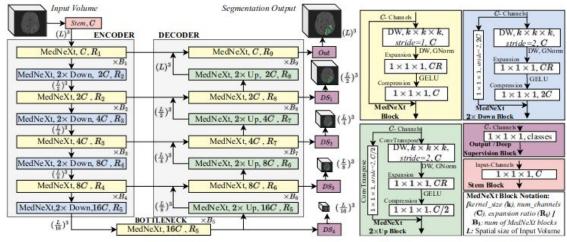




Segmentation task: MedNeXt Model presentation



- Keeps the inductive bias of the convolutional layers
- Compares favorably with SOTA architectures, including transformers, when trained from scratch
- Is de facto integrated in the nnUnet segmentation pipeline



[1] F. Isensee, et al. "nnu-net revisited: A call for rigorous validation in 3d medical image segmentation." *arXiv preprint arXiv:2404.09556* (2024).

[2] S. Roy, et al. "Mednext: transformer-driven scaling of convnets for medical image segmentation." *International Conference on Medical Image Computing and Computer-Assisted Intervention.* Cham: Springer Nature Switzerland, 2023.

2×0 p 100	CK.	•		L	Span	idi Siza	01 11	pui	VOLU	HIRC
	BTCV n=30	ACDC n=200		BraTS n=1251		AMOS n=360		RT [h]	Arch.	nnU
nnU-Net (org.)	83.08	91.54	80.09	91.24	86.04	88.64	7.70	9	CNN	Yes
nnU-Net ResEnc M	83.31	91.99	80.75	91.26	86.79	88.77	9.10	12	CNN	Yes
nnU-Net ResEnc L	83.35	91.69	81.60	91.13	88.17	89.41	22.70	35	CNN	Yes
nnU-Net ResEnc XL	83.28	91.48	81.19	91.18	88.67	89.68	36.60	66	CNN	Yes
MedNeXt L k3 33	84.70	92.65	82.14	91.35	88.25	89.62	17.30	68	CNN	Yes
MedNeXt L k5 34	85.04	92.62	82.34	91.50	87.74	89.73	18.00	233	CNN	Yes
STU-Net S [20]	82.92	91.04	78.50	90.55	84.93	88.08	5.20	10	CNN	Yes
STU-Net B 20	83.05	91.30	79.19	90.85	86.32	88.46	8.80	15	CNN	Yes
STU-Net L 20	83.36	91.31	80.31	91.26	85.84	89.34	26.50	51	CNN	Yes
SwinUNETR 32	78.89	91.29	76.50	90.68	81.27	83.81	13.10	15	TF	Yes
SwinUNETRV2	80.85	92.01	77.85	90.74	84.14	86.24	13.40	15	TF	Yes
nnFormer [43]	80.86	92.40	77.40	90.22	75.85	81.55	5.70	8	TF	Yes
CoTr 💝	81.95	90.56	79.10	90.73	84.59	88.02	8.20	18	TF	Yes
No-Mamba Base	83.69	91.89	80.57	91.26	85.98	89.04	12.0	24	CNN	Yes
U-Mamba Bot 26	83.51	91.79	80.40	91.26	86.22	89.13	12.40	24	Mam	Yes
U-Mamba Enc [20]	82.41	91.22	80.27	90.91	86.34	88.38	24.90	47	Mam	Yes
A3DS SegResNet 3,28	80.69	90.69	79.28	90.79	81.11	87.27	20.00	22	CNN	No
A3DS DINTS DINS	78.18	82.97	69.05	87.75	65.28	82.35	29.20	16	CNN	No
A3DS SwinUNETR A3DS	76.54	82.68	68.59	89.90	52.82	85.05	34.50	9	TF	No

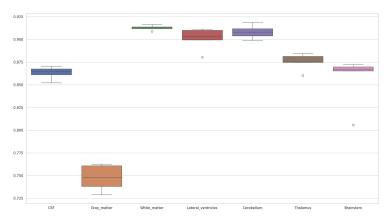




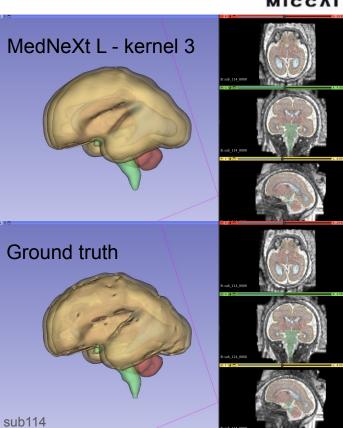
Segmentation task: Results



- Predicts smoother structures than the expert annotations
- Compares on pare with previous FeTA submissions
- Has been evaluated by radiologist experts on a private dataset: 45 reconstructed foetus MRI (1.5T)



[3] K. Payette, et al. "Multi-Center Fetal Brain Tissue Annotation (FeTA) Challenge 2022 Results." *arXiv preprint arXiv:2402.09463* (2024).







Biometry task: Rule based strategy



- Measures correlate well with segmentations bounding boxes
- Estimation of the rotation matrix would certainly improve the results

Measure	ME (%)	R2		
sBIP	5.703417	0.865769		
bBIP	7.969022	0.754252		
HV	39.190029	-2.12880		
TCD	4.48192	0.94305		
LCC	19.55445	-0.73959		

