



UNIVERSIDAD NACIONAL AUTÓNOMA DE MÉXICO

Posgrado en Ciencia e Ingeniería de la Computación

Curso de aprendizaje profundo



DoubleU-Net: A Deep Convolutional Neural Network for Medical Image Segmentation

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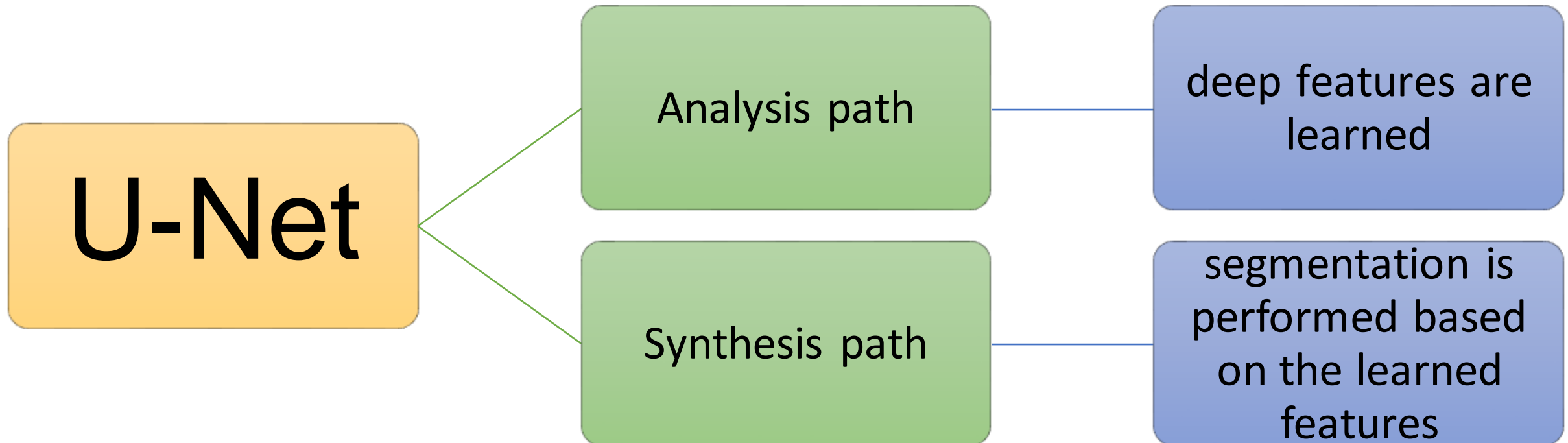
Miguel Ángel Veloz Lucas

Introduction

Segmentation task and U-Net network

In the state-of-the-art Convolutional Neural Networks (CNNs) have shown a good performance for automated medical image segmentation.

For example: U-Net



Related work in the field of medical image segmentation

U-Net Model

U-Net have gained significant popularity among semantic segmentation approach for 2D images.

In the medical imaging, there are many challenging images, which are usually missed out during colonoscopy examination and can develop into cancer if early detection is not performed. Therefore, there is a need for a more accurate medical image segmentation approach to deal with the challenging images

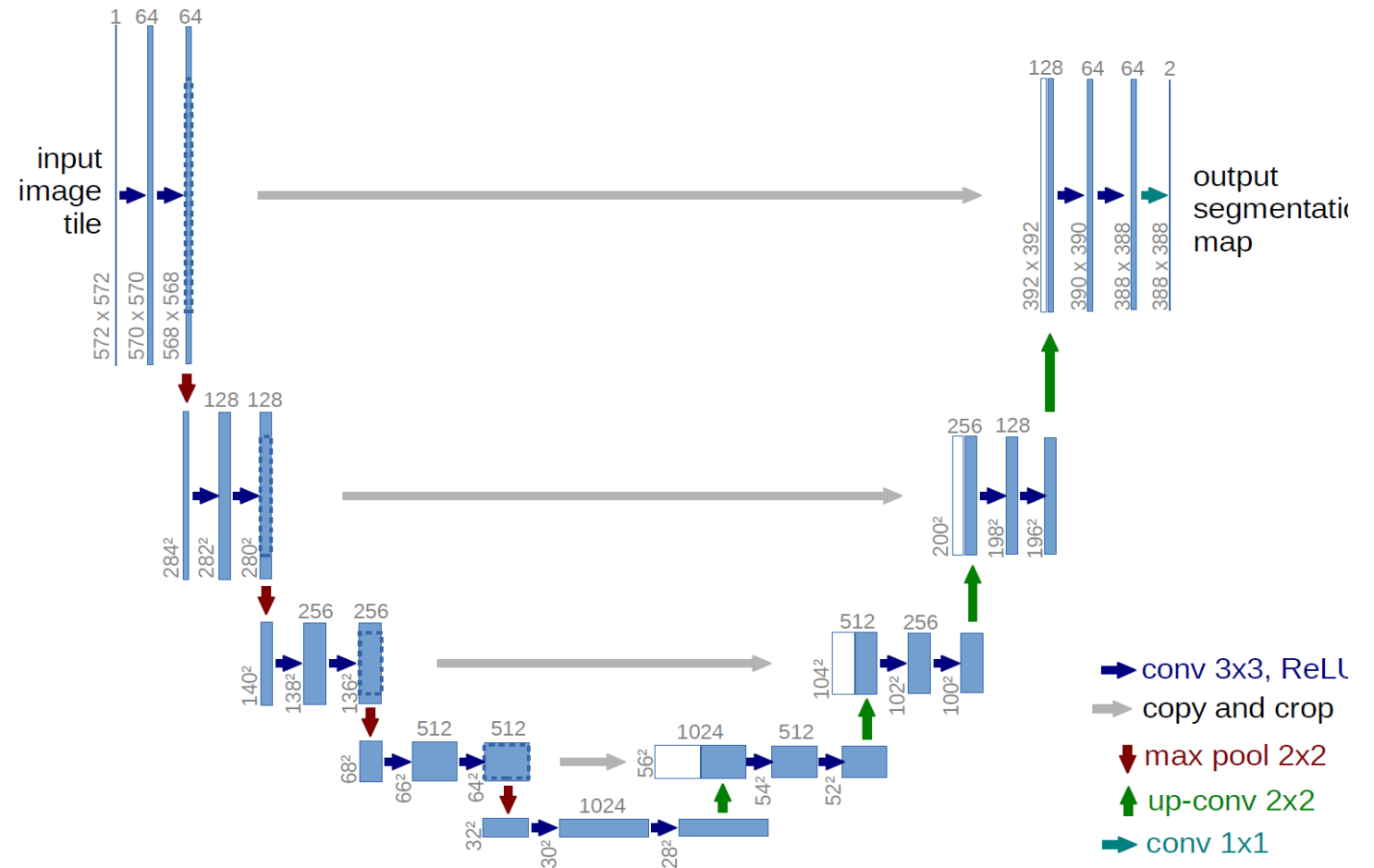
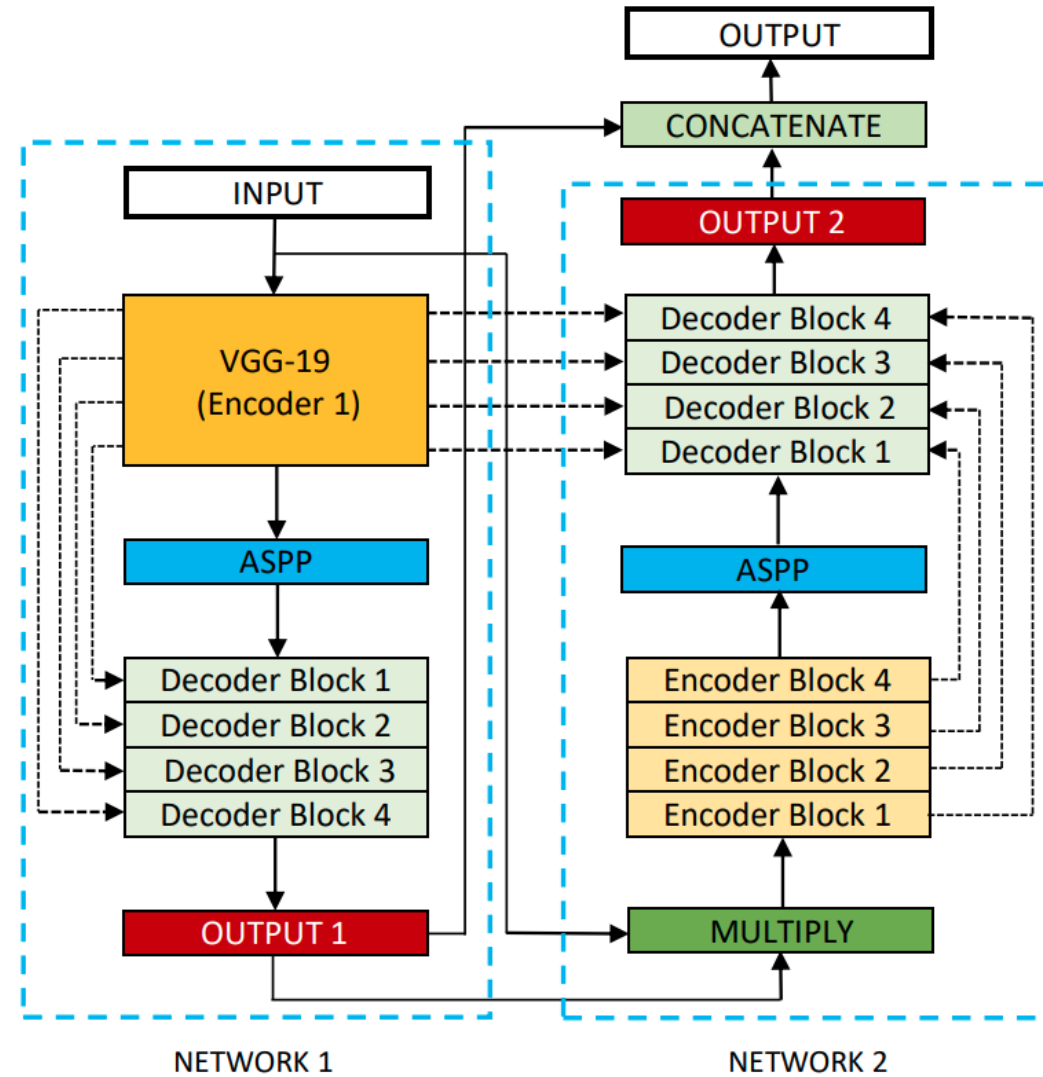


Image: <https://paperswithcode.com/method/u-net>

Double U-Net Architecture

The produced output feature map from NETWORK 1 can and concatenating with Output2 will produce a better segmentation mask than the previous one.



The experiments

Datasets

Dataset	Description	No. of Images	Input size	Aplication
a) 2015 MICCAI sub-challenge automatic polyp detection	It used the CVC-ClinicDB [1] for training and ETIS-Larib [2]	808	384 x 288	Colonoscopy
b) CVC-ClinicDB	For polyp segmentation	612	384 x 288	Colonoscopy
c) Lesion Boundary Segmentation dataset [3, 4]	It contains skin lesions and their corresponding annotations.	2594	Variable	Dermoscopy
d) 2018 Data Science Bowl challenge	For nuclei segmentation	670	256 x 256	Nuclei

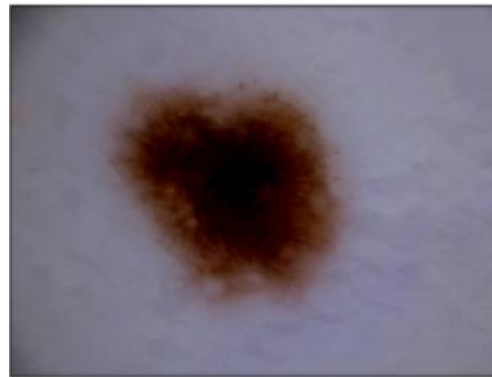
a)



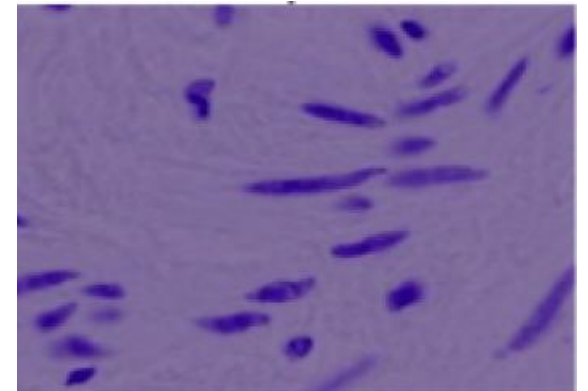
b)



c)



d)



The experiments

Evaluation metrics

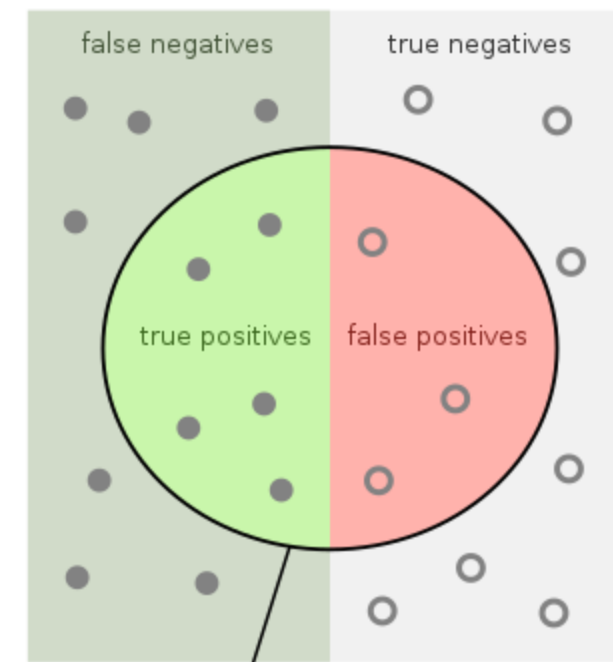
DoubleU-Net is evaluated on the basis of:


1. Precision:


$$PPV = \frac{TP}{TP + FP}$$

2. Recall:

$$TPR = \frac{TP}{P} = \frac{TP}{TP + FN}$$



Precision = 

Recall = 

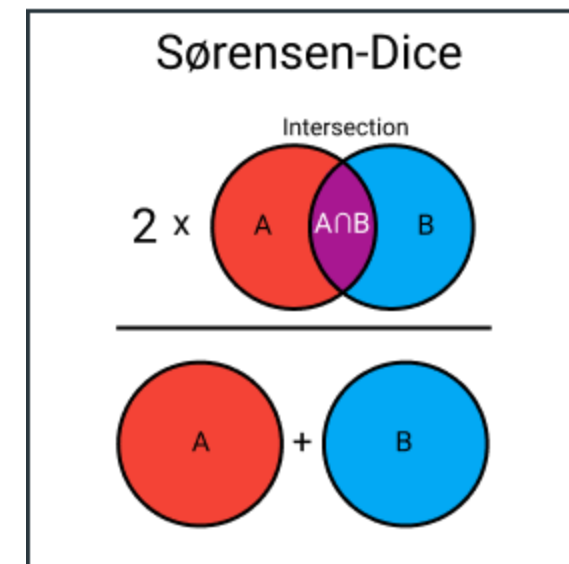
The experiments

Evaluation metrics

DoubleU-Net is evaluated on the basis of:

1. Sørensendice coefficient or Dice similarity coefficient (DSC) :

$$2 |X \cap Y| / (|X| + |Y|)$$



2. Mean Intersection over Union (mIoU):

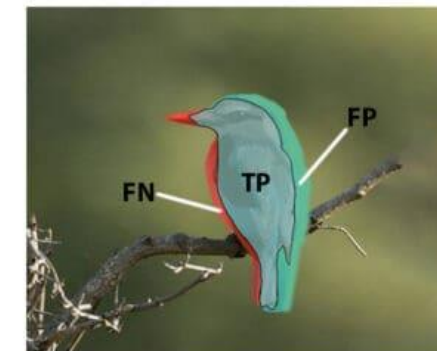
$$IoU = \frac{TP}{(TP + FP + FN)}$$



Ground Truth Mask



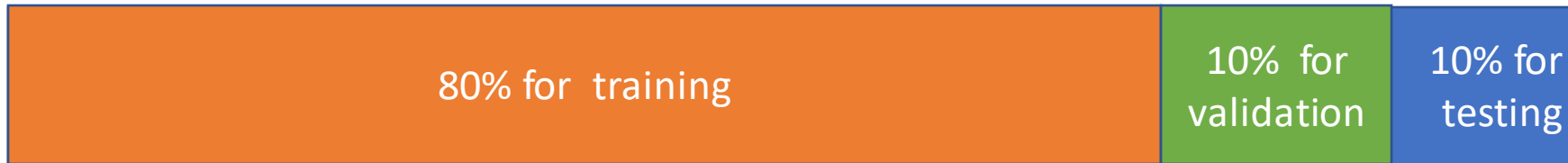
Predicted Mask



The experiments

Experiment setup and configuration

- All models were implemented using Keras framework [5] with Tensorflow 2.1.0 [6] as backend.
- A Volta 100 GPU and an Nvidia DGX-2 AI system were used.
- In all of the datasets:



- The original image size for the smaller dataset was used and resized the images to 384×512 for the Lesion Boundary segmentation challenge dataset to balance between training time and complexity.

Réplica de artículo

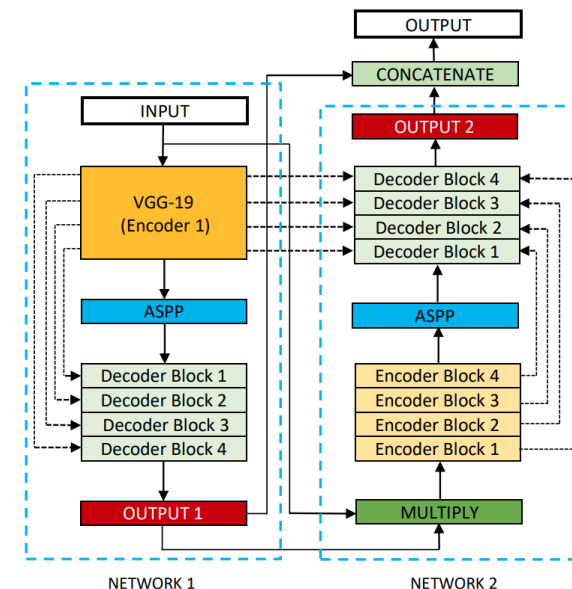
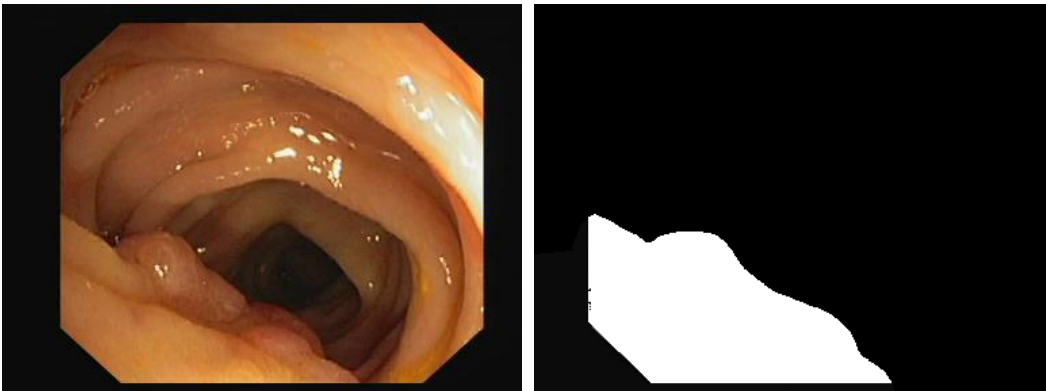
Base de datos ***CVC-ClinicDB***

Configuración de los experimentos

- Se utilizó la entropía binaria cruzada.
- El optimizador Nadam fue empleado.
- El tamaño del batch es 16 y la tasa de aprendizaje de $1e-5$.
- Se entrenó el modelo con 100 épocas.

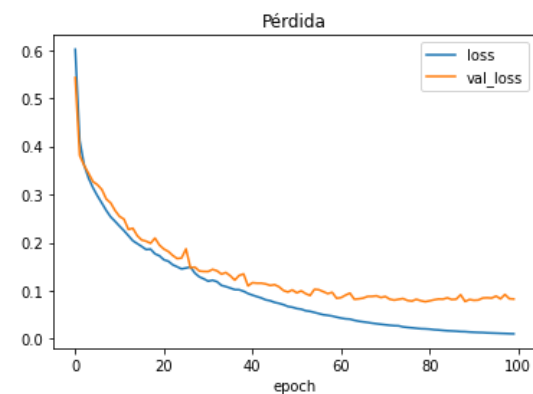
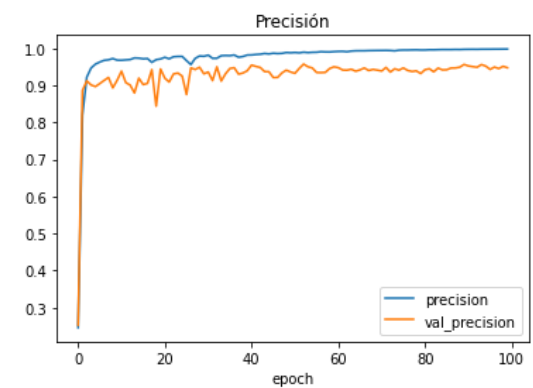
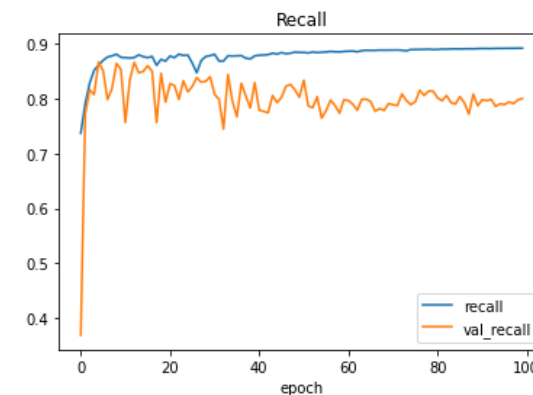
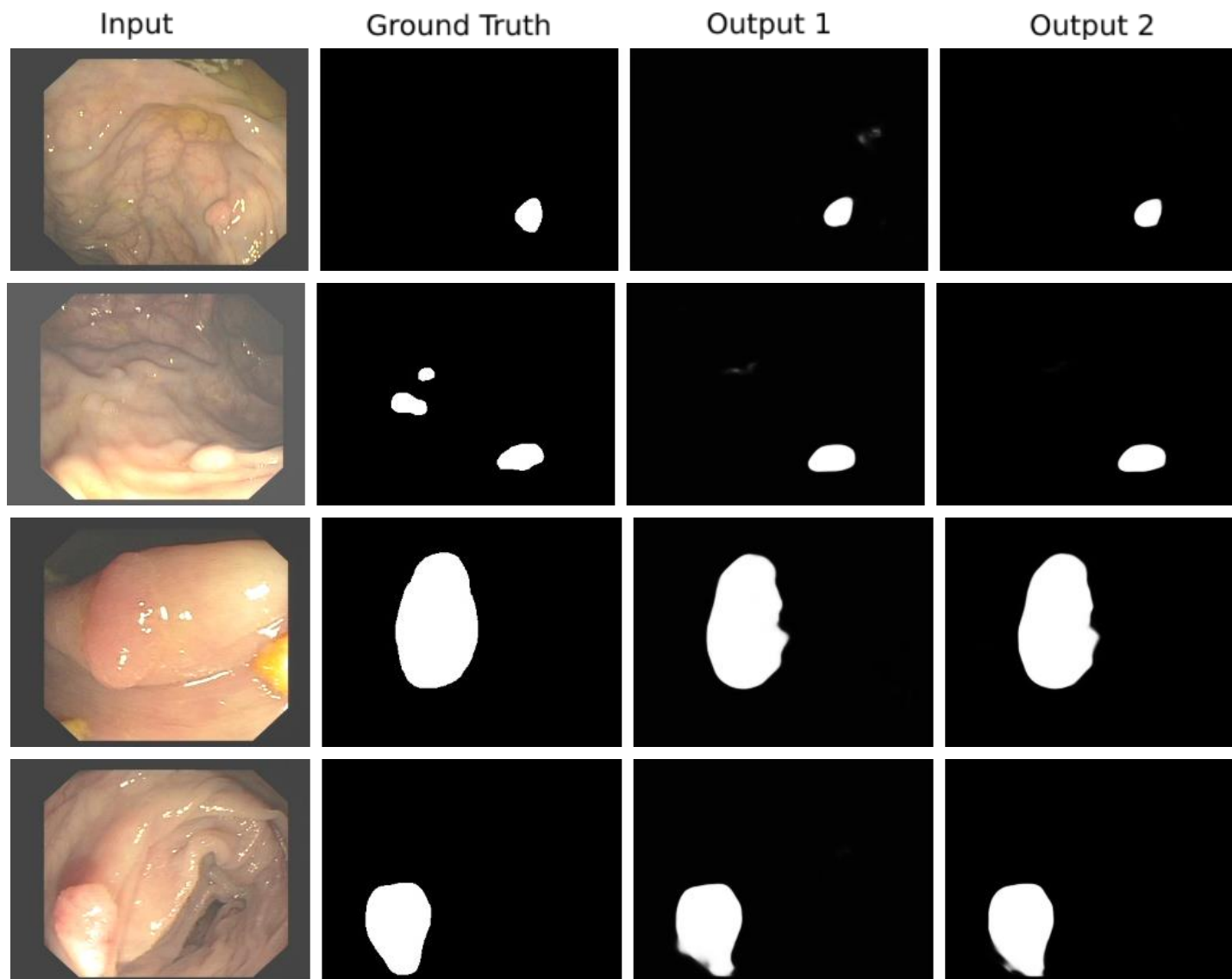
Réplica del modelo DoubleU-Net: Base de datos

- Se utilizó la BD CVC-ClinicDB de colonoscopia.
- Resolución de imágenes de 256x192 de un conjunto base 612:
 - 200 para entrenamiento → 5 000 (25 transformaciones)
 - 61 para validación
 - 61 para prueba



```
from albumentations import (  
    PadIfNeeded,  
    HorizontalFlip,  
    VerticalFlip,  
    CenterCrop,  
    Crop,  
    Compose,  
    Transpose,  
    RandomRotate90,  
    ElasticTransform,  
    GridDistortion,  
    OpticalDistortion,  
    RandomSizedCrop,  
    OneOf,  
    CLAHE,  
    RandomBrightnessContrast,  
    RandomGamma,  
    HueSaturationValue,  
    RGBShift,  
    RandomBrightness,  
    RandomContrast,  
    MotionBlur,  
    MedianBlur,  
    GaussianBlur,  
    GaussNoise,  
    ChannelShuffle,  
    CoarseDropout  
)
```

Desempeño DoubleUnet



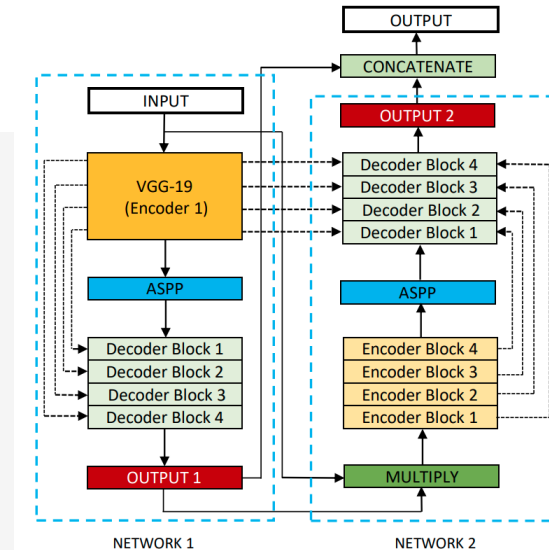
Modificación del modelo

Reducción de filtros en las dos redes que componen la arquitectura *DoubleUnet*

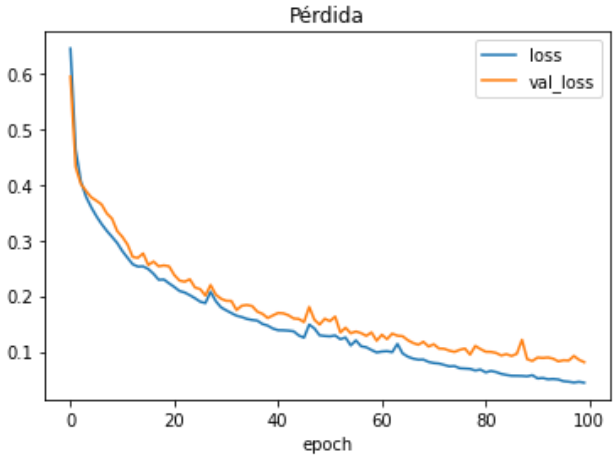
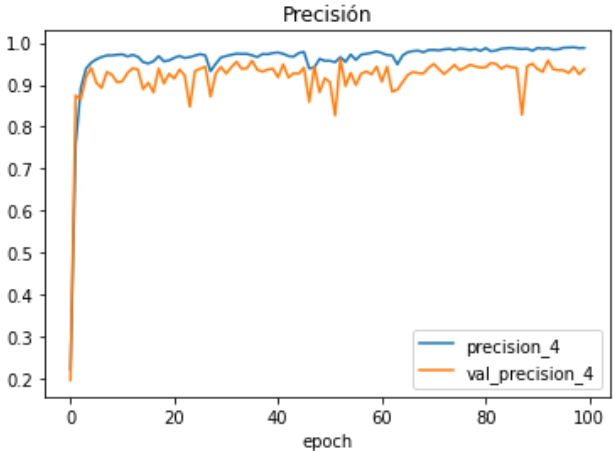
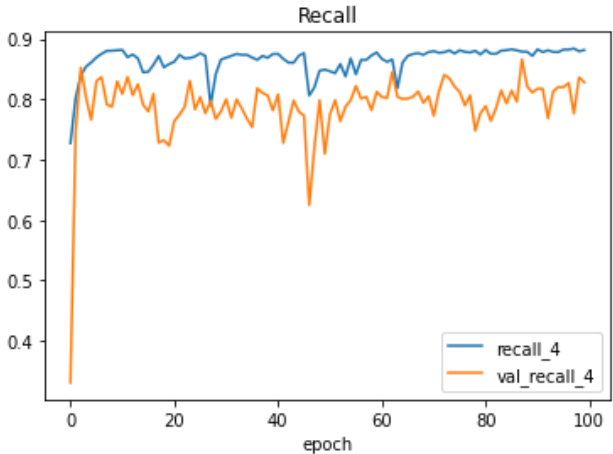
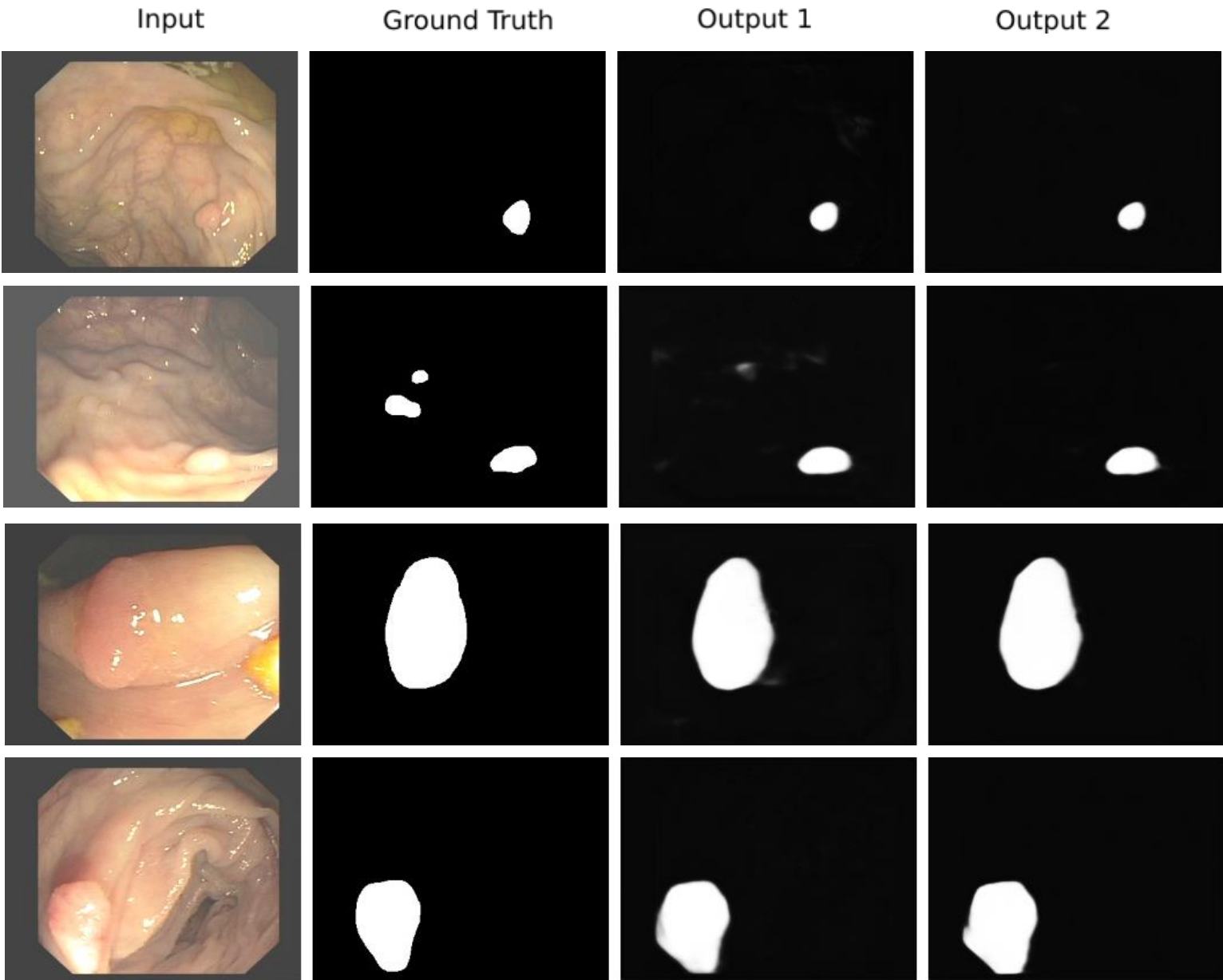
Descripción de la modificación

- Reducción de filtros aplicados por convolución en la etapa de codificación para el modelo UNET.
- Reducción de filtros en la etapa de decodificación para ambos modelos (VGG19, UNET).
- Resultados:
 - Mejora de velocidad de ejecución del modelo.
 - Desempeño parecido al original.

```
def encoder2(inputs):  
    # num_filters = [32, 64, 128, 256]  
    num_filters = [16, 32, 64, 128]  
    skip_connections = []  
    x = inputs  
  
    for i, f in enumerate(num_filters):  
        x = conv_block(x, f)  
        skip_connections.append(x)  
        x = MaxPool2D((2, 2))(x)  
  
    return x, skip_connections  
  
def decoder2(inputs, skip_1, skip_2):  
    # num_filters = [256, 128, 64, 32]  
    num_filters = [128, 64, 32, 16]  
    skip_2.reverse()  
    x = inputs  
  
    for i, f in enumerate(num_filters):  
        x = UpSampling2D((2, 2), interpolation='bilinear')(x)  
        x = Concatenate()([x, skip_1[i], skip_2[i]])  
        x = conv_block(x, f)  
  
    return x
```

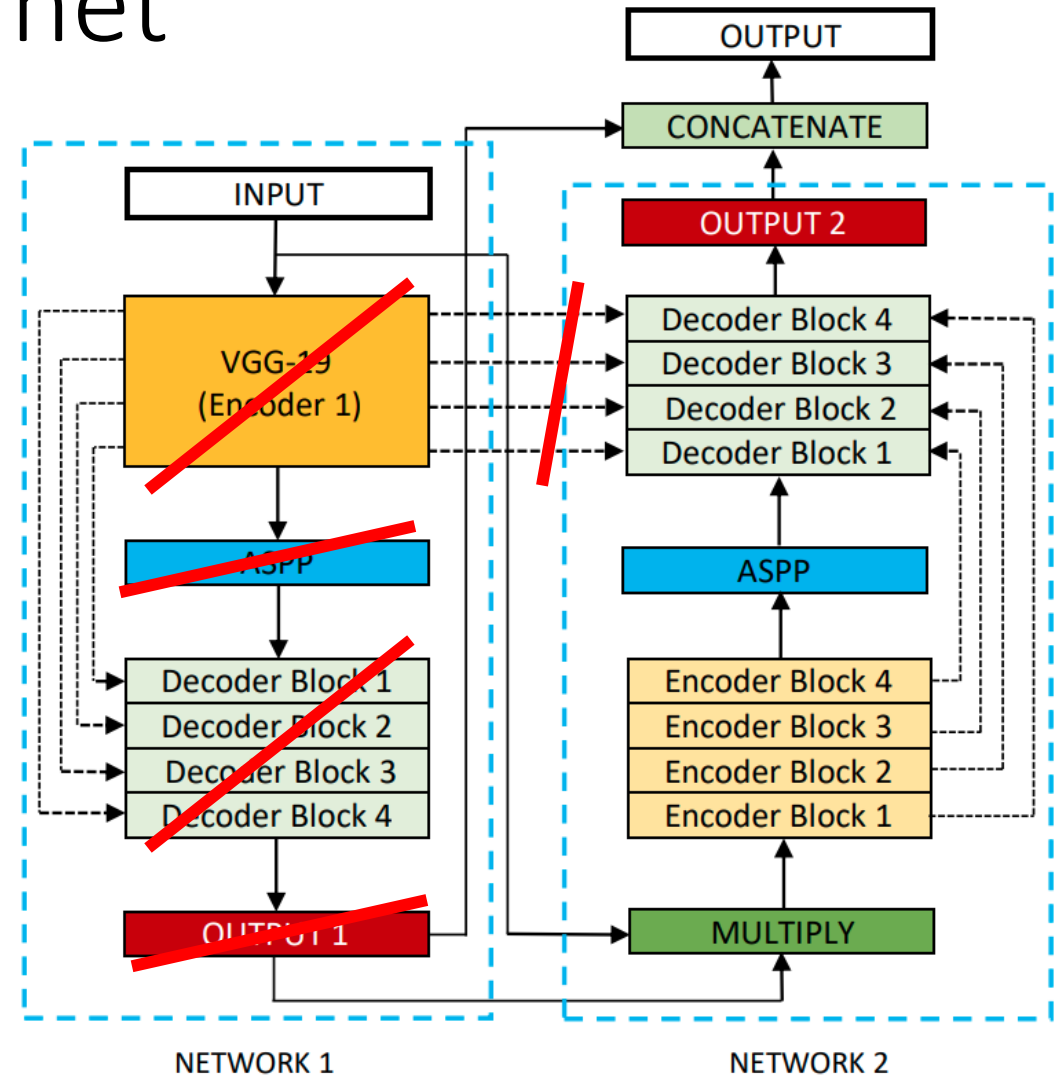


Desempeño DoubleUnet Modificada

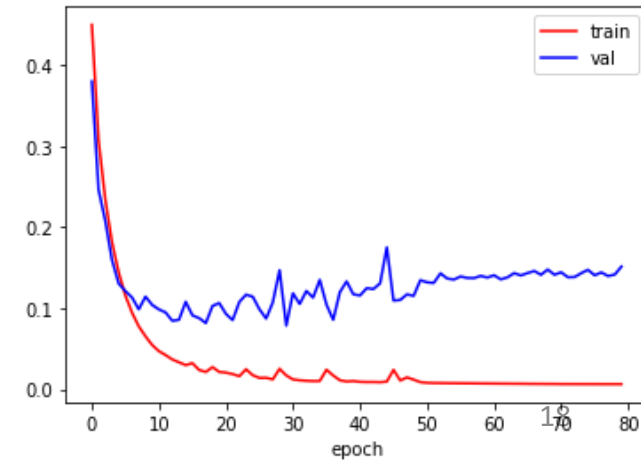
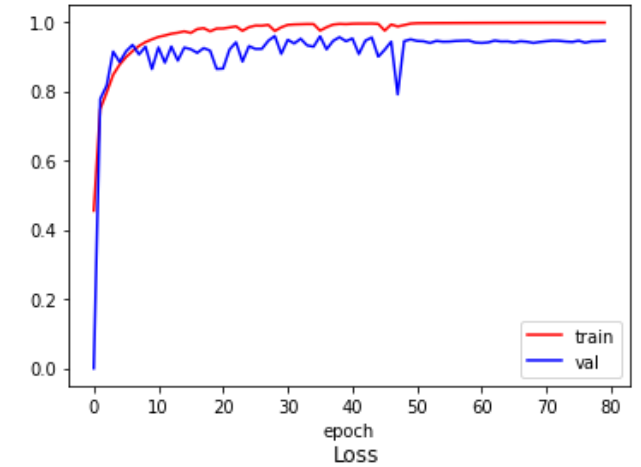
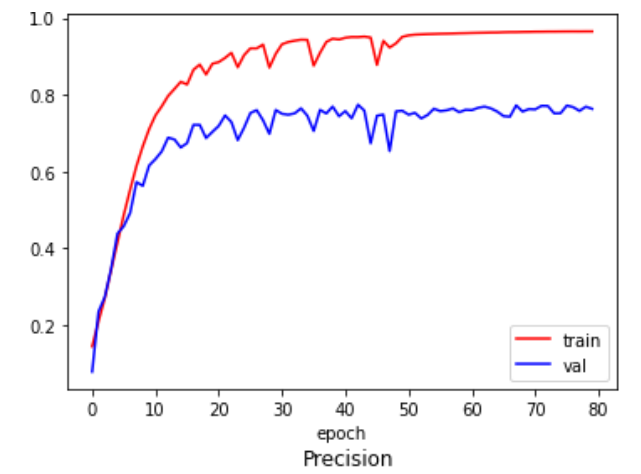
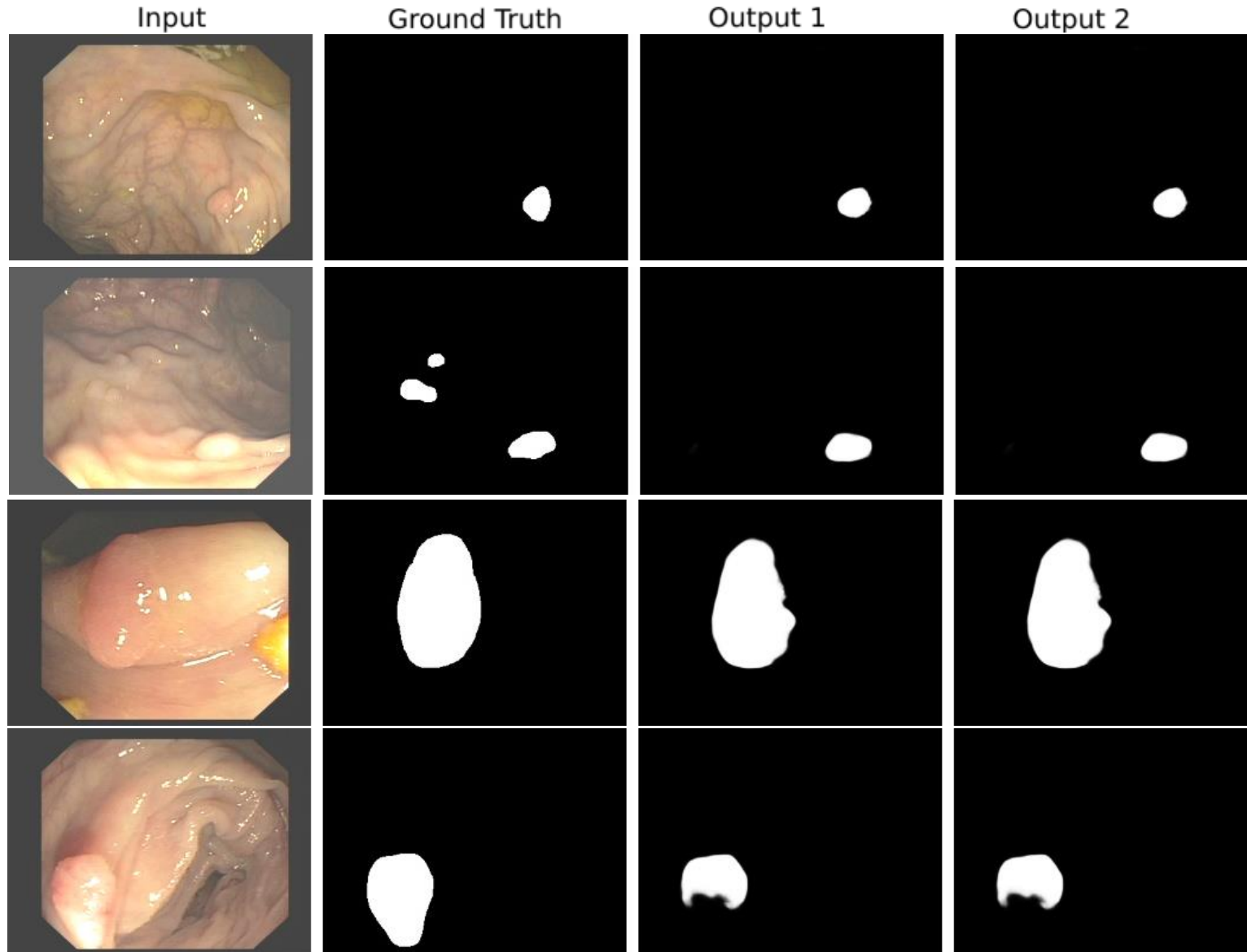


Modelo de referencia Unet

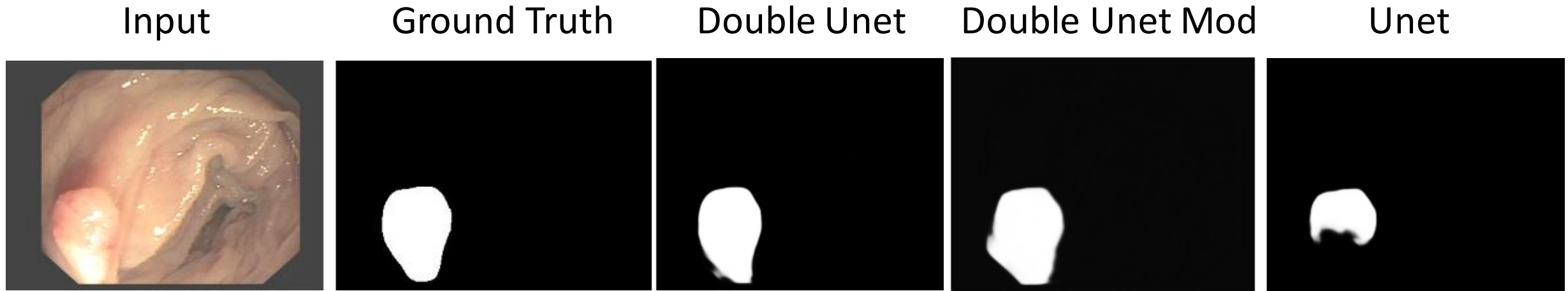
- Para tener una referencia de sobre el desempeño de la arquitectura DoubleUnet modificado, se entreno un modelo Unet basado en el número de filtros aplicados en las etapas de codificación de decodificación del modelo original.



Desempeño modelo Unet de referencia



Discusión: Unet vs Double UNet



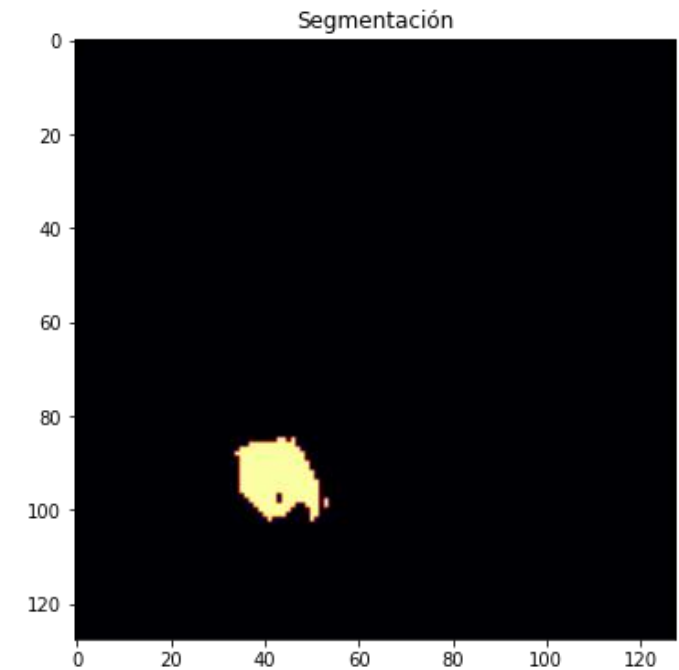
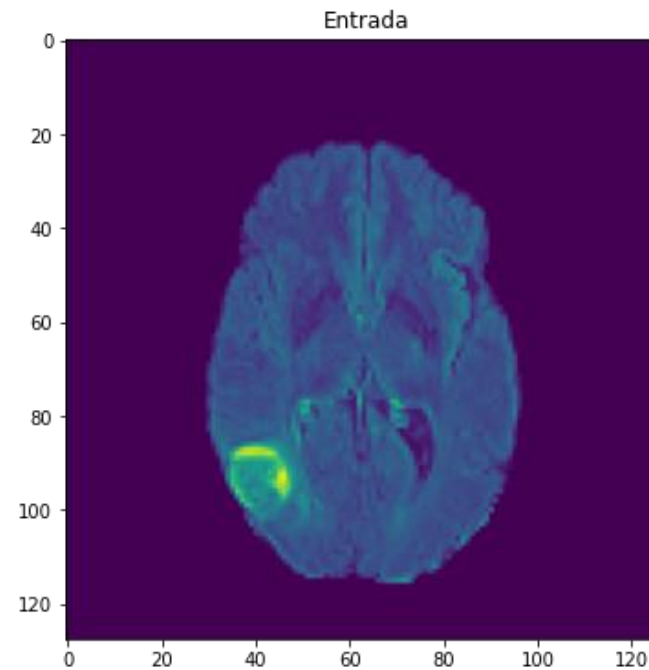
Modelo	loss	precision	val_loss	val_precision
<i>DoubleUnet</i>	0,0113	0,998	0,0826	0,9475
<i>DoubleUnet Mod</i>	0,0443	0,9879	0,0811	0,9376
<i>Unet</i>	0,0364	0,9683	0,1516	0,9461

Segmentación BD BrainTS 2021

Implementación y desempeño de los modelos Unet y DoubleUnet en comparación con el modelo ViT Swin-UNETR (Swin-Unet Transformers) .

BD BrainTS 2021

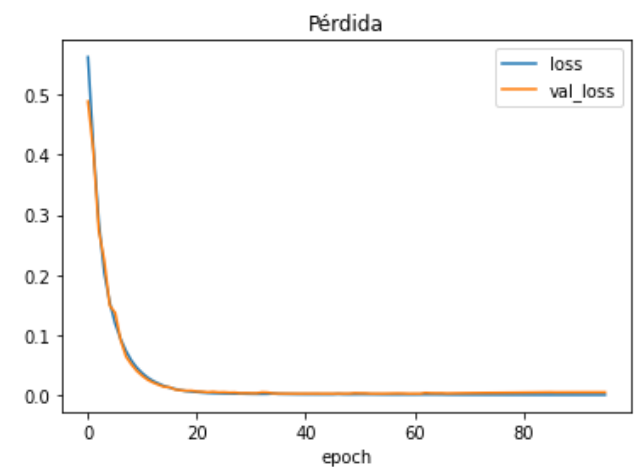
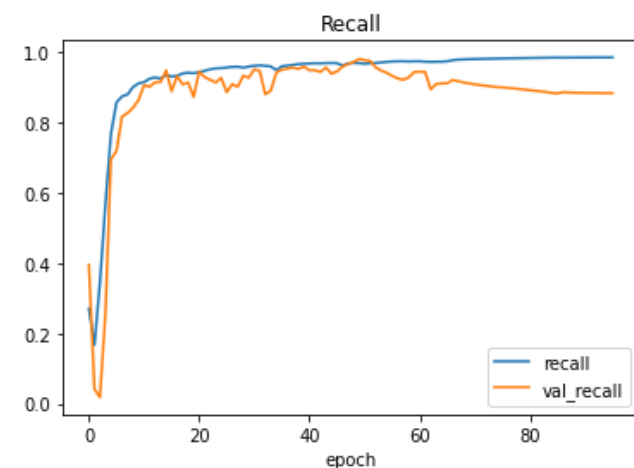
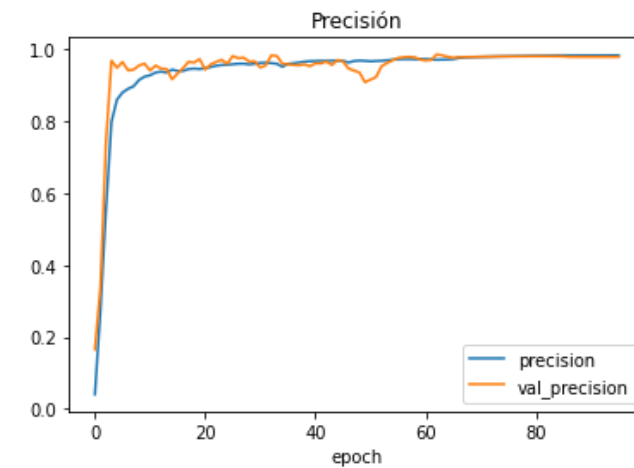
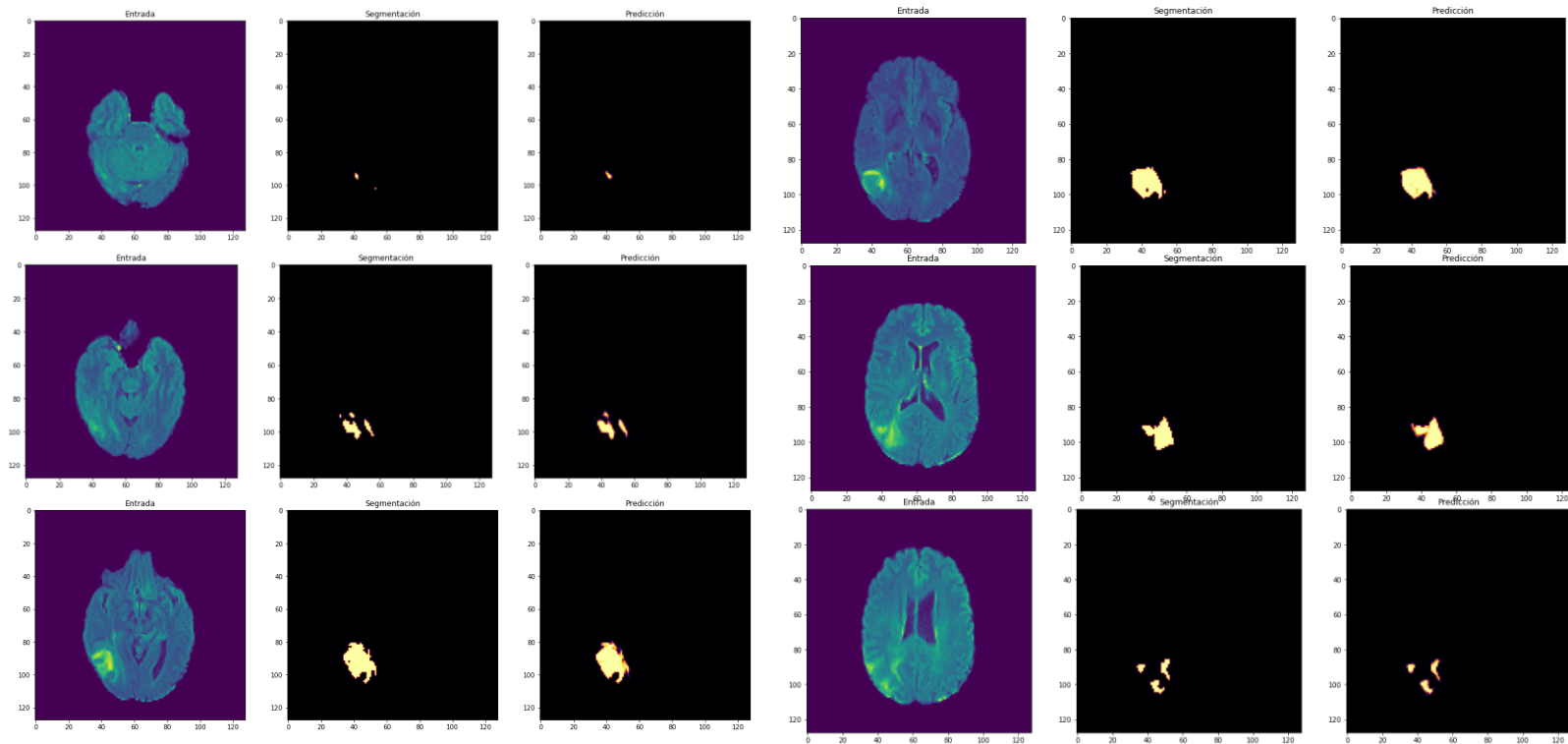
- Se utilizo la BD BraTS 2021 Challenge con
- Resolución de imágenes resolución de 128x128 de un conjunto base 7000:
 - 5 600 para entrenamiento
 - 1 400 para validación
 - 1 para prueba



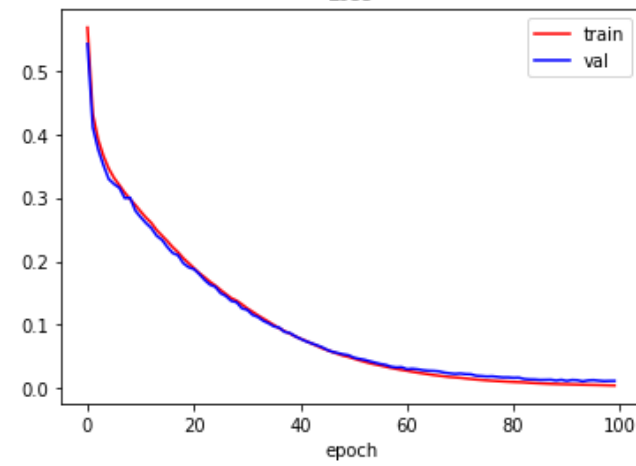
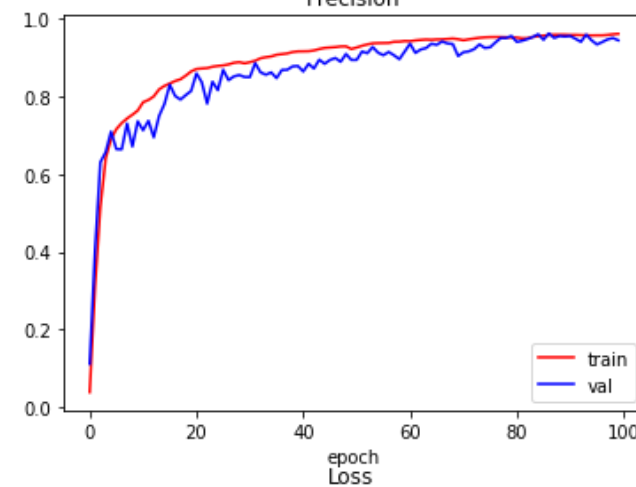
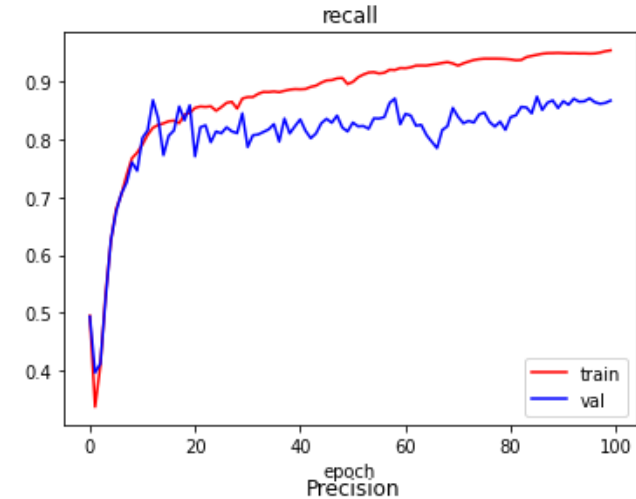
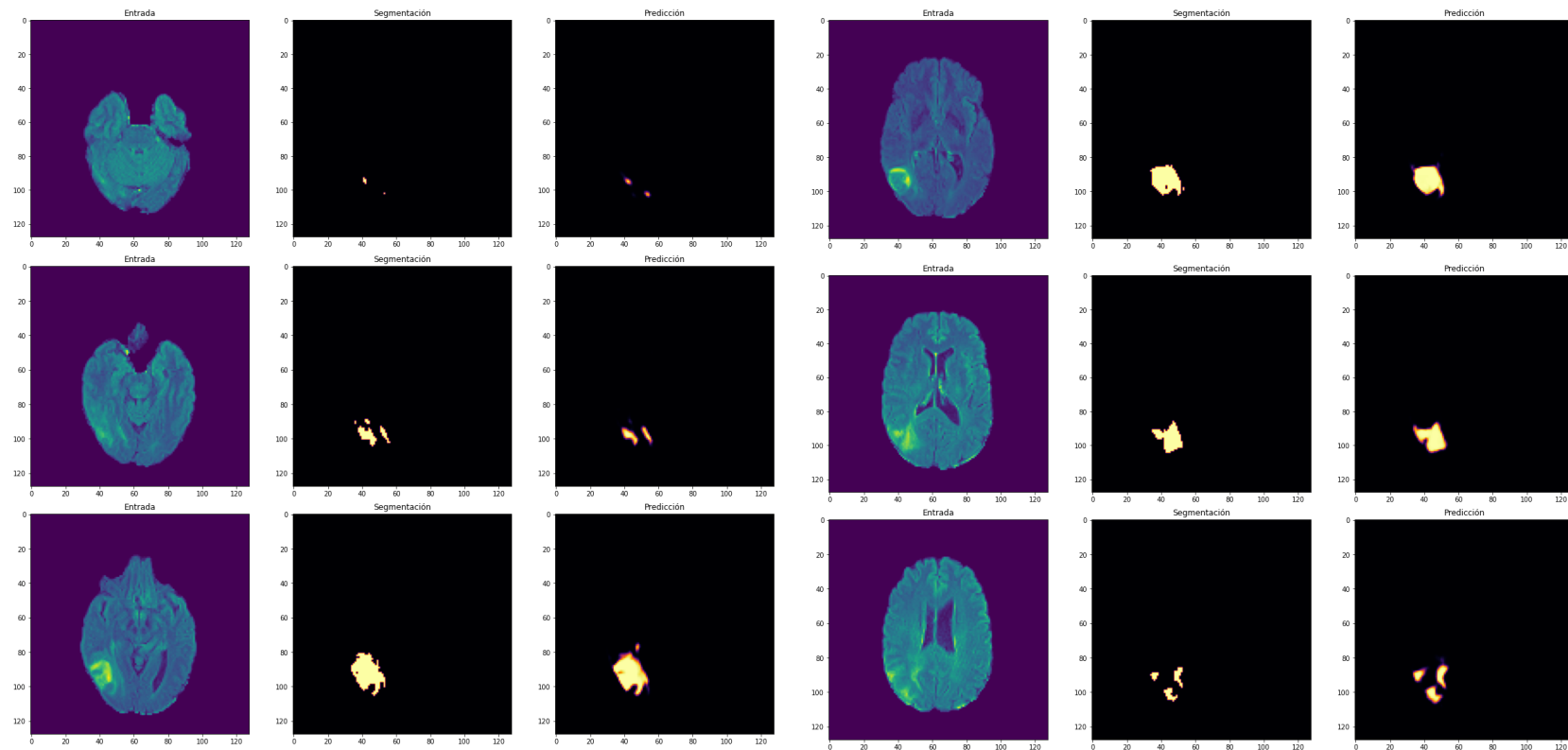
Configuración de los experimentos modelos DoubleUnet y Unet

- Se utilizó la entropía binaria cruzada.
- El optimizador Adam fue empleado.
- El tamaño del batch es 16 y la tasa de aprendizaje de $1e-5$ o $1e-4$.
- Se entrenó el modelo con 100 épocas.

DoubleUNet-Brain

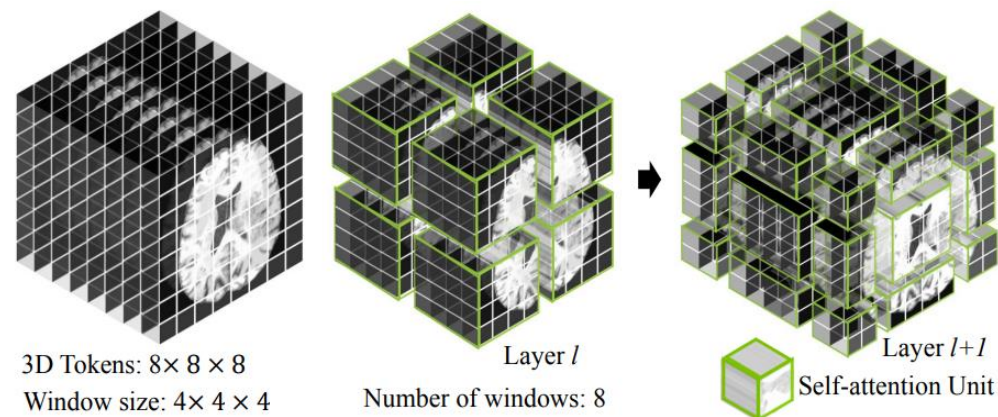
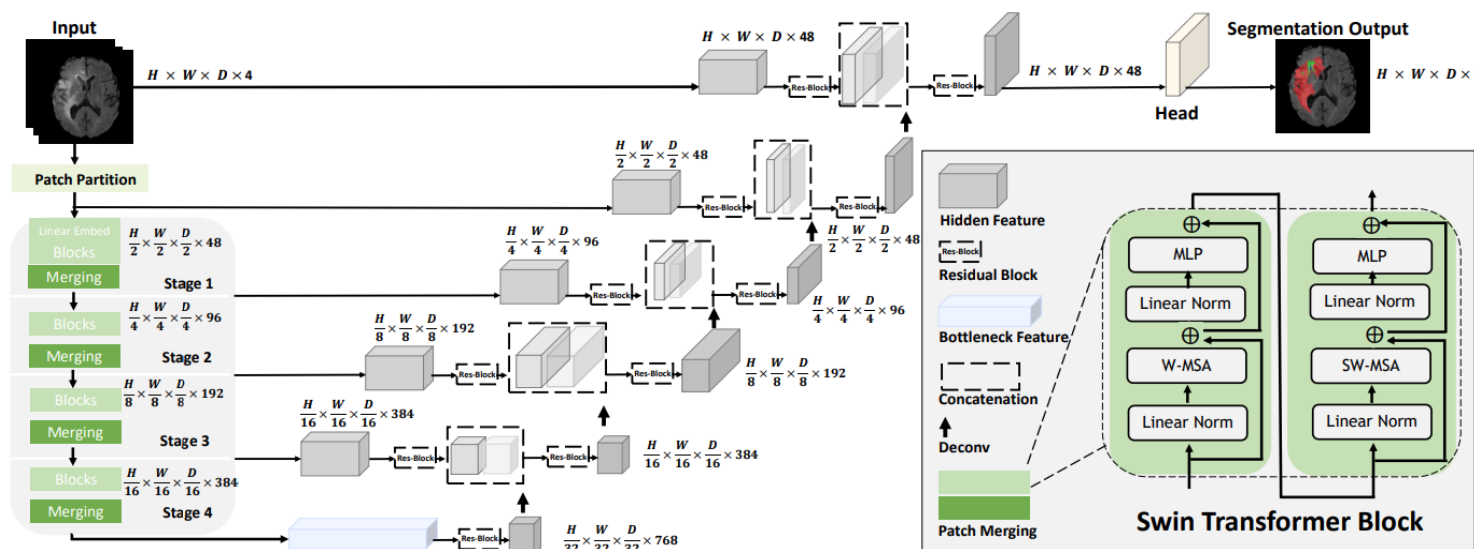


UNet-Brain

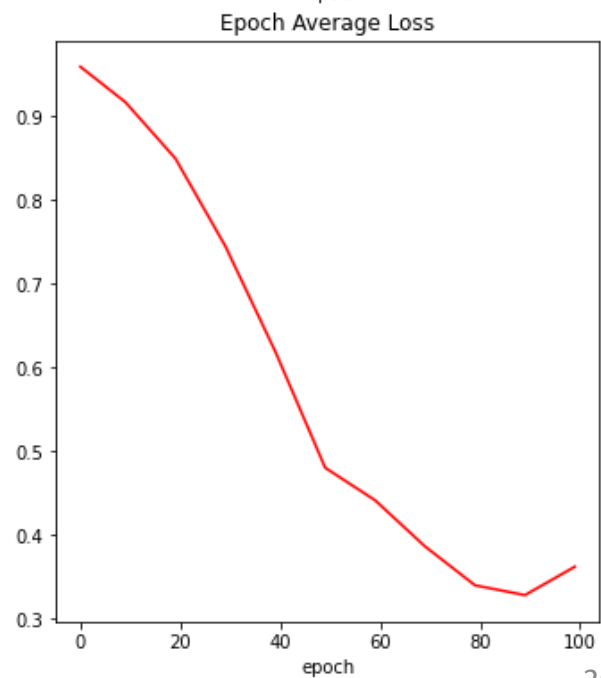
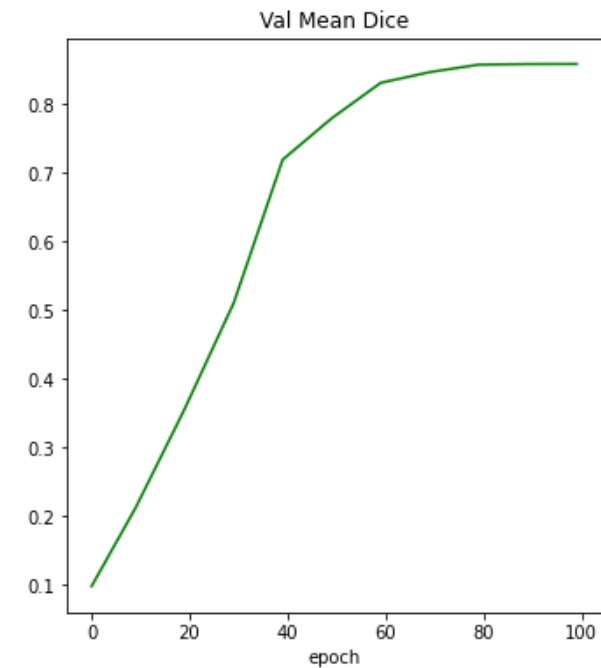
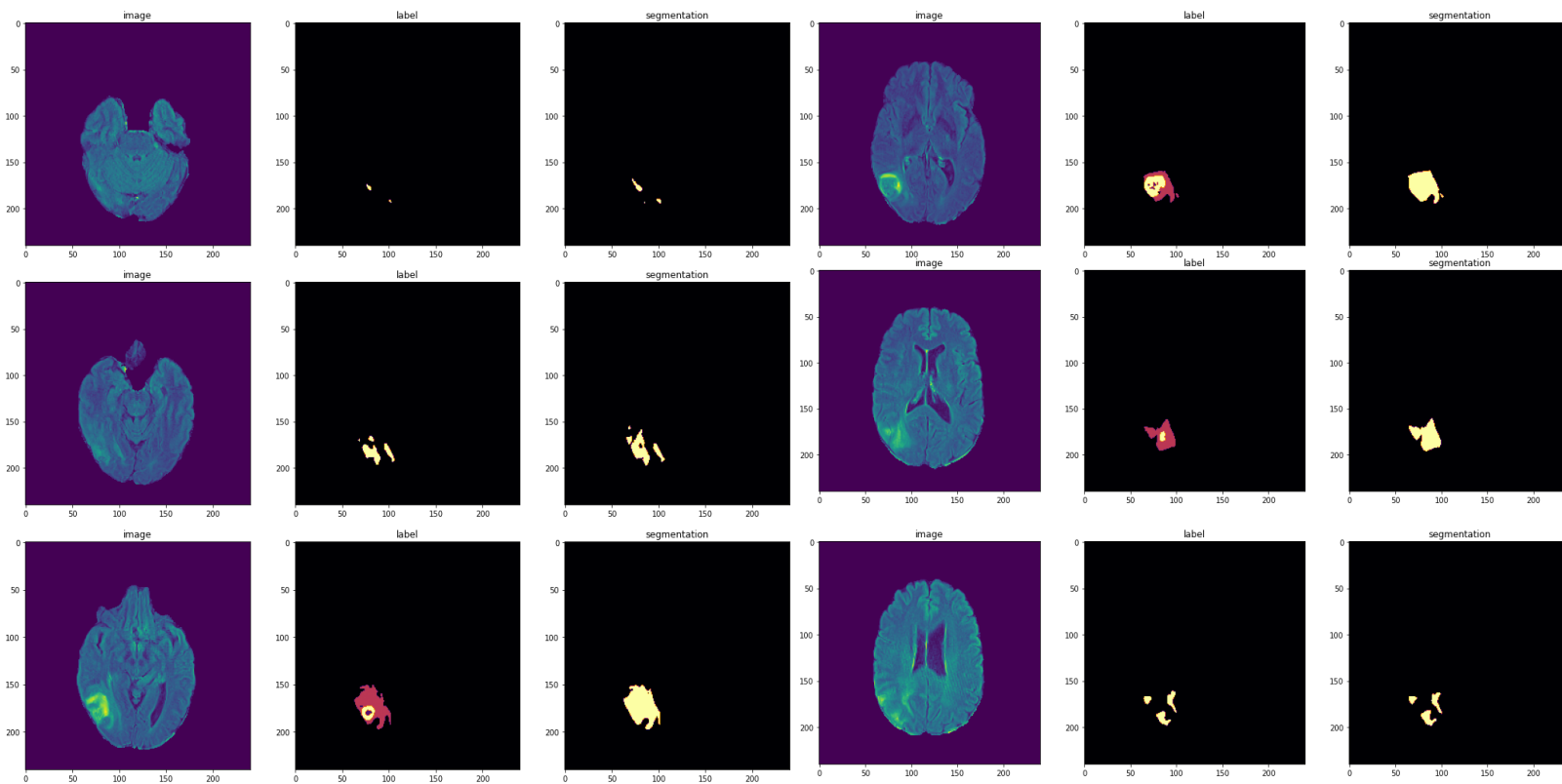


Configuración de los experimentos modelo Swin-UNETR

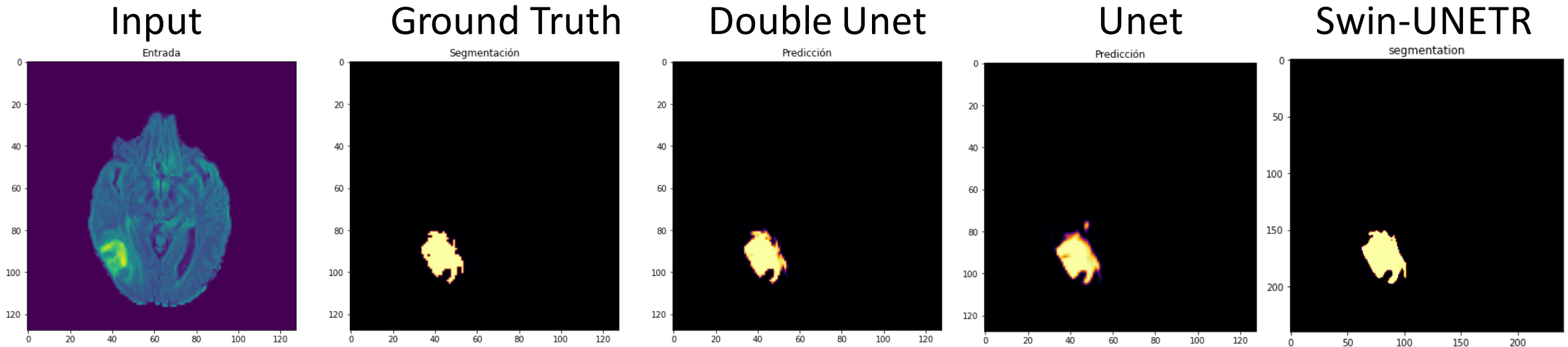
- Se utilizó la entropía binaria cruzada.
- El optimizador AdamW fue empleado.
- El tamaño del batch es 16 y la tasa de aprendizaje de $1e-5$ o $1e-4$.
- Transformaciones
 - RandFlipd
 - NormalizeIntensityd
 - RandScaleIntensityd
 - RandShiftIntensityd
 - ConvertToMultiChannelBasedOnBratsClassesd
 - RandSpatialCropd
 - CropForegroundd
- Se entrenó el modelo con 100 épocas.



Swin-UNET Transformers



Discusión: Unet vs Double Unet VS *Swin UNETR*



Modelo	loss	precision	val_loss	val_precision
<i>DoubleUnet</i>	0.0093	0.9844	0.0052	0.9801
<i>Unet</i>	0,0443	0,9610	0,0115	0,9444
<i>Swin UNETR</i>	0,0364	0,9263	--	0,8583

Referencia

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Referencias

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- [3] N. C. Codella, D. Gutman, M. E. Celebi, B. Helba, M. A. Marchetti, S. W. Dusza, A. Kalloo, K. Liopyris, N. Mishra, H. Kittler et al., “Skin lesion analysis toward melanoma detection: A challenge at the 2017 international symposium on biomedical imaging (isbi), hosted by the international skin imaging collaboration (isic),” in *IEEE International Symposium on Biomedical Imaging (ISBI)*, 2018, pp. 168–172.
- [4] P. Tschandl, C. Rosendahl, and H. Kittler, “The ham10000 dataset, a large collection of multi-source dermatoscopic images of common pigmented skin lesions,” *Scientific data*, vol. 5, p. 180161, 2018. [26] F. Chollet et al., “Keras,” 2015
- [5] F. Chollet et al., “Keras,” 2015.
- [6] M. Abadi, P. Barham, J. Chen, Z. Chen, A. Davis, J. Dean, M. Devin, S. Ghemawat, G. Irving, M. Isard et al., “Tensorflow: A system for large-scale machine learning,” in *Proceeding of {USENIX} Symposium on Operating Systems Design and Implementation ({OSDI})*, 2016, pp. 265–283
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Gracias....