Training report for U-Net (2D) multilabel model (Glio_50E_66im)

Date: 2023-11-11

Training time: 0.0hour(s) 10.0min(s) 35sec(s)

Information for your materials and method:

The U-Net (2D) multilabel model was trained from scratch for 50 epochs on 66 paired image patches (image dimensions: (512, 512), patch size: (512,512)) with a batch size of 4 and acategorical_crossentrop loss function, using the U-Net (2D) multilabel ZeroCostDL4Mic notebook (v 2) (von Chamier & Laine et al., 2020). Key python packages used include tensorflow (v 2.8.0), keras (v 2.8.0), numpy (v 1.23.5), cuda (v 11.8.89 Build cuda_11.8.r11.8/compiler.31833905_0). The training was accelerated using a Tesla T4 GPU.

Augmentation:

The dataset was augmented by

- rotation
- flipping
- random zoom magnification
- shifting
- image shearing

Parameters

Default Advanced Parameters were enabled rameter Value number of epochs 50 512x512 patch_size batch_size 4 number_of_steps 15 10 percentage_validation 0.0003 initial_learning_rate pooling_steps 2 min_fraction 0.02

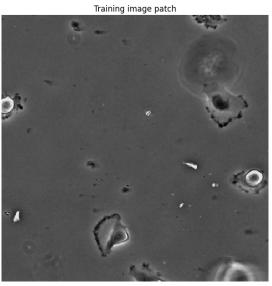
Training Dataset

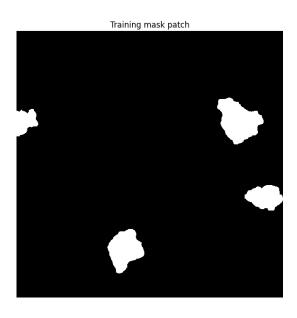
Training_source:/content/gdrive/MyDrive/Colab Notebooks/workshops/datasets/ctc-glioblastoma-66im/train/sources

Training_target: /content/gdrive/MyDrive/Colab Notebooks/workshops/datasets/ctc-glioblastoma-66im/train/masks

Model Path: /content/gdrive/MyDrive/Colab Notebooks/workshops/models//Glio_50E_66im

Example Training pair





References:

- ZeroCostDL4Mic: von Chamier, Lucas & Laine, Romain, et al. "Democratising deep learning for microscopy with ZeroCostDL4Mic." Nature Communications (2021).
- Unet: Ronneberger, Olaf, Philipp Fischer, and Thomas Brox. "U-net: Convolutional networks for biomedical image

segmentation." International Conference on Medical image computing and computer-assisted intervention. Springer, Cham, 2015.

Important:

Remember to perform the quality control step on all newly trained models Please consider depositing your training dataset on Zenodo