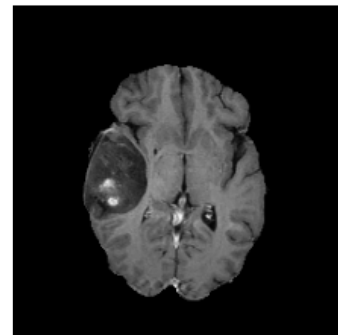
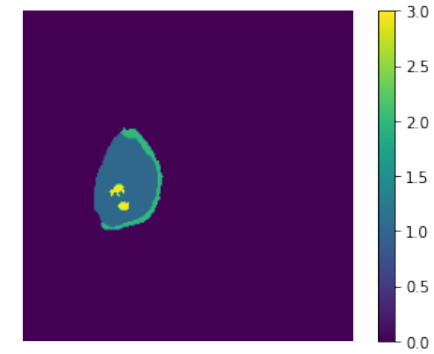
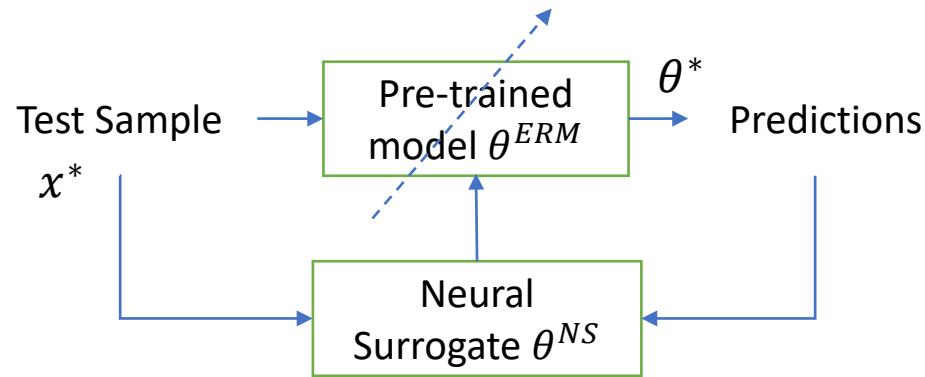


# Test-time adaptation using neural surrogates for medical image segmentation – Hariharan Ravishankar

- Goal : Address generalization of semantic segmentation models by Test-time adaptation of pre-trained model's weights  $\theta^{ERM}$  to  $\theta^*$ , conditioned only on new input sample  $x^*$  in an unsupervised manner.
- Data-set : BRATS 2018 challenge High Grade Glioma (HGG) dataset.



Input 4 channels  
[T1, T1Gd, T2, T2-Flair]



Output 4 channels  
[0- Background, 1 – Core,  
2 – Edema, 3 – EnhT]

- Proposal : Learn  $\theta^{NS}$ - Neural surrogate, which learns to predict quality of segmentation

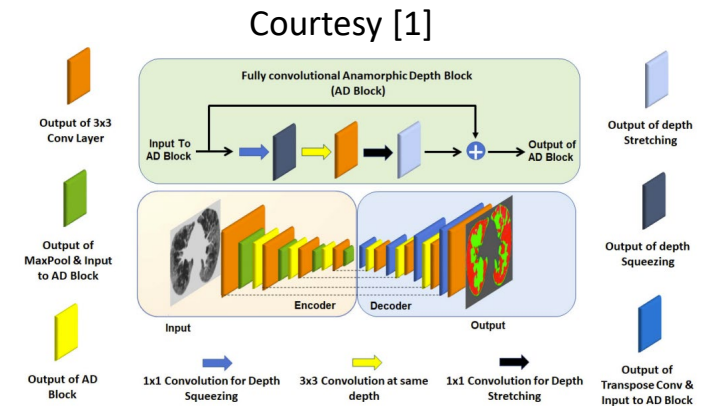
$$g_{\theta^{NS}}(x, y^{\wedge}) \propto l(f_{\theta}(x), y)$$

- During test-time use learnt  $g_{\theta^{NS}}$  for adapting to single sample

# Base Model and Neural Surrogate(s) Details

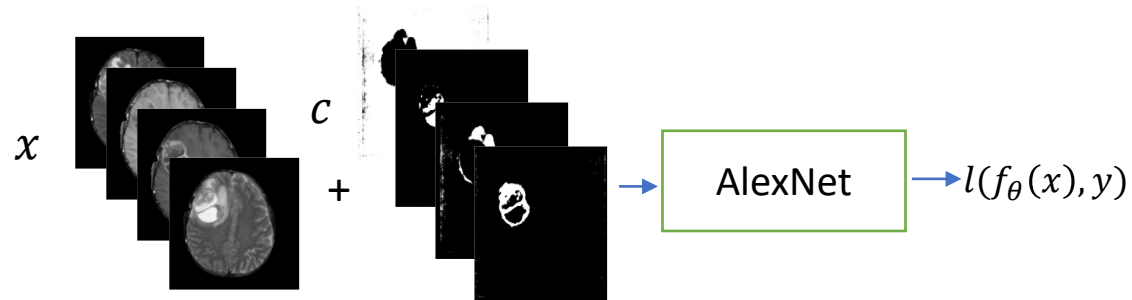
## Base Model Details

- Multi-class segmentation network – AnamNet. Input and output channels = 4.
- Total # of subjects = 210. Training and Validation – 168 subjects, Testing – 42
- Number of slices per subject = 90. Optimizer = Adam, 100 epochs, batch size = 32.



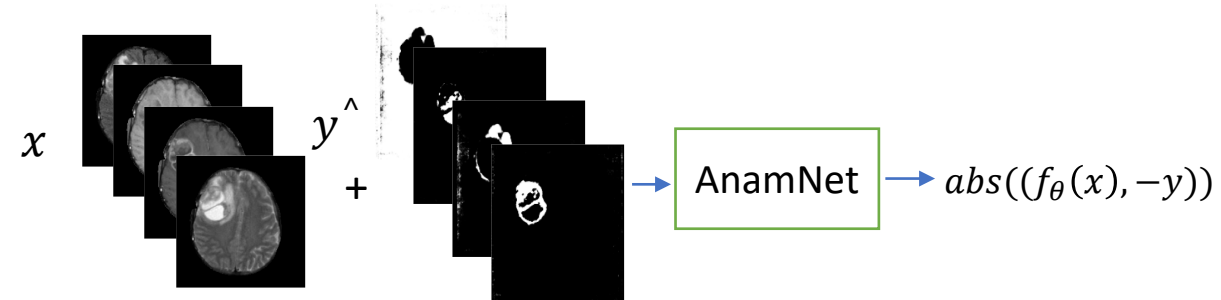
## Neural surrogate 1 – Regression to cumulative loss

- Architecture used : AlexNet
- Input – 8 channels  $\{x, y^\wedge\}$ . Output – single loss value  $l(f_\theta(x), y)$ .
- Hyperparameters same as base model.



## Neural surrogate 2 – Predicting pixel-wise error maps

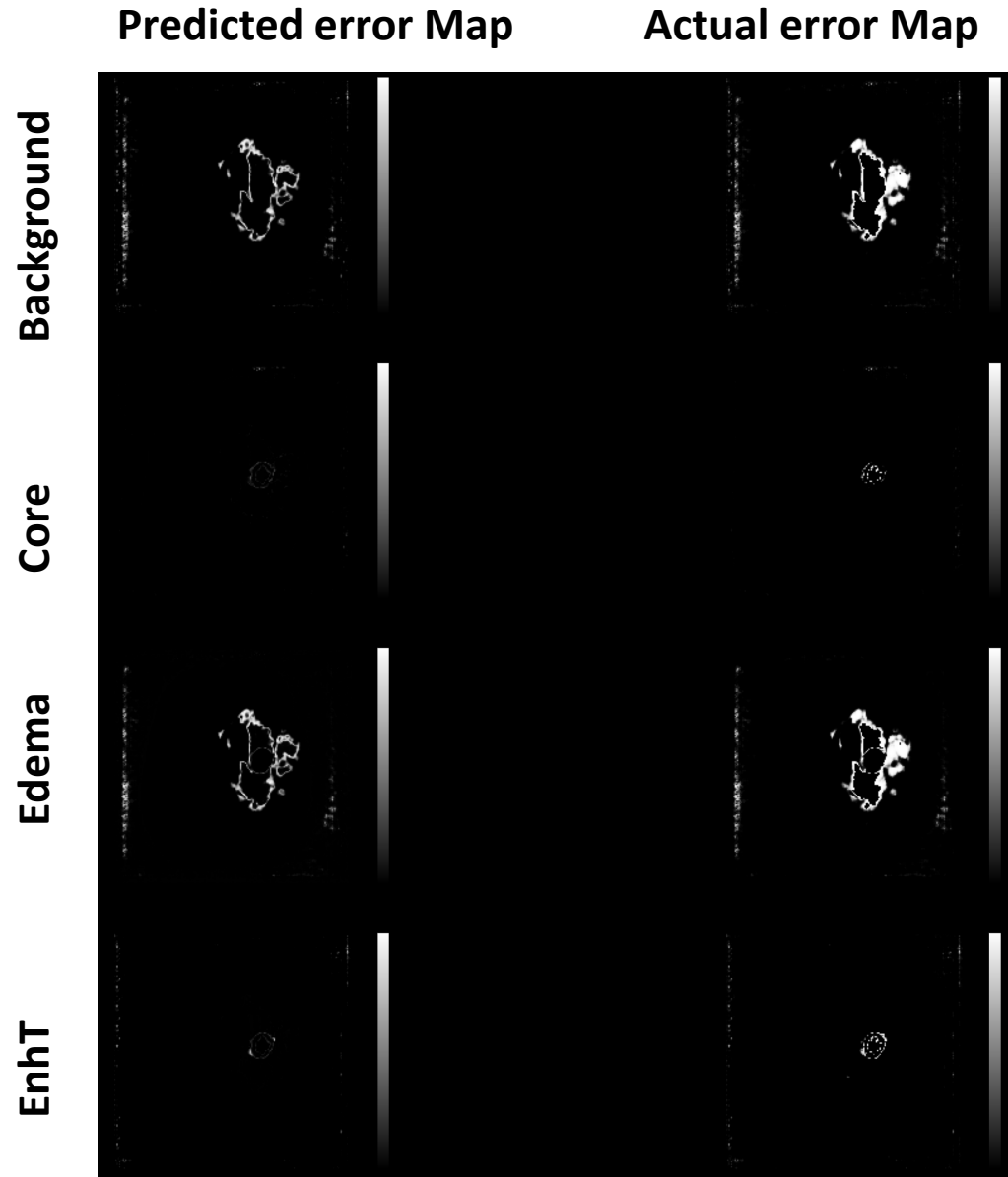
- Architecture used : AlexNet
- Input – 8 channels  $\{x, y^\wedge\}$ . Output – 4 channels -  $abs((f_\theta(x), -y))$ .
- Hyperparameters same as base model.



- Learning was inadequate. Single loss value could not capture pixel-wise mistakes made by base-model.

- NS2 was able to approximate error maps adequately.

# Neural Surrogate 2 performance and test-time adaptation

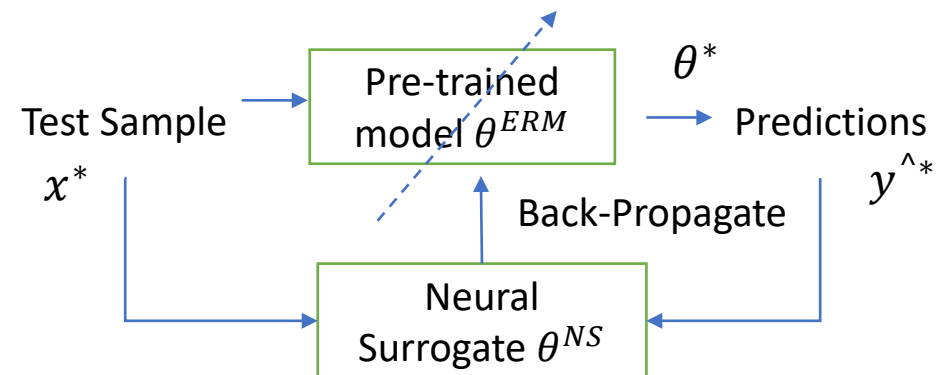


## Neural Surrogate Performance

- $NS2 - g_{\theta^{NS}}(x, \hat{y})$  : was able to capture mistakes made by base Model.
- Note : Overlapping information in background and one of the classes.
- Criterion used : MSE. Training error  $\sim 0.0010$

## Iterative procedure for test-time adaptation of using NS2

- For batch of samples from test-data, initial predictions were made using pre-trained model -  $\theta^{ERM}$
- The output predictions  $\hat{y}^*$  and inputs  $x^*$  were passed to  $g_{\theta^{NS}}$ .
- Output of  $g_{\theta^{NS}}$  was used to backpropagate and update weights.
- Only batch norm-parameters were updated using this procedure.



# Quantitative results with Test-time adaptation using NS2

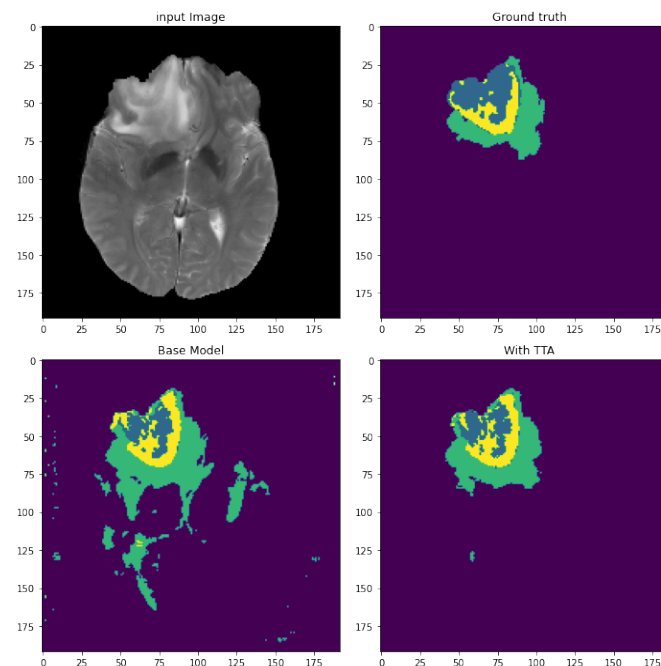
Dice overlap scores on test data-set for different components of tumor and for varying training data sizes

Class / Training Data Fraction	0.25		0.5		1.0	
	Orig Model	With TTA	Orig Model	With TTA	Orig Model	With TTA
Core	0.442	<b>0.527</b>	0.535	<b>0.655</b>	0.69	<b>0.692</b>
Edema	0.650	<b>0.660</b>	0.714	<b>0.724</b>	0.718	<b>0.765</b>
EnhT	0.429	<b>0.692</b>	0.800	<b>0.800</b>	0.780	<b>0.82</b>
Whole Tumor	0.664	<b>0.762</b>	0.799	<b>0.834</b>	0.818	<b>0.864</b>

## Key Take-aways

- TTA with NS2 consistently improves performance on all types of tumor tissues with roughly 5% increase on full training dataset.
- TTA with 50% data beats orig model's performance on full data.
- Increase in performance due to TTA is possible for all training sizes, including 25% of training data with close to 10% increase in dice overlap.

Qualitative visualization of improvements in segmentation results with TTA using neural surrogates



## Other Details

- Resources : MIG Lab server and workstation with RTX A600 gpu cards.
- Orig Model inference – 10ms, Adapt procedure – 1s for batch size of 24.
- Experiments with more architecture and additional problems in progress.