939725



Figure 4.3: Raw T1, Predicted output, Ground truths

Table 4.9: T1 10 iterations

Class	True	Recall	Precision	F-Score
	Positive			
0	39789051	0.99650037	0.998898098	0.997697793
1	3987212	0.989764291	0.97229485	0.9809518
2	34902	0.930223881	0.649436195	0.764874756
3	3904	0.744044216	0.312595084	0.440234551

4.2.2 Two inputs

By combining two of the images to the input layers. The evaluations start looking better. There are still some of the imaging types that perform better than the others. The different combinations of 2 image types are shown in the below sections where the tables show the scoring of evaluation methods discussed in section 3.4.2.

FS and PD

There are some problems when classifying lower classes in data containing two images where the PD image type is present. In this case, none of the lower classes were detected.

Average = 0.492114709 Weighted = 0.993381669

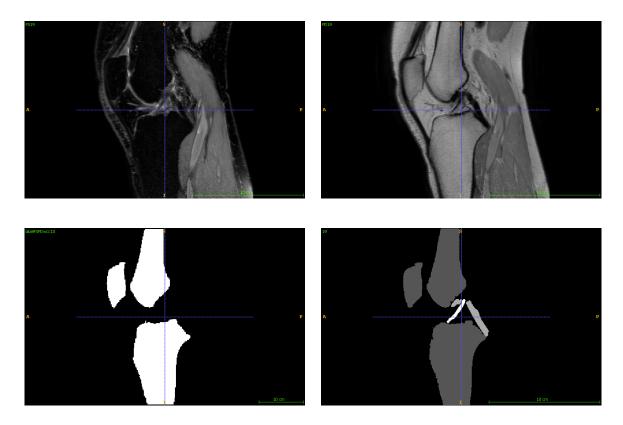


Figure 4.4: Raw FS, Raw PD, Predicted output, Ground truths

Table 4.10: FS, PD 10

Class	True	Recall	Precision	F-Score
	Positive			
0	39796975	0.996698823	0.996539267	0.996619038
1	3932679	0.976227309	0.967491549	0.971839799
2	0	0	Nan	Nan
3	0	0	Nan	Nan

FS and T1

This combination scores the best of the inputs containing two image types.

Average = 0.756220155

Table 4.11: FS, T1 10

Class	True	Recall	Precision	F-Score
	Positive			
0	39795367	0.996658551	0.99865021	0.997653386
1	3975564	0.986872854	0.977049596	0.981936658
2	36510	0.973081023	0.5853682	0.730996786
3	3893	0.74194778	0.199375192	0.314293788

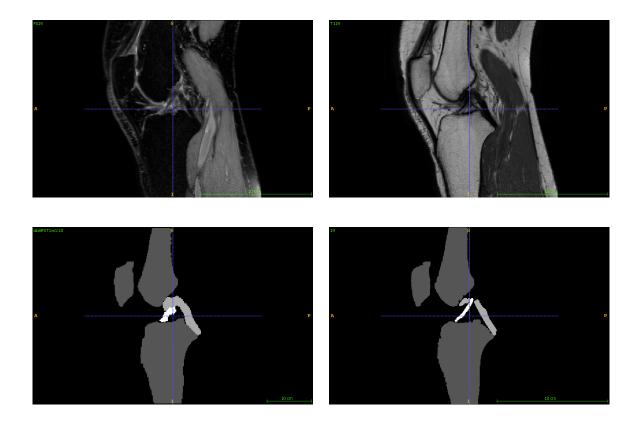


Figure 4.5: Raw FS, Raw T1, Predicted output, Ground truths

PD and T1

There are some still some problems when classifying lower classes in data containing two images where the PD image type is present. Combined with the T1 image type the PD scores a better then whit the FS.

Average = 0.68750736

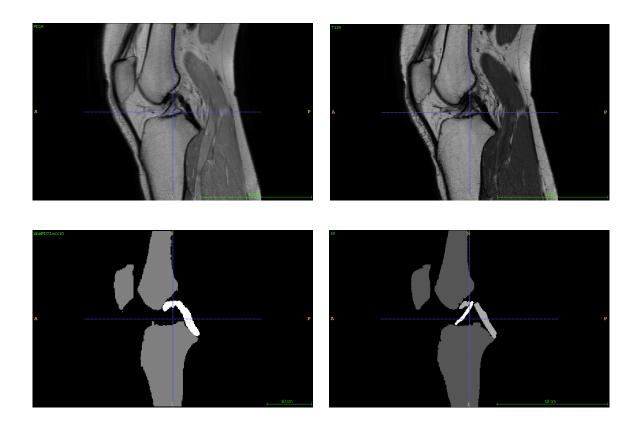


Figure 4.6: Raw PD, Raw FS, Predicted output, Ground truths

Table 4.12: PD, T1 10

Class	True	Recall	Precision	F-Score
	Positive			
0	39813380	0.997109679	0.99858805	0.997848317
1	3979720	0.987904517	0.975905694	0.981868449
2	34614	0.922547974	0.661203438	0.770312674
3	0	0	Nan	Nan

4.2.3 Three inputs

This is the most common way to combine the image-types to represent all three inputs and make the network look at all the image-types in one. Since this image contains the most of information there is easy to conclude that this is the best input for the network. Although there seems to be some imperfections also with this input.

Average = 0.798264327 Weighted = 0.995903632

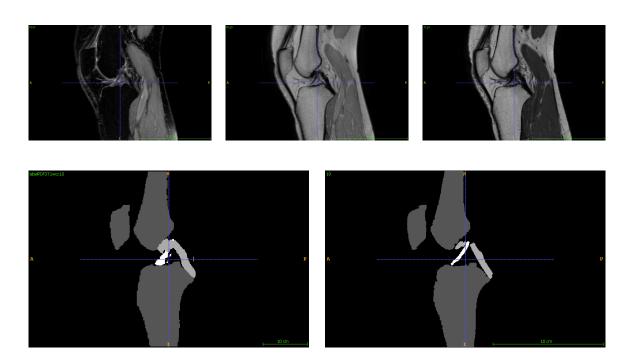


Figure 4.7: Raw FS, Raw PD, Raw T1, Predicted output, Ground truths

Table 4.13: PD, FS, T1, 10

Class	True	Recall	Precision	F-Score
	Positive			
0	39789588	0.996513818	0.998841493	0.997676298
1	3984817	0.989169769	0.973869288	0.981459901
2	34937	0.931156716	0.628216424	0.75026038
3	3901	0.74347246	0.230664617	0.352091701

4.3 Accuracy over Iterations

Show the results of the classification as a 3D image of the entire image. The model reduces its loss within a small number of iterations. It seems to get too detailed and starts introducing some errors. Although it scores better in f1-score the visualization looks more wrong.

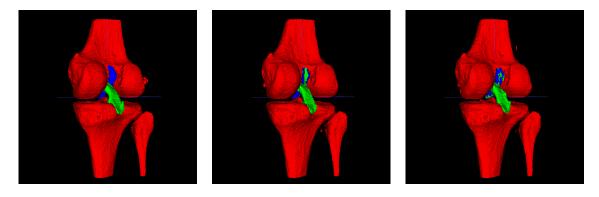


Figure 4.8: 10, 100, 1000

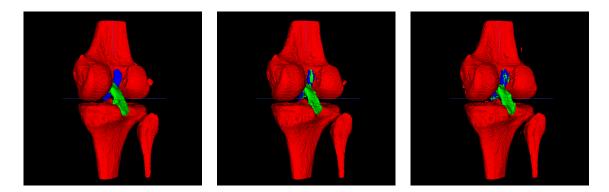


Figure 4.9: Caption for this figure with two images

Chapter 5

Discussion

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5.1 Loss-Function with Weights

The categorical cross-entropy (CC) scores great values on the knee, looks realistic with no obvious fails to recognize visually on its own and not comparing it to the labeled. Overall the categorical cross-entropy maybe the best-looking knee segmentation. But it does not score a good blue value (class 3). It does a god segmentation elsewhere and does not detect tissue remote from where it should be. When training the model for fewer iterations there is a significant issue in the CC's way to identify the lower classes and it regularly does not score any positives in class 2 and 3.

The first weighted categorical cross-entropy seems to be performing somewhere in between CC and the second version of weighted categorical cross-entropy. It does not achieve the best results at any of the classes but always makes a decent detection of the small classes and a satisfying segmentation on detecting the bone tissue values 1.

The second is the best model to predict value 3 and does that one properly. It does, however, find points outside what seems to be a reasonable area to detect the low weighted classes. This loss function had the best performance of them all when training for only 10 iterations. It detected all classes early, already after a small number of iterations with all inputs.

5.2 Best Input to Build a Model

In the earliest stages of training, the PD seems to contain less information than the other two imaging types. It has problems detecting the ACL and PCL, also when combined with other imaging types.

When training with only 10 or 100 iterations, this seems to be the best results combining all of the outputs. Although when training over 1000 iterations the differences seams to even out and there is no big differentiation anymore.

It seems that the 3 inputs introduces more noise than information to the model when

training the model for many iterations. This can be seen when looking at the best models from 1000 iterations. When studying this network the best evaluation seems to come from the segmentation from models created only whit T1 images.

5.3 Best Model

There are a couple ways to explain which model is the best. The first is an overall accuracy. This is model is made by combining the categorical cross-entropy and T1 image over 1000 iterations. This gives a clear and realistic outcome that does score as high as 99,638 % accuracy. Witch is a satisfying accuracy to have on a CNN. Out of a total of 44 000 000 pixels, there are 43 840 759 correct classified pixels. Only missing on 159 241 of the pixels. But there is a problem with this model as well. Although it scores high in accuracy it has a problem detecting the smaller classes. The smallest class contains 5247 pixels, and this model only scores 3050 correct classifications in this class.

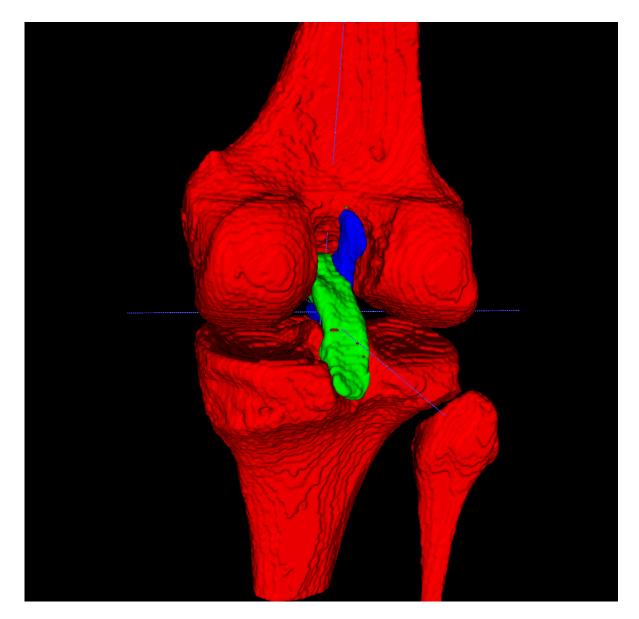


Figure 5.1: Best Result CC

The second way to score the best model is to look at the model that scores the highest accuracy on each of the classes combined. This model is created using one of the inputs. It uses T1 images and have the second weighted cross-entropy trained over 1000 iterations. It gives a realistic outcome with some missed boundaries in the ACL and PCL. It still has a good accuracy with 99,60 % and scoring 43 825 612 out of 44 000 000 leaving only 174 388 missed pixels. This model scores way better on the lower weighted classes and hits on 4436 of the total 5247 pixels in the class with value 3.

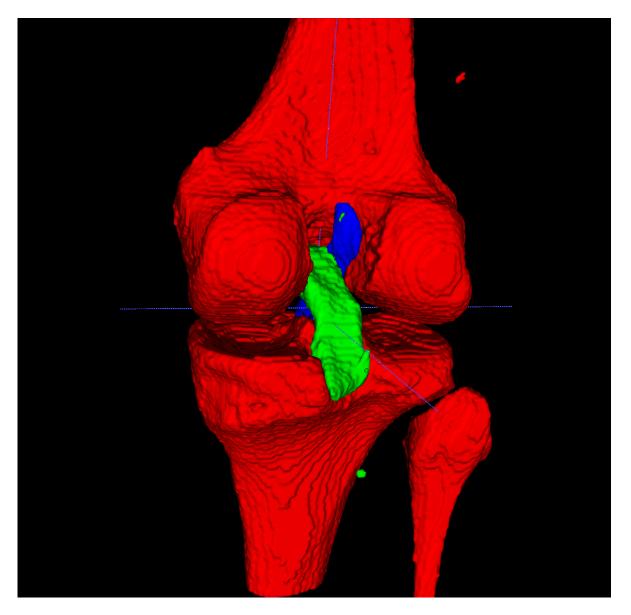


Figure 5.2: Best Result WCC2

5.4 Model error vs. Human error

We are scoring the accuracy of the predictions and on most of the cases, it is better then 99%. This brings up the question: How accurate are these predictions against the actual true values and does the old techniques also have some error when classifying the ground truth. The ACL and PCL are often drawn by hand by a human that is qualified to know where it starts, ends and where it should go in between these points. But the pixels that are marked are pixels that often are dark and other pixels surrounding the ligaments are also dark. This makes it hard for a human to see the difference and it can introduce errors to the labeled ground truth. By looking at the ground truth you can also detect some unrealistic pixels and shapes in the way the ligaments look. After viewing some of the results and applying the new predicted segmentation's on top of the original image, the output from this model seems to be as possible as the ground truth.

Chapter 6

Conclusion

6.1 Conclusion and Future Work

The segmentation of bodyparts are a valuable tool to both understand and examen the human body. You can add a segmented bodyparts in a 3-dimensional world, look at it from different angles and discover something you could not have seen from a 2-dimensional view.

The designed python script named MartiNet is a Convolutional Neural Network (CNN) using Keras with TensorFlow as backend. It is a CNN created to make detailed segmentation of the knee joint. It is a U-Net inspired network that can classify each pixel in an MRI scan as a tissue.

This CNN is a tool performs this task very well. It can make a precise prediction and outputs a segmentation that is over 99% accurate and looks as real as the ground truth. It is probably not a perfect segmentation and it will require few corrections. These corrections can be performed by a professional in a short amount of time. This is a significant decrease in workload taken into a count that the CNN segments the entire knee in only seconds compared with the old method that used hours to complete.

The biggest issue was when detecting small tissues that do not have a lot of pixels in the images. This was solved by adding a weighted loss function. This had a big impact on how it performed on the smaller classes an this could be considered further when segmenting more labels on the knee. The loss function named WCC2 sowed a promising result on classifying these lower weighted classes. For further work, there should be a task finding a ratio or an optimal description for the weights which could help a lot when classifying more labels.

When looking at the different input images and how they performed, there were no significant differences. The finding in this thesis shows results that can point in the direction of using only one input image (T1) when classifying just three labels as the bone, PCL and ACL.

This could also work on several other joints on the human body and a CNN like this

can almost without any modifications be trained to classify other bodyparts.

As further work, this CNN could detect more labels. The rate of learning in this CNN was fast and it had an accuracy of better than 95 % after only a few iterations. This shows that there is space for more complex problems and it is possible to add more labels to the segmentation.

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Appendix A

All Results

cc FS		10	66					
13	0	10		2	support	Recall	Precision	f-score
Λ	39703283	225504		0			0.994514392	0.99443336
1		3845321		0		0.95454202		0.94873987
2	30632	6888		0	37520	0.93434202	0.943007843	0.54673567
3	5241	6		0	5247	0	0	
3	3241	U	U	U	3247	0.48722359		0.4857933
						0.48722333		
						0.363741	0.368632034	0.36326330
PD	1	10	СС					
	0	1			support	Recall	Precision	f-score
0	39769067	147818	11902	0	39928787	0.99599988	0.997355704	0.9966773
1	93922	3934524	0	0	4028446	0.9766853	0.96356618	0.97008138
2	8831	931	27758		37520	0.73981876	0.657788099	0.69639609
3	2687	21	2539	0	5247	0	0	
						0.67812599		0.66578870
						0.9938943	0.993853591	0.99386741
T1		10	СС					
	0	1			support	Recall	Precision	f-score
0	39847412	81375		0			0.997456958	0.99770941
1		3969431		0	4028446	0.98535043	0.979865445	0.98260028
2	37336	184		0	37520	0	0	
3	5241	6	0	0	5247	0	0	
						0.49582811	0.494330601	0.49507742
						0.99583734	0.994876851	0.99535633
FS,	,PD	10						
_	0	1			support		Precision	f-score
	39780681	148106					0.996938789	
1		3948327		0			0.963672224	
2	36803	717		0			0	
3	5229	18	0	0	5247	0	0	
							0.490152753	
						0.99384109	0.992924047	0.99337610
FS,	T1	10		~		Dagall	Dua sisis :-	£
_	0	1			support		Precision	f-score
	39858451	70336			39928787		0.997300778	
1		3963113		0			0.982507745	
2	37304	216		0			0	
3	5241	6	0	0	5247	0	0	0.40500047
							0.494952131	
						0.99594464	0.994977039	0.995460

0 1 2 3 support Recall Precision f						CC	10	,T1	PD
	f-score	Precision	Recall	support	3	2	1	0	
0 39827330 99815 1642 0 39928787 0.99745905 0.998088118	0.997773486	0.998088118	0.99745905	39928787	0	1642	99815	39827330	0
1 47701 3980745 0 0 4028446 0.98815896 0.975031376	0.981551276	0.975031376	0.98815896	4028446	0	0	3980745	47701	1
2 23794 2119 11607 0 37520 0.30935501 0.847535597	0.453265645	0.847535597	0.30935501	37520	0	11607	2119	23794	2
3 4796 5 446 0 5247 0 0	O	0	0	5247	0	446	5	4796	3
0.57374325 0.705163773	0.608147602	0.705163773	0.57374325						
0.99590186 0.995729742	0.99570495	0.995729742	0.99590186						
		Description	D II		2	_	_	•	FS _.
0 1 2 3 support Recall Precision f	f-score					2	1	0	
0 1 2 3 support Recall Precision f 0 39818777 110010 0 0 39928787 0.99724484 0.997809963	0.997527324	0.997809963	0.99724484	39928787	0	2	1 110010	0 39818777	
0 1 2 3 support Recall Precision f 0 39818777 110010 0 0 39928787 0.99724484 0.997809963		0.997809963	0.99724484	39928787	0	2	1 110010	0 39818777	0
0 1 2 3 support Recall Precision f 0 39818777 110010 0 0 39928787 0.99724484 0.997809963	0.997527324	0.997809963 0.97307385	0.99724484 0.98886667	39928787 4028446	0 0	2 0 0	1 110010 3983596	0 39818777 44850	0
0 1 2 3 support Recall Precision f 0 39818777 110010 0 0 39928787 0.99724484 0.997809963 1 44850 3983596 0 0 4028446 0.98886667 0.97307385	0.997527324 0.980906699	0.997809963 0.97307385 0	0.99724484 0.98886667 0	39928787 4028446 37520	0 0 0	2 0 0 0	1 110010 3983596 214	0 39818777 44850 37306	0 1 2
0 1 2 3 support Recall Precision f 0 39818777 110010 0 0 39928787 0.99724484 0.997809963 0 1 44850 3983596 0 0 4028446 0.98886667 0.97307385 2 37306 214 0 0 37520 0 0 3 5240 7 0 0 5247 0 0	0.997527324 0.980906699	0.997809963 0.97307385 0 0	0.99724484 0.98886667 0	39928787 4028446 37520	0 0 0	2 0 0 0	1 110010 3983596 214	0 39818777 44850 37306	0 1 2

	wcc							
FS		10	wcc					
	0	1	2	3	support	Recall	Precision	f-score
0	39669131	222629	18391	18636	39928787	0.993497023	0.994866632	0.9941813
1	200707	3825669	600	1470	4028446	0.949663716	0.944144318	0.9468959
2	2552	3692	31276	0	37520	0.83358209	0.611851243	0.7057098
3	1428	6	850	2963	5247	0.56470364	0.128440765	0.2092809
						0.835361617	0.669825739	0.7140170
						0.989296341	0.989792791	0.989512
PD)	10	wcc					
	0	1	2	3	support	Recall	Precision	f-score
0	39777731	139826	11230	0	39928787	0.996216865	0.997099323	0.9966578
1	92616	3935749	81	0	4028446	0.97698939	0.965525878	0.971223
2	18382	692	18446	0	37520	0.49163113	0.609261461	0.5441618
3	4720	8	519	0	5247	0	0	
						0.616209346	0.642971666	0.6280109
						0.993907409	0.993758974	0.993824
T1		10	wcc					
	0	1	2	3	support	Recall	Precision	f-score
0	39789051	113445		8266			0.998898098	
1		3987212	339	4	4028446	0.989764291	0.97229485	0.9809
2	2140	163	34902	315	37520	0.930223881		
3	861	6	476	3904	5247	0.744044216	0.312595084	
						0.915133189		
						0.995797023		
FS,	,PD		wcc					
_	0	1	2		support	Recall	Precision	f-score
	39796975	131812	0	0			0.996539267	
1		3932679	0	0			0.967491549	
2	37225	295	0	0		0	0	
3	5213	34	0	0	5247	0	0	
							0.491007704	
						0.993855773	0.992911172	0.9933810
FS,	,T1	10	wcc					
	0	1	2		support	Recall	Precision	f-score
0	39795367	93188	24736	15496	39928787	0.996658551	0.99865021	0.997653
1	52676	3975564	191	15	4028446	0.986872854	0.977049596	0.981936
2	698	190	36510	122	37520	0.973081023		0.730996
3	414	6	934	3893	5247	0.74194778	0.199375192	0.314293
						0.924640052	0.690110799	0.756220
						0.995712136	0.996224822	0.995905

),T1	10	wcc					
0	1	2	3	support	Recall	Precision	f-score
39813380	98102	17305	0	39928787	0.997109679	0.99858805	0.997848317
48513	3979720	213	0	4028446	0.987904517	0.975905694	0.981868449
2757	149	34614	0	37520	0.922547974	0.661203438	0.770312674
5024	5	218	0	5247	0	0	0
					0.726890543	0.658924296	0.68750736
					0.996084409	0.996104575	0.996072251
,PD,T1	10	wcc					
0	1	2	3	support	Recall	Precision	f-score
39789588	106659	19705	12835	39928787	0.996513818	0.998841493	0.997676298
43528	3984817	92	9	4028446	0.989169769	0.973869288	0.981459901
2157	259	34937	167	37520	0.931156716	0.628216424	0.75026038
465	2	879	3901	5247	0.74347246	0.230664617	0.352091701
					0.915078191	0.707897955	0.798264327
					0.995755523	0.9961475	0.995903632
	39813380 48513 2757 5024 ,PD,T1 0 39789588 43528 2157	0 1 39813380 98102 48513 3979720 2757 149 5024 5 ,,PD,T1 10 0 1 39789588 106659 43528 3984817 2157 259	0 1 2 39813380 98102 17305 48513 3979720 213 2757 149 34614 5024 5 218 ,,PD,T1 10 wcc 0 1 2 39789588 106659 19705 43528 3984817 92 2157 259 34937	0 1 2 3 39813380 98102 17305 0 48513 3979720 213 0 2757 149 34614 0 5024 5 218 0 ,,PD,T1 10 wcc 0 1 2 3 39789588 106659 19705 12835 43528 3984817 92 9 2157 259 34937 167	0 1 2 3 support 39813380 98102 17305 0 39928787 48513 3979720 213 0 4028446 2757 149 34614 0 37520 5024 5 218 0 5247 ,,PD,T1 10 wcc 0 1 2 3 support 39789588 106659 19705 12835 39928787 43528 3984817 92 9 4028446 2157 259 34937 167 37520	0 1 2 3 support Recall 39813380 98102 17305 0 39928787 0.997109679 48513 3979720 213 0 4028446 0.987904517 2757 149 34614 0 37520 0.922547974 5024 5 218 0 5247 0 0.726890543 0.996084409 7,PD,T1 10 wcc 0 1 2 3 support Recall 39789588 106659 19705 12835 39928787 0.996513818 43528 3984817 92 9 4028446 0.989169769 2157 259 34937 167 37520 0.931156716 465 2 879 3901 5247 0.74347246 0.915078191	0 1 2 3 support Recall Precision 39813380 98102 17305 0 39928787 0.997109679 0.99858805 48513 3979720 213 0 4028446 0.987904517 0.975905694 2757 149 34614 0 37520 0.922547974 0.661203438 5024 5 218 0 5247 0 0 0.726890543 0.658924296 0.996084409 0.996104575 ,PD,T1 10 wcc 0 1 2 3 support Recall Precision 39789588 106659 19705 12835 39928787 0.996513818 0.998841493 43528 3984817 92 9 4028446 0.989169769 0.973869288 2157 259 34937 167 37520 0.931156716 0.628216424 465 2 879 3901 5247 0.74347246 0.230664617 0.915078191 0.707887955

FS		10	wcc2					
гэ	0			2	cupport	Recall	Drasision	facer
^	0	1	2		support			f-score
	39678430	212172				0.993729912		
1		3834083	1747	2111		0.951752363	0.94689197	
2	3755		30845	54		0.822094883		
3	2621	3	397	2226	5247			0.151903917
							0.658338734	
						0.989672364	0.990206835	0.989910978
PD			wcc2					
	0	1	2	3	support	Recall	Precision	f-score
0	39821088	89203	0	18496	39928787	0.997302723	0.995490417	0.996395746
1	142531	3885744	0	171	4028446	0.964576415	0.977504839	0.970997595
2	37030	148	0	342	37520	0	0	0
3	829	71	0	4347	5247	0.828473413	0.186119198	0.303954131
						0.697588138	0.539778613	0.567836868
						0.993435886	0.992898337	0.993138175
T1		10	wcc2					
	0	1	2	3	support	Recall	Precision	f-score
0	39820557	66461	26086	15683	39928787	0.997289424	0.998165827	0.997727433
1	71267	3956394	781	4	4028446	0.982114195	0.983431476	0.982772394
2	1820	189	35418	93	37520	0.943976546	0.565936436	0.70763114
3	85	6	298	4858	5247	0.925862398	0.235391026	0.375352521
						0.962310641	0.695731191	0.765870872
						0.995846068		0.996036625
FS,	PD	10	wcc2					
	0	1	2	3	support	Recall	Precision	f-score
0	39689988	200406	24678			0.994019378		
1	58982	3969054	263	147	4028446	0.985256846	0.951771568	0.968224778
2	1740		35083	0		0.935047974		
3	842		1119			0.622832095		
								0.741927281
							0.993719053	
FS,	T1	10	wcc2					
	0	1	2	3	support	Recall	Precision	f-score
0	39769603	111674	30139	17371	39928787	0.996013302	0.998911786	0.997460438
1	42075	3986144	115	112	4028446	0.989499177	0.972686932	0.98102103
2	1018	251	36251	0	37520	0.966178038	0.53496746	0.688639192
3	232		1258			0.714884696		
								0.73760453
							0.996017081	
								

PD),T1	10	wcc2					
	0	1	2	3	support	Recall	Precision	f-score
0	39741163	116756	38606	32262	39928787	0.995301034	0.998824192	0.997059501
1	46515	3980308	1330	293	4028446	0.988050479	0.971487819	0.979699152
2	185	56	36989	290	37520	0.985847548	0.475602073	0.641652138
3	83	6	848	4310	5247	0.821421765	0.116000538	0.203292298
						0.947655207	0.640478655	0.705425772
						0.994608409	0.995769951	0.995072341
FS,	,PD,T1	10	wcc2					
FS,	,PD,T1 0	10 1	wcc2 2	3	support	Recall	Precision	f-score
FS,		_	2		support 39928787	Recall 0.996170107	Precision 0.998921414	f-score 0.997543863
	0	1	2				0.998921414	
0	0 39775864	1 110154	2 22720	20049	39928787	0.996170107	0.998921414	0.997543863
0	0 39775864 41659	1 110154 3986218	2 22720 485	20049 84	39928787 4028446	0.996170107 0.989517546	0.998921414 0.973081108	0.997543863 0.981230501
0 1 2	0 39775864 41659 1207	1 110154 3986218 113	2 22720 485 36055	20049 84 145	39928787 4028446 37520	0.996170107 0.989517546 0.960954158	0.998921414 0.973081108 0.603774533	0.997543863 0.981230501 0.741597762

T1		100	СС					
	0	1	2	3	support	Recall	Precision	f-score
0	39834825	80632	6807	6523	39928787	0.99764676	0.998202401	0.997924503
1	63581	3964825	40	0	4028446	0.984207061	0.979798457	0.981997811
2	6110	923	30431	56	37520	0.811060768	0.802568769	0.806792423
3	2045	192	639	2371	5247	0.451877263	0.264916201	0.334014228
						0.811197963	0.761371457	0.780182241
						0.996192091	0.99626315	0.996224171
PC),T1	100	СС					
	0	1	2	3	support	Recall	Precision	f-score
0	39828632	88115	5397	6643	39928787	0.997491659	0.998433636	0.997962425
1	52615	3975822	9	0	4028446	0.986936898	0.978245128	0.982571792
2	7940	301	28950	329	37520	0.771588486	0.836632662	0.802795225
3	1929	1	247	3070	5247	0.585096245	0.305715993	0.401595919
						0.835278322	0.779756855	0.79623134
						0.9962835	0.996364687	0.996315786
FS	, T1	100	СС					
	0	1	2	3	support	Recall	Precision	f-score
0	39827545	86802	8257	6183	39928787	0.997464436	0.998333985	0.997899021
1	56912	3971504	29	1	4028446	0.985865021	0.978490719	0.982164028
2	6576	499	30223	222	37520	0.805517058	0.782776483	0.793983975
3	2976	1	101	2169	5247	0.413379074	0.252944606	0.31384749
						0.800556397	0.753136448	0.771973628
						0.996169114	0.996244524	0.996202938
PD	FS,T1	100	СС					
	0	1	2		support	Recall		f-score
0	39841868	80724	128		39928787			0.997678316
1	58267		4	0	4028446		0.980019348	0.982769493
2	37009	216	196	99	37520	0.005223881	0.428884026	0.010322037
3	3236	4	129	1878	5247	0.357918811	0.233465937	0.282597246
						0.586625239	0.659975709	0.568341773
						0.995775386	0.995353985	0.99538611
FS		100	66					
гэ	0	100	2	2	support	Recall	Precision	f-score
Λ	39686334	231360			39928787		0.995086117	0.994506654
1		3847679	29	71	4028446		0.942035039	0.948536025
2	11193		20883	75	37520		0.798218791	0.655852517
3	4117	25	304	801	5247		0.738218731	0.129811198
Э	411/	25	304	901	3247	0.152658662		0.682176599
						0.004374201	0.989955921	0.082170399
						0.303302203	0.303333321	0.50550551

T1	_	100		_				_
_	0	1	2		support	Recall	Precision	f-score
0	39825430	83229	13269	6859		0.997411467		0.997896372
1		3970016	345	1	4028446	0.985495648	0.979425003	0.982450948
2	5408		31866	82	37520	0.849307036	0.684819049	0.758244896
3	1060	6	1052	3129	5247	0.596340766	0.310694072	0.408538974
						0.857138729	0.743329968	0.786782797
						0.996146386	0.996296763	0.996207619
PD,		100		_				
_	0	1	2		support	Recall	Precision	f-score
0	39814900	94311		7849	39928787	0.997147747	0.998501805	0.997824316
1		3974876	166	1	4028446	0.986702068	0.976777526	0.981714715
2	4778		32469	90	37520	0.865378465	0.724334092	0.788599325
3	1559	7	464	3217	5247	0.613112255	0.288339159	0.392221409
						0.865585134	0.746988146	0.790089941
						0.996033227	0.996194348	0.996098762
FS,		100		•		5 II		
•	0	1	2		support	Recall	Precision	f-score
0	39810128	99234	13549	5876	39928787	0.997028234		0.997818041
1		3978670	60	0	4028446	0.987643871	0.975570943	0.981570285
2	4337	389	32762	32	37520	0.873187633	0.690584094	0.771224444
3	1396	6	1070	2775	5247	0.528873642	0.319590003	0.398420675
						0.846683345	0.746088535	0.787258361
						0.996007614	0.996156194	0.996065768
PD.	FS,T1	100	WCC					
,	0	1	2	3	support	Recall	Precision	f-score
0	39822307	84467	15296	6717	39928787	0.997333252	0.998486385	0.997909486
1	54357	3973751	337	1	4028446	0.986422804	0.979139378	0.982767597
2	4957	187	32355	21	37520	0.862340085	0.659767537	0.747573937
3	1053	7	1052	3135	5247	0.597484277		0.414655115
						0.860895105		0.785726534
						0.996171545		0.99624014
FS		100	wcc					
	0	1	2	3	support	Recall	Precision	f-score
0	39670670	232467	13345	12305	39928787	0.993535566	0.995097491	0.994315915
1	186796	3841219	256	175	4028446	0.953523766	0.941836513	0.947644106
2	5748	4735	27017	20	37520	0.720069296	0.662165143	0.689904368
3	2900	14	183	2150	5247	0.409757957	0.146757679	0.216112982
						0.769221646	0.686464207	0.711994343
						0.989569455	0.989836085	0.989690469

T1		100	wcc2					
11	0	100	wcc2 2	2	cupport	Recall	Precision	facero
Λ	39816573		17751			0.997189647		f-score 0.99780324
1		3969809	198	3	4028446	0.985444263	0.93841739	0.98213006
2	3992		33343	45	37520	0.888672708		0.74244044
3	678	4		3557	5247		0.037333401	0.4034252
,	070	7	1000	3337	3247	0.887304451	0.725486255	0.78144974
						0.995983682		0.99607964
РΓ),T1	100	wcc2			0.555505002	0.550252425	0.55007504
	0	1	2	3	support	Recall	Precision	f-score
0	39808425		15291					0.99777693
1		3976521	133	0	4028446	0.987110414		0.98131562
2	4337		32989	13	37520	0.87923774	0.661102204	0.75472432
3	897	6	1487	2857	5247	0.54450162	0.331207976	0.41187919
						0.851958839	0.741617045	0.78642401
						0.995927091	0.996098142	0.99599268
FS	, T1	100	wcc2					
	0	1	2	3	support	Recall	Precision	f-score
0	39807808	96036	15699	9244	39928787	0.996970131	0.9985613	0.99776508
1	53009	3975313	124	0	4028446	0.986810547	0.976357792	0.98155634
2	3711	219	33271	319	37520	0.886753731	0.671774991	0.7644376
3	634	6	433	4174	5247	0.795502192	0.303850914	0.43973873
						0.91650915	0.737636249	0.79587444
						0.995921955	0.996166941	0.99601557
PD),FS,T1	100	wcc2					
	0	1	2	3	support	Recall	Precision	f-score
0	39811619	96749	15339	5080	39928787	0.997065576	0.998606423	0.9978354
1	49595	3978806	45	0	4028446	0.987677631	0.976209649	0.98191016
2	4998	212	32231	79	37520	0.859035181	0.661094474	0.7471776
3	965	3	1139	3140	5247	0.598437202	0.378358838	0.46360549
						0.860553897	0.753567346	0.79763216
						0.996040818	0.996194102	0.99609991
FS		100	wcc2					
	0	1	2	3	support	Recall	Precision	f-score
0	39669658	234912					0.994821866	0.99416561
1	197730	3830104	498	114	4028446	0.950764637	0.941020732	0.94586759
2	6400	5140	25961	19	37520	0.691924307	0.612865911	0.65
3	2354	3	876	2014	5247	0.383838384	0.177617074	0.24285542
						0.755009387	0.681581396	0.70822216
						0.98926675	0.989472916	0.98936058

1 54245 3974149 52 0 4028446 0.9865216 0.979428 0.9829	940119 961931 607329 883006 083486
1 54245 3974149 52 0 4028446 0.9865216 0.979428 0.9829	961931 607329 883006 083486
	607329 883006 083486
	883006 083486
2 6088 281 31011 140 37520 0.82651919 0.763216 0.7936	083486
3 2195 11 680 2361 5247 0.44997141 0.191671 0.268	
0.81511512 0.733187 0.760	
0.99623682 0.996396 0.9963	30/596
FS, T1 0 1 2 3 support Recall Precision f-score	
0 39822537 91080 9205 5965 39928787 0.99733901 0.998106 0.9977	722209
1 67336 3961046 64 0 4028446 0.98326898 0.977429 0.9803	340212
2 6331 375 30688 126 37520 0.81791045 0.764658 0.7903	388008
3 1912 15 176 3144 5247 0.59919954 0.340444 0.4341	194172
0.8494295 0.770159 0.800	066115
43817415 0.99585034 0.995935 0.995	588679
T1 0 1 2 3 support Recall Precision f-score	
0 39840970 73521 8501 5795 39928787 0.99780066 0.998237 0.9980	018753
1 62198 3966179 68 1 4028446 0.98454317 0.981687 0.9831	113203
2 6227 446 30560 287 37520 0.81449893 0.776304 0.7949	943163
3 1941 19 237 3050 5247 0.58128454 0.333954 0.4242	200278
0.84453183 0.772546 0.8000	068849
43840759 0.99638089 0.996453 0.9964	412471

	0	1	2	3	support	Recall	Precision	f-score
0	39821311	87671	12952	6853	39928787	0.99730831	0.998482868	0.99789524
1	53799	3974473	174	0	4028446	0.98660203	0.978368421	0.98246798
2	4661	193	32523	143	37520	0.8668177	0.701349953	0.77535403
3	2046	11	723	2467	5247	0.47017343	0.260699567	0.33541808
						0.83022537	0.734725202	0.77278383
						0.99615395	0.996299924	0.99621402
	0	1	2	3	support	Recall	Precision	f-score
0	39827414	81804	14408	5161	39928787	0.99746116	0.998388161	0.99792444
1	57210	3971015	221	0	4028446	0.98574363	0.979776125	0.98275082
2	5170	157	32178	15	37520	0.8576226	0.670948102	0.75288667
3	1919	6	1152	2170	5247	0.41356966	0.295398857	0.34463591
						0.81359926	0.736127811	0.76954946
						0.99619948	0.996321077	0.99624836
	0	1	2	3	support	Recall	Precision	f-score
0	39822228	84083	15074	7402	39928787	0.99733127	0.998225854	0.99777836
1	63932	3964347	141	26	4028446	0.98408841	0.979197336	0.98163678
2	6386	127	30804	203	37520	0.82100213	0.650587142	0.72592732
3	458	11	1329	3449	5247	0.657328	0.311281588	0.42249035
						0.86493745	0.73482298	0.7819582
						0.99592791	0.996105328	0.99600009

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	0	1	2	3	support	Recall	Precision	f-score
0	39819616	90841	13400	4930	39928787	0.99726586	0.99846848	0.99786681
1	53742	3974554	150	0	4028446	0.98662214	0.97759879	0.98208974
2	5360	224	31909	27	37520	0.85045309	0.6875606	0.7603808
3	1976	10	950	2311	5247	0.44044216	0.31796918	0.36931682
						0.81869581	0.74539926	0.77741354
						0.99609977	0.99621148	0.99614486
	0	1	2	3	support	Recall	Precision	f-score
0	39808156	96839	15501	8291	39928787	0.99697885	0.99850318	0.99774043
1	51741	3976530	175	0	4028446	0.98711265	0.97615442	0.98160295
2	6188	294	30966	72	37520	0.82531983	0.65461695	0.73012355
3	1746	6	662	2833	5247	0.53992758	0.2530368	0.34458432
						0.83733473	0.72057784	0.76351281
						0.99587466	0.99607489	0.99595686
	0	1	2	3	support	Recall	Precision	f-score
0	39818221	81858	18464	10244	39928787	0.99723092	0.99842119	0.9978257
1	60349	3967726	368	3	4028446	0.98492719	0.97975825	0.98233592
2	2155	109	35229	27	37520	0.93893923	0.6475324	0.76647267
3	461	6	344	4436	5247	0.84543549	0.30156356	0.44455579
						0.94163321	0.73181885	0.79779752
				43825612		0.99603664	0.99633018	0.99614426

Appendix B

CNN Summary

	Layer (type)	Output	Sha	ре ====		Param #	Connected to
3	img (InputLayer)	(None,	256	, 25	5, 2)	0	
	conv2d_1 (Conv2D)	(None,	256	, 25	6, 16)	304	img[0][0]
6 7 8	batch_normalization_1 (BatchNor	(None,	256	, 25	5, 16)	64	conv2d_1[0][0]
	activation_1 (ReLu)	(None,	256	, 25	5, 16)	0	batch_normalization_1[0][0]
	conv2d_2 (Conv2D)	(None,	256	, 25	5, 16)	2320	activation_1[0][0]
	batch_normalization_2 (BatchNor	(None,	256	, 25	6, 16)	64	conv2d_2[0][0]
	activation_2 (ReLu)	(None,	256	, 25	6, 16)	0	batch_normalization_2[0][0]
	max_pooling2d_1 (MaxPooling2D)	(None,	128	, 12	3, 16)	0	activation_2[0][0]
	dropout_1 (Dropout)	(None,	128	, 12	3, 16)	0	max_pooling2d_1[0][0]
0	conv2d_3 (Conv2D)	(None,	128	, 12	3, 32)	4640	dropout_1[0][0]
22	batch_normalization_3 (BatchNor	(None,	128	, 12	3, 32)	128	conv2d_3[0][0]
4 5	activation_3 (Relu)	(None,	128	, 12	3, 32)	0	batch_normalization_3[0][0]
6	conv2d_4 (Conv2D)	(None,	128	, 12	3, 32)	9248	activation_3[0][0]
8	batch_normalization_4 (BatchNor						conv2d_4[0][0]
0	activation_4 (ReLu)	(None,					batch_normalization_4[0][0]
2	max pooling2d 2 (MaxPooling2D)					0	activation 4[0][0]
4	dropout 2 (Dropout)	(None,				0	max pooling2d 2[0][0]
6	_	(None,				18496	dropout 2[0][0]
8	batch normalization 5 (BatchNor					256	conv2d 5[0][0]
0	activation 5 (ReLu)	(None,				0	
2							batch_normalization_5[0][0]
4	conv2d_6 (Conv2D)	(None,				36928	activation_5[0][0]
6	batch_normalization_6 (BatchNor					256	conv2d_6[0][0]
8	activation_6 (ReLu)	(None,				0	batch_normalization_6[0][0]
0		(None,				0	activation_6[0][0]
2	dropout_3 (Dropout)	(None,				0	max_pooling2d_3[0][0]
4	conv2d_7 (Conv2D)	(None,				73856	dropout_3[0][0]
6	batch_normalization_7 (BatchNor					512	conv2d_7[0][0]
8	activation_7 (ReLu)	(None,				0	batch_normalization_7[0][0]
0	conv2d_8 (Conv2D)	(None,				147584	activation_7[0][0]
1	batch_normalization_8 (BatchNor	(None,	32,	32,	128)	512	conv2d_8[0][0]
3	activation_8 (ReLu)	(None,	32,	32,	128)	0	batch_normalization_8[0][0]
		(None,	16,	16,	128)	0	activation_8[0][0]
	dropout_4 (Dropout)	(None,	16,	16,	128)	0	max_pooling2d_4[0][0]
	conv2d_9 (Conv2D)	(None,	16,	16,	256)	295168	dropout_4[0][0]
	batch_normalization_9 (BatchNor	(None,	16,	16,	256)	1024	conv2d_9[0][0]
	activation_9 (ReLu)	(None,	16,	16,	256)	0	batch_normalization_9[0][0]
	conv2d_10 (Conv2D)	(None,	16,	16,	256)	590080	activation_9[0][0]
7	batch_normalization_10 (BatchNo	(None,	16,	16,	256)	1024	conv2d_10[0][0]
	activation_10 (ReLu)	(None,	16,	16,	256)	0	batch_normalization_10[0][0]
	conv2d_transpose_1 (Conv2DTrans	(None,	32,	32,	128)	295040	activation_10[0][0]
	concatenate_1 (Concatenate)	(None,	32,	32,	256)	0	conv2d_transpose_1[0][0]
5	durant F (F	/27	20	20	25.00		activation_8[0][0]
7	dropout_5 (Dropout)	(None,				0	concatenate_1[0][0]
9	conv2d_11 (Conv2D)	(None,				295040	dropout_5[0][0]
	batch normalization 11 (BatchNo	(None.	32,	32,	128)	512	conv2d_11[0][0]

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	O: 100010 Waltim Desitep inet of						
	activation_11 (ReLu)	(None,	32,	32, 128	8)	0	batch_normalization_11[0][0]
	conv2d_12 (Conv2D)	(None,	32,	32, 128	8)	147584	activation_11[0][0]
	batch_normalization_12 (BatchNo	(None,	32,	32, 128	8)	512	conv2d_12[0][0]
97 98 99	activation_12 (ReLu)	(None,	32,	32, 128	8)	0	batch_normalization_12[0][0]
	conv2d_transpose_2 (Conv2DTrans	(None,	64,	64, 64))	73792	activation_12[0][0]
	concatenate_2 (Concatenate)	(None,	64,	64, 128	8)	0	conv2d_transpose_2[0][0] activation_6[0][0]
	dropout_6 (Dropout)	(None,	64,	64, 128	8)	0	concatenate_2[0][0]
	conv2d_13 (Conv2D)	(None,	64,	64, 64))	73792	dropout_6[0][0]
	batch_normalization_13 (BatchNo	(None,	64,	64, 64))	256	conv2d_13[0][0]
	activation_13 (ReLu)	(None,	64,	64, 64))	0	batch_normalization_13[0][0]
	conv2d_14 (Conv2D)	(None,	64,	64, 64))	36928	activation_13[0][0]
	batch_normalization_14 (BatchNo	(None,	64,	64, 64))	256	conv2d_14[0][0]
	activation_14 (ReLu)	(None,	64,	64, 64))	0	batch_normalization_14[0][0]
119 120	conv2d_transpose_3 (Conv2DTrans	(None,	128,	128, 3	32)	18464	activation_14[0][0]
121 122 123	concatenate_3 (Concatenate)	(None,	128,	128, 6	64)	0	conv2d_transpose_3[0][0] activation_4[0][0]
	dropout_7 (Dropout)	(None,	128,	128,	64)	0	concatenate_3[0][0]
	conv2d_15 (Conv2D)	(None,	128,	128, 3	32)	18464	dropout_7[0][0]
	batch_normalization_15 (BatchNo	(None,	128,	128, 3	32)	128	conv2d_15[0][0]
	activation_15 (ReLu)	(None,	128,	128, 3	32)	0	batch_normalization_15[0][0]
132 133	conv2d_16 (Conv2D)	(None,	128,	128, 3	32)	9248	activation_15[0][0]
134 135	batch_normalization_16 (BatchNo	(None,	128,	128,	32)	128	conv2d_16[0][0]
136 137	activation_16 (ReLu)	(None,	128,	128, 3	32)	0	batch_normalization_16[0][0]
138 139	conv2d_transpose_4 (Conv2DTrans	(None,	256,	256, 1	16)	4624	activation_16[0][0]
140 141 142	concatenate_4 (Concatenate)	(None,	256,	256, 3	32)	0	conv2d_transpose_4[0][0] activation_2[0][0]
	dropout_8 (Dropout)	(None,	256,	256, 3	32)	0	concatenate_4[0][0]
145 146	conv2d_17 (Conv2D)	(None,	256,	256,	16)	4624	dropout_8[0][0]
147 148	batch_normalization_17 (BatchNo	(None,	256,	256, 1	16)	64	conv2d_17[0][0]
	activation_17 (ReLu)	(None,	256,	256, 1	16)	0	batch_normalization_17[0][0]
	conv2d_18 (Conv2D)	(None,	256,	256, 3	16)	2320	activation_17[0][0]
	batch_normalization_18 (BatchNo	(None,	256,	256,	16)	64	conv2d_18[0][0]
	activation_18 (sigmoid)	(None,	256,	256,	16)	0	batch_normalization_18[0][0]
157	conv2d_19 (Conv2D)			256,			activation_18[0][0]
159 160	Total params: 2,164,500 Trainable params: 2,161,556 Non-trainable params: 2,944						

Appendix C

Code

File - C:\Users\User\Desktop\Martin\DL\scripts\load.py

```
import random
 2 import numpy as np
 3 from scripts.unet import get unet, get unet2, get unet3
 4 from scripts.load data import get_data, get_data2, get_test_data, get_test_data2, get_test_data3
5 from scripts.plot import plot_img, plot_categorical, plot_categorical2
 6 from scripts.weighted_categorical_crossentropy import weighted_categorical_crossentropy
 7 from scripts.save_data import save_array, save_array_from_categorical
 8 from scripts.score import score
10 from sklearn.model_selection import train_test_split
11 from PIL import Image
13 from keras.models import Model
14 from keras.layers import Input, BatchNormalization, Activation, Dropout
15 from keras.optimizers import Adam
16 from keras.utils import multi_gpu_model
17 from keras.layers.convolutional import Conv2D, Conv2DTranspose
18 from keras.layers.pooling import MaxPooling2D
19 from keras.layers.merge import concatenate
20 from keras.callbacks import EarlyStopping, ModelCheckpoint, ReduceLROnPlateau
21 from keras.models import load_model
23 IMG_HEIGHT = 256
24 IMG_WIDTH = 256
25 IMG_DEPTH = 256
26 IMG CHANNELS = 1
27 NR CLASSES = 4
28 path_train2 = '/Users/User/Desktop/Martin/DL/input2/train/'
29 path_test2 = '/Users/User/Desktop/Martin/DL/input2/test/'
30
31 def load():
      print("
32
                                     RUNNING LOAD
33
       X= get_test_data(path_test2)
34
35
36
       # Split train and valid
37
       input_img = Input((IMG_HEIGHT, IMG_WIDTH, 1), name='img')
38
       model = get_unet(input_img, n_filters=16, dropout=0.02, batchnorm=True)
39
       model = multi_gpu_model(model, gpus=2)
40
41
       # Load best model
42
       #keras.losses.custom_loss = weighted_categorical_crossentropy
43
        #model = load_model('unet_model.h5', custom_objects={'weighted_categorical_crossentropy':
   weighted_categorical_crossentropy})
44
       model.load_weights('unet_100_FS_wcc_weights.h5')
45
46
47
        # Predict on train, val and test
48
49
       preds_train = model.predict(X, verbose=1) #X_train)
50
51
       save_array_from_categorical(preds_train,"labelFSwcc100")
52
53
54
55
       model.load_weights('unet_100_FS_wcc2_weights.h5')
       preds_train = model.predict(X, verbose=1) # X_train
       save_array_from_categorical(preds_train, "labelFSwcc2100")
57
       model.load_weights('unet_100_FS_cc_weights.h5')
       preds_train = model.predict(X, verbose=1) # X_train
save_array_from_categorical(preds_train, "labelFScc100")
58
59
```

File - C:\Users\User\Desktop\Martin\DL\scripts\plot.py

```
import random
 2 import numpy as np
 3 import matplotlib.pyplot as plt
 5 IMG HEIGHT = 256
 6 IMG_WIDTH = 256
 7 IMG DEPTH = 256
 8 IMG CHANNELS = 1
 9 NR_CLASSES = 4
1.0
11 def plot_categorical(nr,X_train, y_train):
       y train = y train[nr]
13
      y_train = y_train.reshape(IMG_WIDTH * IMG_DEPTH, NR_CLASSES)
y_train = np.argmax(y_train, axis=1)
y_train = y_train.reshape(IMG_WIDTH, IMG_DEPTH)
15
16
17
18
       has_mask = y_train.max() > 0
19
20
21
22
       fig, ax = plt.subplots(1, 2, figsize=(20, 10))
       ax[0].imshow(X_train[nr, ..., 0], cmap='gray', interpolation='bilinear')
23
       if has mask:
24
            ax[0].contour(y_train.squeeze(), colors='k', levels=[0.5])
25
26
       ax[0].set_title('First')
27
       ax[1].imshow(y train.squeeze(), interpolation='bilinear', cmap='gray')
28
       #ax[1].imshow(y_train[ix], interpolation='bilinear', cmap='gray')
29
       ax[1].set_title('Second');
30
       plt.show()
31
32 def plot categorical2(nr, X train, y train, pred, pred2):
34
        fig, ax = plt.subplots(1, 4, figsize=(20, 10))
35
       y_train = y_train[nr]
36
37
       y_train = y_train.reshape(IMG_WIDTH * IMG_DEPTH, NR_CLASSES)
38
       y_train = np.argmax(y_train, axis=1)
39
40
       y_train = y_train.reshape(IMG_WIDTH, IMG_DEPTH)
41
       has mask = y train.max() > 0
42
43
       ax[0].imshow(X_train[nr, ..., 0], cmap='gray', interpolation='bilinear')
44
       if has_mask:
45
           ax[0].contour(y_train.squeeze(), colors='k', levels=[0.5])
46
       ax[0].set_title('Input + Countour')
47
48
       ax[1].imshow(y_train.squeeze(), interpolation='bilinear', cmap='gray')
49
       \verb|#ax[1].imshow(y\_train[ix], interpolation='bilinear', cmap='gray')|\\
       ax[1].set_title('Ground Truth');
50
51
52
53
       pred2 = pred2[nr]
54
55
       pred2 = pred2.reshape(IMG_WIDTH * IMG_DEPTH, NR_CLASSES)
56
       pred2 = np.argmax(pred2, axis=1)
57
       pred2 = pred2.reshape(IMG_WIDTH, IMG_DEPTH)
58
59
60
       has_{mask} = pred2.max() > 0
61
       ax[2].imshow(pred[nr, ..., 0], cmap='gray', interpolation='bilinear')
62
63
64
            ax[2].contour(pred2.squeeze(), colors='k', levels=[0.5])
       ax[2].set_title('Scores')
65
66
       ax[3].imshow(pred2.squeeze(), interpolation='bilinear', cmap='gray')
67
       ax[3].set_title('Result');
68
       plt.show()
69
70
71 def plot_categorical_argmax(nr, X_train, y_train, pred, pred2):
72
73
74
       fig, ax = plt.subplots(1, 4, figsize=(20, 10))
       y_train = y_train[nr]
75
       y_train = y_train.reshape(IMG_WIDTH * IMG_DEPTH, NR_CLASSES)
77
       y_train = np.argmax(y_train, axis=1)
78
       y_train = y_train.reshape(IMG_WIDTH, IMG_DEPTH)
79
80
       has mask = y train.max() > 0
81
82
        ax[0].imshow(X_train[nr, ..., 0], cmap='gray', interpolation='bilinear')
83
84
       if has_mask:
           ax[0].contour(y_train.squeeze(), colors='k', levels=[0.5])
85
       ax[0].set title('Input + Countour')
86
87
        ax[1].imshow(y_train.squeeze(), interpolation='bilinear', cmap='gray')
88
       #ax[1].imshow(y_train[ix], interpolation='bilinear', cmap='gray')
ax[1].set_title('Ground Truth');
89
90
```

File - C:\Users\User\Desktop\Martin\DL\scripts\plot.py

```
pred2 = pred2[nr]
 92
 93
        pred2 = pred2.reshape(IMG_WIDTH * IMG_DEPTH, NR_CLASSES)
 95
        pred2 = np.argmax(pred2, axis=1)
        pred2 = pred2.reshape(IMG_WIDTH, IMG_DEPTH)
 96
 97
 98
        has mask = pred2.max() > 0
100
        ax[2].imshow(pred[nr, ..., 0], cmap='gray', interpolation='bilinear')
101
        if has_mask:
102
            ax[2].contour(pred2.squeeze(), colors='k', levels=[0.5])
103
        ax[2].set_title('Scores')
104
105
        ax[3].imshow(pred2.squeeze(), interpolation='bilinear', cmap='gray')
106
107
        ax[3].set_title('Result');
        plt.show()
108
109
110 def plot_img(X_train, y_train):
        ix = random.randint(0, len(X_train))
has_mask = y_train[ix].max() > 0
111
112
113
114
        fig, ax = plt.subplots(1, 2, figsize=(20, 10))
115
116
117
        ax[0].imshow(X_train[ix, ..., 0], cmap='seismic', interpolation='bilinear')
        if has_mask:
118
            ax[0].contour(y_train[ix].squeeze(), colors='k', levels=[0.5])
119
        ax[0].set_title('Seismic')
120
        ax[1].imshow(y_train[ix].squeeze(), interpolation='bilinear', cmap='gray')
ax[1].set_title('Salt');
121
122
123
124
        print("Done! img show")
```

File - C:\Users\User\Desktop\Martin\DL\scripts\unet.py

```
from keras.models import Model
  2 from keras.layers import Input, BatchNormalization, Activation, Dropout
  3 from keras.layers.convolutional import Conv2D, Conv2DTranspose, Conv3D, Conv3DTranspose
   4 from keras.layers.pooling import MaxPooling2D, MaxPooling3D
  5 from keras.layers.merge import concatenate
  8 def conv2d block(input tensor, n filters, kernel size=3, batchnorm=True):
10
                {\tt x = Conv2D(filters=n\_filters, \ kernel\_size=(kernel\_size, \ kernel\_size), \ kernel\_initializer="he\_normal", leaves to be a size of the large term of th
11
                                           padding="same")(input_tensor)
               if batchnorm:
12
                        x = BatchNormalization()(x)
13
                 x = Activation("relu")(x)
1.5
                 # second layer
               16
17
18
                if batchnorm:
19
                       x = BatchNormalization()(x)
20
                x = Activation("relu")(x)
21
                return x
23 def conv3d_block(input_tensor, n_filters, kernel_size=3, batchnorm=True):
25
                x = \texttt{Conv3D}(\texttt{filters=n\_filters}, \ \texttt{kernel\_size=(kernel\_size}, \ \texttt{kernel\_size}), \ \texttt{kernel\_initializer="he\_normal"}, \\ \texttt{kernel\_size}) = \texttt{kernel\_initializer="he\_normal"}, \\ \texttt{kernel\_size}) = \texttt{kernel\_size} = \texttt{kernel\_s
26
                                         padding="same") (input tensor)
27
                if batchnorm:
28
                        x = BatchNormalization()(x)
29
                x = Activation("relu")(x)
30
                 # second layer
31
               x = Conv3D(filters=n filters, kernel size=(kernel size, kernel size, kernel size), kernel initializer="he normal",
32
                                        padding="same")(x)
 33
                if batchnorm:
                x = BatchNormalization()(x)
x = Activation("relu")(x)
34
35
36
                return x
37
39 def get_unet2(input_img, n_filters=16, dropout=0.5, batchnorm=True):
40
                # contracting path
41
                \verb|c1 = conv2d_block(input_img, n_filters=n_filters * 1, kernel_size=3, batchnorm=batchnorm)|\\
               p1 = MaxPooling2D((2, 2))(c1)
p1 = Dropout(dropout * 0.5)(p1)
42
43
44
45
               c2 = conv2d_block(p1, n_filters=n_filters * 2, kernel_size=3, batchnorm=batchnorm)
p2 = MaxPooling2D((2, 2))(c2)
46
                p2 = Dropout(dropout)(p2)
48
49
                \verb|c3| = \verb|conv2d_block(p2, n_filters=n_filters * 4, kernel_size=3, batchnorm=batchnorm)|\\
                p3 = MaxPooling2D((2, 2))(c3)
50
51
                p3 = Dropout (dropout) (p3)
53
                \verb|c4| = \verb|conv2d_b| | \verb|ck(p3, n_filters=n_filters * 8, kernel_size=3, batchnorm=batchnorm||
54
55
                p4 = MaxPooling2D(pool_size=(2, 2))(c4)
                p4 = Dropout (dropout) (p4)
56
57
                c5 = conv2d_block(p4, n_filters=n_filters * 16, kernel_size=3, batchnorm=batchnorm)
58
59
60
                 # expansive path
                u6 = Conv2DTranspose(n filters * 8, (3, 3), strides=(2, 2), padding='same')(c5)
61
                u6 = concatenate([u6, c4])
62
                u6 = Dropout(dropout)(u6)
63
                c6 = conv2d_block(u6, n_filters=n_filters * 8, kernel_size=3, batchnorm=batchnorm)
64
65
                u7 = Conv2DTranspose(n filters * 4, (3, 3), strides=(2, 2), padding='same')(c6)
66
                u7 = concatenate([u7, c3])
67
                u7 = Dropout(dropout)(u7)
68
                c7 = conv2d_block(u7, n_filters=n_filters * 4, kernel_size=3, batchnorm=batchnorm)
69
70
                u8 = Conv2DTranspose(n_filters * 2, (3, 3), strides=(2, 2), padding='same')(c7)
 71
                u8 = concatenate([u8, c2])
72
                u8 = Dropout(dropout)(u8)
 73
                c8 = conv2d_block(u8, n_filters=n_filters * 2, kernel_size=3, batchnorm=batchnorm)
74
75
                u9 = Conv2DTranspose(n filters * 1, (3, 3), strides=(2, 2), padding='same')(c8)
                u9 = concatenate([u9, c1], axis=3)
77
                u9 = Dropout(dropout)(u9)
78
                \verb|c9 = conv2d_block(u9, n_filters=n_filters * 1, kernel_size=3, batchnorm=batchnorm||
79
80
                outputs = Conv2D(1, (1, 1), activation='sigmoid')(c9)
                model = Model(inputs=[input_img], outputs=[outputs])
82
                return model
83
84 ## Unte for categorical data
86 def get_unet(input_img, n_filters=16, dropout=0.5, batchnorm=True):
87
                 # contracting path
88
                \verb|c1| = \verb|conv2d_block(input_img, n_filters=n_filters * 1, | kernel_size=3, | batchnorm=batchnorm)|
                p1 = MaxPooling2D((2, 2))(c1)
p1 = Dropout(dropout * 0.5)(p1)
89
90
```

File - C:\Users\User\Desktop\Martin\DL\scripts\unet.py

```
c2 = conv2d_block(p1, n_filters=n_filters * 2, kernel_size=3, batchnorm=batchnorm)
 92
 93
        p2 = MaxPooling2D((2, 2))(c2)
        p2 = Dropout(dropout)(p2)
 95
 96
        \verb|c3| = \verb|conv2d_b| \verb|lock(p2|, n_filters=n_filters * 4|, kernel_size=3|, batchnorm=batchnorm||
 97
        p3 = MaxPooling2D((2, 2))(c3)
 98
        p3 = Dropout(dropout)(p3)
100
        {\tt c4 = conv2d\_block(p3, n\_filters=n\_filters * 8, kernel\_size=3, batchnorm=batchnorm)}
101
        p4 = MaxPooling2D(pool_size=(2, 2))(c4)
        p4 = Dropout(dropout)(p4)
103
104
        c5 = conv2d_block(p4, n_filters=n_filters * 16, kernel_size=3, batchnorm=batchnorm)
105
106
        # expansive path
107
        u6 = Conv2DTranspose(n filters * 8, (3, 3), strides=(2, 2), padding='same')(c5)
108
        u6 = concatenate([u6, c4])
109
        u6 = Dropout(dropout)(u6)
110
        c6 = conv2d_block(u6, n_filters=n_filters * 8, kernel_size=3, batchnorm=batchnorm)
112
        u7 = Conv2DTranspose(n filters * 4, (3, 3), strides=(2, 2), padding='same')(c6)
113
        u7 = concatenate([u7, c3])
114
        u7 = Dropout(dropout)(u7)
115
        \verb|c7 = conv2d_block(u7, n_filters=n_filters * 4, kernel_size=3, batchnorm=batchnorm||
116
        u8 = Conv2DTranspose(n filters * 2, (3, 3), strides=(2, 2), padding='same')(c7)
117
118
        u8 = concatenate([u8, c2])
119
        u8 = Dropout(dropout)(u8)
120
        \verb|c8| = \verb|conv2d_block| (u8, n_filters=n_filters * 2, kernel_size=3, batchnorm=batchnorm)|
121
        u9 = Conv2DTranspose(n_filters * 1, (3, 3), strides=(2, 2), padding='same')(c8)
123
        u9 = concatenate([u9, c1], axis=3)
124
        u9 = Dropout(dropout)(u9)
125
        \verb|c9 = conv2d_block(u9, n_filters=n_filters * 1, kernel_size=3, batchnorm=batchnorm)|\\
126
127
        outputs = Conv2D(4, (1, 1), activation='sigmoid')(c9)
128
        model = Model(inputs=[input_img], outputs=[outputs])
        model.summary()
129
130
        return model
131
132 def get_unet3(input_img, n_filters=16, dropout=0.5, batchnorm=True):
133
        # contracting path
134
        c1 = conv3d_block(input_img, n_filters=n_filters * 1, kernel_size=3, batchnorm=batchnorm)
        p1 = MaxPooling3D((2, 2, 2))(c1)
p1 = Dropout(dropout * 0.5)(p1)
135
136
137
138
        c2 = conv3d_block(p1, n_filters=n_filters * 2, kernel_size=3, batchnorm=batchnorm)
139
        p2 = MaxPooling3D((2, 2, 2))(c2)
140
        p2 = Dropout(dropout)(p2)
141
142
        c3 = conv3d_block(p2, n_filters=n_filters * 4, kernel_size=3, batchnorm=batchnorm)
143
        p3 = MaxPooling3D((2, 2, 2))(c3)
144
        p3 = Dropout(dropout)(p3)
145
146
        {\tt c4 = conv3d\_block(p3, n\_filters=n\_filters * 8, kernel\_size=3, batchnorm=batchnorm)}
147
        p4 = MaxPooling3D(pool_size=(2, 2, 2))(c4)
148
        p4 = Dropout(dropout) (p4)
149
150
        c5 = conv3d_block(p4, n_filters=n_filters * 16, kernel_size=3, batchnorm=batchnorm)
151
152
153
         u6 = \texttt{Conv3DTranspose} ( \texttt{n\_filters * 8, (3, 3, 3), strides} = (2, 2, 2), \texttt{padding} = \texttt{'same'}) (c5) 
154
        u6 = concatenate([u6, c4])
155
        u6 = Dropout(dropout)(u6)
156
        c6 = conv3d_block(u6, n_filters=n_filters * 8, kernel_size=3, batchnorm=batchnorm)
157
158
         u7 = \texttt{Conv3DTranspose}(n\_filters * 4, (3, 3, 3), \texttt{strides} = (2, 2, 2), \texttt{padding} = \texttt{'same'}) (c6) 
159
        u7 = concatenate([u7, c3])
160
        u7 = Dropout(dropout)(u7)
161
        c7 = conv3d_block(u7, n_filters=n_filters * 4, kernel_size=3, batchnorm=batchnorm)
162
163
         u8 = Conv3DTranspose(n\_filters * 2, (3, 3, 3), strides=(2, 2, 2), padding='same') (c7) 
164
        u8 = concatenate([u8, c2])
        u8 = Dropout(dropout)(u8)
165
166
        c8 = conv3d_block(u8, n_filters=n_filters * 2, kernel_size=3, batchnorm=batchnorm)
167
         \verb"u9 = Conv3DTranspose(n_filters * 1, (3, 3, 3), strides=(2, 2, 2), padding='same') (c8) 
168
        u9 = concatenate([u9, c1])#, axis=3
u9 = Dropout(dropout)(u9)
169
170
171
        c9 = conv3d_block(u9, n_filters=n_filters * 1, kernel_size=3, batchnorm=batchnorm)
172
173
        \verb"outputs" = \verb"Conv3D(4, (1, 1, 1), activation="sigmoid")(c9)
        model = Model(inputs=[input_img], outputs=[outputs])
174
175
        return model
176
```

File - C:\Users\User\Desktop\Martin\DL\scripts\score.py

```
2 import os
 3 import sys
 4 import numpy as np
 5 import matplotlib.pyplot as plt
 6 import nibabel as nib
 7 plt.style.use("ggplot")
 9 from tqdm import tqdm_notebook
10 from skimage.transform import resize
11 from nibabel.testing import data_path
13 IMG_HEIGHT = 400
14 IMG_WIDTH = 400
15 IMG_DEPTH = 275
16 IMG CHANNELS = 1
17 NR_CLASSES = 4
19
20 def score():
21
       print('score')
22
       # Get train and test IDs
23
       train_ids = next(os.walk('/Users/User/Desktop/Martin/DL/input2/score/'))[2]
25
26
       # Get and resize train images and masks
       Y_train = np.zeros((len(train_ids), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), dtype=np.float32)
27
       print('Getting and resizing train images and masks ... ')
28
       sys.stdout.flush()
29
30
31
       for n, id_ in tqdm_notebook(enumerate(train_ids), total=len(train_ids)):
    example_filename2 = os.path.join(data_path, '/Users/User/Desktop/Martin/DL/input2/score/' + id_)
32
            print(example filename2)
33
            img = nib.load(example_filename2)
34
35
36
           data = img.get_fdata()
           data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve range=True)
37
           data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH)
            data = np.rint(data)
38
39
           print(Y_train.shape)
40
            print(data.shape)
41
           Y train[n] = data
42
43
       score_value(0,Y_train[0],Y_train[1])
44
       score_value(1, Y_train[0], Y_train[1])
       score_value(2, Y_train[0], Y_train[1])
score_value(3, Y_train[0], Y_train[1])
45
46
47
48
49
50 def score_all_values(true_data, predicted_data):
51
       print(true data.shape)
       print(np.unique(true_data))
53
       print(predicted_data.shape)
54
55
       print(np.unique(predicted_data))
       print("All values equals:")
rigth_predictions = np.equal(true_data, predicted_data)
56
57
       print(np.sum(rigth_predictions))
58
       print("/")
59
60
       print(true_data.size)
       print('==
61
       print(np.sum(rigth predictions)/true data.size)
63 def score_value(value, true_data, predicted_data):
64
       true_values = np.isin(true_data, [value])
65
       true_values_predicted=predicted_data[true_data == [value]]
       predicted0=np.isin(true_values_predicted,[0])
predicted1=np.isin(true_values_predicted,[1])
66
67
68
       predicted2=np.isin(true_values_predicted,[2])
       predicted3=np.isin(true_values_predicted,[3])
all_prediction = np.sum(predicted0)+np.sum(predicted1)+np.sum(predicted2)+np.sum(predicted3)
69
70
```

File - C:\Users\User\Desktop\Martin\DL\scripts\train.py

```
import random
  2 import numpy as np
  3 import matplotlib.pyplot as plt
  5 plt.style.use("ggplot")
  8 from scripts.load data import get data, get data2, get data3
  9 from scripts.unet import get_unet, get_unet2, get_unet3
10 from scripts.plot import plot_img, plot_categorical
{\tt 11} \ {\tt from} \ {\tt scripts.weighted\_categorical\_crossentropy} \ {\tt import} \ {\tt weighted\_categorical\_crossentropy},
     weighted categorical crossentropy2
12
13 from sklearn.model_selection import train_test_split
15 from keras.layers import Input, BatchNormalization, Activation, Dropout
16 from keras.callbacks import EarlyStopping, ModelCheckpoint, ReduceLROnPlateau
 17 from keras.optimizers import Adam
18 from keras.utils import multi_gpu_model
20 # Set parameters for dimentions and data
21 IMG HEIGHT = 256
22 IMG_WIDTH = 256
23 IMG_DEPTH = 256
24 IMG CHANNELS = 1
25 NR CLASSES = 4
26 path train2 = '/Users/User/Desktop/Martin/DL/input2/train/'
27 path_test2 = '/Users/User/Desktop/Martin/DL/input2/test/'
28
29
30
31 def train():
            print("
                                                               RUNNING TRAIN
33
34
             \texttt{\#Get data from folder, see scipts} \\ \\ \texttt{load\_data.py for more}
35
            X, y = get data(path train2, train=True)
36
 37
             print(X.shape)
38
39
40
             #Split the data into train and validation
41
             X_train, X_valid, y_train, y_valid = train_test_split(X, y, test_size=0.14, random_state=2018)
42
43
             #Set image dimentions and size
             input_img = Input((IMG_HEIGHT, IMG_WIDTH, 1), name='img')
44
45
46
             #Create CNN unet model
47
             model = get_unet(input_img, n_filters=16, dropout=0.02, batchnorm=True)
48
49
             #RUN multiple GPUS, remove if only one gpu
50
             model = multi_gpu_model(model, gpus=2)
51
52
             #Compilie with loss function
     model.compile(optimizer=Adam(), loss=weighted_categorical_crossentropy(), metrics=["accuracy"]) # '
binary_crossentropy', weighted_categorical_crossentropy()
             #Print model summary
55
             model.summarv()
56
57
58
             {\it \#MODEL~SAVE~does~not~work~with~multiple~GPUS}
             #model.save('unet 100 PD model.h5')
59
60
             #Set early stopping, reduce learning rate, and save weights if better
61
             callbacks = [
                    EarlyStopping(patience=30, verbose=1),
ReduceLROnPlateau(factor=0.1, patience=6, min_lr=0.000001, verbose=1),
62
63
64
                   ModelCheckpoint('unet_100_FS_wcc_weights.h5', verbose=1, save_best_only=True, save_weights_only=True)
65
66
67
             #Train model
             results = model.fit(X_train, y_train, batch_size=32, epochs=100, class_weight='auto', callbacks=callbacks, validation_data=(X_valid, y_valid))
68
69
70
71
             72
73
             #Create CNN unet model
             model1 = get_unet(input_img, n_filters=16, dropout=0.02, batchnorm=True)
75
76
             #RUN multiple GPUS, remove if only one gpu
 77
             model1 = multi gpu model(model1, gpus=2)
78
 79
             #Compille with loss function
80
             modell.compile(optimizer=Adam(), loss=weighted_categorical_crossentropy2(), metrics=["accuracy"]) # '
     categorical\_crossentropy() \ , \ weighted\_categorical\_crossentropy() \ , weighted\_ca
81
             #Print model summary
82
             model1.summary()
 83
84
             #MODEL SAVE does not work with multiple GPUS
85
86
             #model.save('unet 100 PD model.h5')
             #Set early stopping, reduce learning rate, and save weights if better
```

File - C:\Users\User\Desktop\Martin\DL\scripts\train.py

```
callbacks1 = [
             EarlyStopping(patience=30, verbose=1),
ReduceLROnPlateau(factor=0.1, patience=6, min_lr=0.000001, verbose=1),
ModelCheckpoint('unet_100_FS_wcc2_weights.h5', verbose=1, save_best_only=True, save_weights_only=True)
 89
 90
 91
 92
 93
 94
         #Train model
 95
         results1 = model1.fit(X_train, y_train, batch_size=32, epochs=100, class_weight='auto', callbacks=callbacks1,
                                validation_data=(X_valid, y_valid))
 97
 98
        99
100
101
         plt.figure(figsize=(8, 8))
102
         plt.title("Learning curve")
    plt.plot(results.history["loss"], label="loss")
plt.plot(results.history["val_loss"], label="val_loss")
plt.plot( np.argmin(results.history["val_loss"]), np.min(results.history["val_loss"]), marker="x", color="r", label="best model")
="best model")
103
104
105
         plt.xlabel("Epochs")
plt.ylabel("log_loss")
106
107
108
         plt.legend
109
         plt.show()
110
111
112
113
         #Load best weights
         model.load_weights('unet_100_FST1_wcc_weights.h5')
114
115
         116
117
         \verb| model.evaluate(X_valid, y_valid, verbose=1)|\\
118
119
         print("DONE")
120
```

File - C:\Users\User\Desktop\Martin\DL\scripts_	_init_	py
1 #####		

```
import os
 2 import sys
 3 import numpy as np
 4 import matplotlib.pyplot as plt
 5 import nibabel as nib
 7 from keras.utils import to categorical
 8 from tgdm import tgdm notebook, thrange
 9 from skimage.transform import resize
10 from nibabel.testing import data_path
11 from keras.preprocessing.image import img_to_array, load_img
13 plt.style.use("ggplot")
15 # Set some parameters
16 IMG_HEIGHT = 256
17 IMG WIDTH = 256
18 IMG_DEPTH = 256
19 IMG CHANNELS = 1
20 NR CLASSES = 4
21 path train = '/Users/User/Desktop/Martin/DL/input2/train/'
22 path test = '/Users/User/Desktop/Martin/DL/input2/test/'
25 def get_data3(path, train=True):
        # Get train and test IDs
train_ids1 = next(os.walk(path + "imagesPD"))[2]
26
29
        # Get and resize train images and masks
30
         \textbf{X\_train = np.zeros((len(train\_ids1), IMG\_HEIGHT, IMG\_WIDTH, IMG\_DEPTH, 3), dtype=np.float32) } \\
        Y train = np.zeros((len(train_idsl), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES), dtype=np.float32)
31
32
        print('Getting and resizing train images and masks ... ')
33
        sys.stdout.flush()
34
35
36
        for n, id_ in tqdm_notebook(enumerate(train_ids1), total=len(train_ids1)):
            example_filename1 = os.path.join(data_path, path + 'imagesPD/' + id_)
example_filename2 = os.path.join(data_path, path + 'imagesFS/' + id_)
37
38
             example_filename3 = os.path.join(data_path, path + 'imagesT1/' + id_)
39
            print(example_filename1)
40
            print(example_filename2)
41
            print(example filename3)
42
            img1 = nib.load(example_filename1)
43
            data1 = imgl.get_fdata()
44
45
46
            img2 = nib.load(example filename2)
            data2 = img2.get_fdata()
48
            img3 = nib.load(example_filename3)
49
            data3 = img3.get_fdata()
51
            data1 = resize(data1, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve range=True)
            data1 = data1.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
53
54
55
56
            data2 = resize(data2, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
            data2 = data2.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
57
            data3 = resize(data3, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
58
59
60
            data3 = data3.reshape(IMG HEIGHT, IMG WIDTH, IMG DEPTH, IMG CHANNELS)
            data4 = np.concatenate((data1, data2, data3), axis=3)
61
            X train[n] = data4
63
64
65
            example_filename2 = os.path.join(data_path, path + 'masks/' + id_)
66
            img = nib.load(example_filename2)
67
            data = img.get_fdata()
68
            data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
69
70
            data = np.rint(data)
72
            data = to_categorical(data)
73
74
            data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES)
            Y train[n] = data
75
        X_train = X_train.reshape(len(train_ids1) * IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, 3)
Y_train = Y_train.reshape(len(train_ids1) * IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES)
77
78
79
        return X train, Y train
80
82 def get_data2(path, train=True):
83
        train ids1 = next(os.walk(path + "imagesFS"))[2]
84
        # Get and resize train images and masks
        X_train = np.zeros((len(train_ids1), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, 2), dtype=np.float32)
Y_train = np.zeros((len(train_ids1), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES), dtype=np.float32)
87
88
89
        print('Getting and resizing train images and masks ... ')
        sys.stdout.flush()
```

```
for n, id_ in tqdm_notebook(enumerate(train_ids1), total=len(train_ids1)):
 92
             example_filename1 = os.path.join(data_path, path + 'imagesFS' + id_)
example_filename2 = os.path.join(data_path, path + 'imagesT1' + id_)
 93
 95
             print(example_filename1)
 96
             print(example_filename2)
 97
             img1 = nib.load(example filename1)
 98
            data1 = img1.get_fdata()
100
             img2 = nib.load(example_filename2)
101
            data2 = img2.get_fdata()
103
             datal = resize(datal, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
104
            data1 = data1.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
105
            data2 = resize(data2, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
data2 = data2.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
106
107
108
109
             data3 = np.concatenate((data1, data2), axis=3)
110
            X train[n] = data3
112
113
             example_filename2 = os.path.join(data_path, path + 'masks/' + id_)
114
115
             img = nib.load(example_filename2)
            data = img.get_fdata()
116
117
118
             data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
119
             data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
120
             data = np.rint(data)
121
            data = to_categorical(data)
data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES)
123
             Y train[n] = data
124
        X_train = X_train.reshape(len(train_ids1) * IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, 2)
Y_train = Y_train.reshape(len(train_ids1) * IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES)
125
126
127
128
        return X train, Y train
129
130 ### to categorical get data
131
132 def get_data(path, train=True):
133
          Get train and test IDs
134
        train_ids = next(os.walk(path + "imagesFS"))[2]
135
136
        # Get and resize train images and masks
137
        X_train = np.zeros((len(train_ids), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS), dtype=np.float32)
138
        Y_train = np.zeros((len(train_ids), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES), dtype=np.float32)
139
        \verb|print('Getting and resizing train images and masks ... ')|\\
140
        sys.stdout.flush()
141
142
        for n, id_ in tqdm_notebook(enumerate(train_ids), total=len(train_ids)):
143
             example_filename = os.path.join(data_path, path + 'imagesFS/' + id_ )
144
             print(example_filename)
145
             img = nib.load(example_filename)
146
            data = img.get_fdata()
147
148
            data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
149
            data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
150
151
            X train[n] = data
152
153
154
             example_filename2 = os.path.join(data_path, path + 'masks/' + id_ )
             img = nib.load(example filename2)
156
            data = img.get_fdata()
157
158
             data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
             data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
159
160
             data = np.rint(data)
161
             data = to_categorical(data)
162
             data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES)
163
             Y train[n] = data
164
        X train = X train.reshape(len(train ids) * IMG HEIGHT, IMG WIDTH, IMG DEPTH, IMG CHANNELS)
165
        Y_train = Y_train.reshape(len(train_ids) * IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES)
166
167
168
        return X_train,Y_train
169
170
171 def get_test_data(path, train=True):
172
         # Get IDs
        test_ids = next(os.walk(path + "imagesFS"))[2]
173
174
175
         # Get and resize train images and masks
176
        X_test = np.zeros((len(test_ids), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS), dtype=np.float32)
177
        print('Getting and resizing train images and masks ... ')
178
         svs.stdout.flush()
179
180
        for n, id_ in tqdm_notebook(enumerate(test_ids), total=len(test_ids)):
```

File - C:\Users\User\Desktop\Martin\DL\scripts\load data.py

```
example_filename = os.path.join(data_path, path + 'imagesFS/' + id_
             print(example filename)
182
183
             img = nib.load(example filename)
184
             data = img.get_fdata()
185
186
             data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
187
             data = data.reshape(IMG HEIGHT, IMG WIDTH, IMG DEPTH, IMG CHANNELS)
188
189
             X test[n] = data
190
191
         X test = X test.reshape(len(test ids) * IMG HEIGHT, IMG WIDTH, IMG DEPTH, IMG CHANNELS)
192
193
194
         return X test
195
196 def get_test_data2(path, train=True):
197
         # Get IDs
198
         test_ids = next(os.walk(path + "imagesFS"))[2]
199
200
         \# Get and resize train images and masks
         X_test = np.zeros((len(test_ids), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, 2), dtype=np.float32)
2.01
202
         print('Getting and resizing train images and masks ...
203
         sys.stdout.flush()
204
205
         for n, id_ in tqdm_notebook(enumerate(test_ids), total=len(test_ids)):
             example_filename = os.path.join(data_path, path + 'imagesFS'' + id_ )
example_filename2 = os.path.join(data_path, path + 'imagesT1' + id_)
206
207
208
             print(example_filename)
209
             print(example_filename2)
210
             img = nib.load(example_filename)
211
             data = img.get_fdata()
img2 = nib.load(example filename2)
212
213
             data2 = img2.get_fdata()
214
215
             data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
216
             data = data.reshape(IMG HEIGHT, IMG WIDTH, IMG DEPTH, IMG CHANNELS)
217
             data2 = resize(data2, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
218
219
             data2 = data2.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
220
            data3 = np.concatenate((data, data2), axis=3)
data3 = data3.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, 2)
221
222
223
224
             X_{test[n]} = data3
225
226
        X test = X test.reshape(len(test ids) * IMG HEIGHT, IMG WIDTH, IMG DEPTH, 2)
227
228
229
         return X_test
230
231 def get_test_data3(path, train=True):
232
233
         test_ids = next(os.walk(path + "imagesPD"))[2]
234
235
         X_test = np.zeros((len(test_ids), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, 3), dtype=np.float32)
236
         \texttt{print}(\,\text{'Getting and resizing train images and masks ... ')}
237
         svs.stdout.flush()
238
239
240
         \label{lem:condition} \mbox{for n, id\_in tqdm\_notebook(enumerate(test\_ids), total=len(test\_ids)):}
             example_filename = os.path.join(data_path, path + 'imagesPD/' + id_
             example filename2 = os.path.join(data path, path + 'imagesFS/' + id )
241
242
             example_filename3 = os.path.join(data_path, path + 'imagesT1/' + id_)
243
             print(example_filename)
244
             print(example_filename2)
print(example_filename3)
245
246
             img = nib.load(example_filename)
247
             data = img.get_fdata()
248
             img2 = nib.load(example_filename2)
             data2 = img2.get_fdata()
249
250
             img3 = nib.load(example filename3)
251
             data3 = img3.get_fdata()
252
253
             data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
254
             data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
255
256
             data2 = resize(data2, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
257
             data2 = data2.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
258
259
             data3 = resize(data3, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
data3 = data3.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
260
261
262
             data4= np.concatenate((data, data2, data3), axis=3)
263
264
             X test[n] = data4
265
         X_test = X_test.reshape(len(test_ids) * IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, 3)
266
267
268
269
         return X test
270
```

File - C:\Users\User\Desktop\Martin\DL\scripts\load_data.py

```
272 def get_data3D(path, train=True):
273
274
          train_ids = next(os.walk(path + "images"))[2]
275
         X_train = np.zeros((len(train_ids), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS), dtype=np.float32)
Y_train = np.zeros((len(train_ids), IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES), dtype=np.float32)
276
277
278
          print('Getting and resizing train images and masks ... ')
279
          sys.stdout.flush()
280
281
282
          for n, id_ in tqdm_notebook(enumerate(train_ids), total=len(train_ids)):
    example_filename = os.path.join(data_path, path + 'images/' + id_ )
283
              print(example_filename)
284
               img = nib.load(example_filename)
285
              data = img.get_fdata()
286
287
              data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
288
289
290
291
292
              X_train[n] = data
293
294
               example_filename2 = os.path.join(data_path, path + 'masks/' + id_ + ".gz")
295
296
297
              img = nib.load(example_filename2)
data = img.get_fdata()
298
299
               data = resize(data, (IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH), mode='constant', preserve_range=True)
300
               data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, IMG_CHANNELS)
               data = np.rint(data)
302
               data = to categorical(data)
303
               data = data.reshape(IMG_HEIGHT, IMG_WIDTH, IMG_DEPTH, NR_CLASSES)
304
               Y_train[n] = data
305
306
          return X train, Y train
307
```

File - C:\Users\User\Desktop\Martin\DL\scripts\save_data.py

```
1 import numpy as np
 2 import nibabel as nib
 3 {f from} skimage.transform {f import} resize
 6 ORG_HEIGHT = 400
7 ORG_WIDTH = 400
 8 ORG_DEPTH = 275
10 IMG_HEIGHT = 256
11 IMG_WIDTH = 256
12 IMG_DEPTH = 256
13 IMG_CHANNELS = 1
14 NR_CLASSES = 4
False, order=0)
21
22
       empty_header = nib.NiftilHeader()
23
24
25
26
        \texttt{affine = np.array}([[0,\ 0,\ 1,\ 0],\ [-1,\ 0,\ 0,\ 0],\ [0,\ -1,\ 0,\ 0],\ [0,\ 0,\ 0,\ 1]]) 
       print(affine)
       clipped_img = nib.NiftilImage(data, affine, empty_header)
nib.save(clipped_img, '%s.nii' % (file_string))
27
28
29
30 def save_array(array, file_string):
31
32
        empty_header = nib.NiftilHeader()
33
34
35
        affine = np.diag([1, 1, 1, 1])
       clipped_img = nib.NiftilImage(array, affine, empty_header)
nib.save(clipped_img, '%s.nii' % (file_string))
36
```

File - C:\Users\User\Desktop\Martin\DL\scripts\weighted_categorical_crossentropy.py

```
1 from keras import backend as {\tt K}
   2 import numpy as np
  5 def weighted_categorical_crossentropy():
            weights = np.array([1.0, 1.05, 6.9, 11.8])
           weights = K.variable(weights)
 10
 11
12
            def loss(y_true, y_pred):
                 # Scale predictions so that the class probas of each sample sum to 1 y_pred /= K.sum(y_pred, axis=-1, keepdims=True)
 13
14
15
16
17
18
                  #Clip
                 y_pred = K.clip(y_pred, K.epsilon(), 1 - K.epsilon())
                  #Calculate
                 loss = y_true * K.log(y_pred) * weights
loss = -K.sum(loss, -1)
 19
                 return loss
 20
21
22
            return loss
22
23 def weighted_categorical_crossentropy2():
24
25 weights = np.array([1.0, 1.1, 12.0, 19.2])
26
27 weights = K.variable(weights)
28
29 def loss(y_true, y_pred):
30  # scale predictions so that the cl
31  y_pred /= K.sum(y_pred, axis=-1, k
32  # Clip
33  y_pred = K.clip(y_pred, K.epsilon()
34  # Calculate
35  loss = y_true * K.log(y_pred) * well
36  loss = -K.sum(loss, -1)
37 return loss
            weights = np.array([1.0, 1.1, 12.0, 19.7])
                  # scale predictions so that the class probas of each sample sum to 1
                  y_pred /= K.sum(y_pred, axis=-1, keepdims=True)
# Clip
                  # Calculate
" Calculate
" Calculate
" Calculate
" Calculate
                  # Calculate
loss = y_true * K.log(y_pred) * weights
loss = -K.sum(loss, -1)
  37
                  return loss
  38
 39
             return loss
```