

#### Optimizing U-Net Architecture Using Differential Evolution for Brain Tumor Segmentation

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# Why Brain Tumor Segmentation?

- Accurate segmentation critical for diagnosis and treatment
- Manual segmentation is time-consuming and inconsistent
- Need for robust automatic models that generalize across MRI modalities



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# **Key Contributions**

- Proposed DE-UNet: U-Net optimized using Differential Evolution (DE)
- Automatically tunes hyperparameters: learning rate, dropout, batch size, filter size
- Outperforms state-of-the-art models on FBTS and BraTS 2021 datasets



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### **DE-UNet Framework**



 Base model: U-Net with skip connections
 DE tunes 4 key hyperparameters

• Optimization minimizes composite loss:  $L_{\text{total}} = 1 - \alpha \cdot \text{DSC} - (1 - \alpha) \cdot \text{JI}$ 



# Differential Evolution Strategy

- Population-based metaheuristic
- Mutation, crossover, and selection
- 10 generations
  with population
  size = 5

Algorithm 1 DE Algorithm for U-Net hyperparameter optimization.

Require: NP, F, CR, G

**Ensure:** Optimized hyperparameter configuration  $x^*$ 

- 1: Initialize a population of NP candidate solutions within predefined bounds.
- 2: for generation g = 1 to G do
- 3: for each candidate solution  $x_i$  in the population do
- 4: **Mutation:** Randomly select 3 distinct solutions  $(x_a, x_b, x_c)$  from the population.
- 5: Generate mutant vector:  $v_i = x_a + F \cdot (x_b x_c)$
- 6: **Crossover:** Generate trial vector  $u_i$  as:

 $u_{ij} = \begin{cases} v_{ij} & \text{if rand}(0,1) < CR \text{ or } j = j_{\text{rand}} \\ x_{ij} & \text{otherwise} \end{cases}$ 

- 7: Selection: Replace  $x_i$  with  $u_i$  if  $f(u_i) < f(x_i)$
- 8: end for
- 9: end for
- 10: Return the best-performing solution  $x^*$



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## **Datasets and Preprocessing**

- FBTS: 3064 slices, 3 tumor types (Meningioma, Glioma, Pituitary)
- BraTS 2021: 1251 slices, 4 modalities (T1, T1-CE, T2, FLAIR)
- Resized to 256×256, normalized to [0,1]







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# Performance Summary

Tumor Type	Training				Validation			
	Accuracy	Loss	DSC	JI	Accuracy	Loss	$\operatorname{DSC}$	JI
Meningioma	0.9983	0.0042	0.9286	0.8677	0.9984	0.0038	0.9348	0.8784
Glioma	0.9971	0.0070	0.9023	0.8231	0.9968	0.0080	0.8943	0.8103
Pituitary	0.9991	0.0022	0.9183	0.8509	0.9991	0.0021	0.9200	0.8539

Modality-	Training				Validation				
	Accuracy (pp)	Loss (pp)	DSC (pp)	JI (pp)	Accuracy (pp)	Loss (pp)	DSC (pp)	JI (pp)	
FLAIR	0.9956 (+0.18)	0.0110 (-0.47)	0.8941 (+4.31)	0.8103(+7.21)	0.9961 (+0.15)	0.0095 (-0.37)	$0.9068 \ (+3.31)$	0.8304 (+5.69)	
T1	0.9935 (+0.32)	0.0157 (-0.79)	0.8464 (+7.58)	0.7353(+12.12)	0.9930 (+0.39)	0.0168 (-0.96)	$0.8327 \ (+9.07)$	0.7154 (+14.3)	
T1-CE	0.9950 (+0.34)	0.0119 (-0.82)	0.8823 (+8.05)	0.7900(+13.86)	0.9940 (+0.43)	0.0147 (-1.06)	0.8576 (+10.37)	0.7524(+17.34)	
T2	0.9946 (+0.28)	0.0135(-0.72)	$0.8707 \ (+6.65)$	0.7714(+11.1)	$0.9942 \ (+0.34)$	0.0147 (-0.89)	0.8602 (+7.97)	0.7550(+13.23)	

Note: "+" indicate performance improvement in pp. "-" loss values represent reduction.

- FBTS: DSC = 0.9160, JI = 0.8472
- BraTS 2021: DSC = 0.9094, JI = 0.8371
- Consistent gains across all tumor types and modalities



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### **Qualitative Segmentation Examples**

#### FBTS





 $\begin{array}{ccccc} Meningioma & Glioma & Pituitary \\ DSC: 0.9368 & DSC: 0.9582 & DSC: 0.9567 \\ JI: 0.8811 & JI: 0.9198 & JI: 0.9169 \end{array}$ 

#### BraTS 2021







FLAIRT1T1-CET2DSC: 0.9694DSC: 0.9556DSC: 0.9699DSC: 0.9762JI: 0.9405JI: 0.9150JI: 0.9416JI: 0.9536

### Red = Ground Truth, Green = Prediction

Strong boundary alignment, especially in T1-CE and FLAIR



#### DE-UNet vs. State-of-the-Art

Method	FBTS Dataset Method			BraTS 2021		
Method	DSC	JI	Wethod	$\operatorname{DSC}$	JI	
Proposed DE-UNet	0.9160	0.8472	Proposed DE-UNet	0.9094	0.8371	
${\it DeepLabV3+Xception}$	0.8115	0.8018	UNet	0.8600	0.7807	
KFCM-CNN	0.8884	0.8204	U-Net base	0.9080	-	
U-Net based	0.8900	0.8100	SPPNet-2	0.9040	-	
MST-based	0.8469	0.7443	UNCE-NODE	0.8949	-	
U-Net with ResNet	0.9011	-	nnU-Net	0.8900	-	

- DE-UNet outperforms models like U-Net, DeepLabV3+, UNCE-NODE
- Higher DSC and JI without manual tuning



## Conclusion & Future Directions

- DE-UNet provides accurate, robust brain tumor segmentation
- Effective hyperparameter tuning across modalities
- Future work: hybrid metaheuristics, more datasets, clinical deployment