

27<sup>TH</sup> INTERNATIONAL CONFERENCE ON MEDICAL IMAGE COMPUTING  
AND COMPUTER ASSISTED INTERVENTION  
6-10 OCTOBER 2024  
PALMERAIE ROTANA RESORT  
MARRAKESH / MOROCCO

## Fetal Tissue Annotation and Segmentation Challenge (FeTA) Overview and results

October 6<sup>th</sup> 2024

@FetaChallenge

<https://fetachallenge.github.io>





# AGENDA



**11:00 - 11:15** FeTA Challenge overview – M. Bach Cuadra

**11:15 - 11:50** Power pitch presentations by top 5 teams:

- **Abdul Qayyum:** “Hybrid Cross Attention Transformer and CNN model for Feta Segmentation Task” (2 min)
- **Maximilian Zenk:** “Exploring the out-of-distribution generalization of nnU-Net for fetal brain tissue segmentation” (2 min)
- Rachika E. Hamadache: “NIC-VICOROB Team: Fetal Tissue Annotation and Segmentation challenge (FeTA 2024)”

Q/A

- Robin Cremese: “Mednext adoption for fetal brain MRI segmentation”
- Tong Lyuyang: “Team FeTA\_Sigma: Prenatal Brain Segmentation and Biometry Algorithm Overview” (4 min)
- Denis Peruzzo: “CeSNE DIGAIR team”
- Param Ahir: power-pitch available at FeTA webpage

Q/A

**11:50 - 12:00** FeTA winners ceremony – K. Payette

**12:00 - 13:30** Joint PIPPI & FeTA Poster Session



# ORGANIZERS



M. Bach Cuadra



K. Payette



A. Jakab



R. Licandro



M. Barkovich



H. B. Li



J. Hutter



T. Sanchez



V. Zalevskyi



M. Roulet



M. Kaandorp



L. Li



D. Fajardo-Rojas



Centre hospitalier  
universitaire vaudois

UCSF Benioff Children's Hospitals



HARVARD  
MEDICAL SCHOOL



Imperial College  
London



MEDICAL UNIVERSITY  
OF VIENNA



Uniklinikum  
Erlangen



UNIL | Université de Lausanne



University of  
Zurich

# OVERVIEW

## Motivation

- In support to clinical risk stratification tools for early interventions, treatments, and clinical counselling.
- In vivo portray of complex neurodevelopmental events during human gestation

## Challenges

- Data scarcity
- Data access
- Domain shifts

Acquisition, scanner, etc  
Pre-processing  
Pathology  
Gestational age  
SR methods

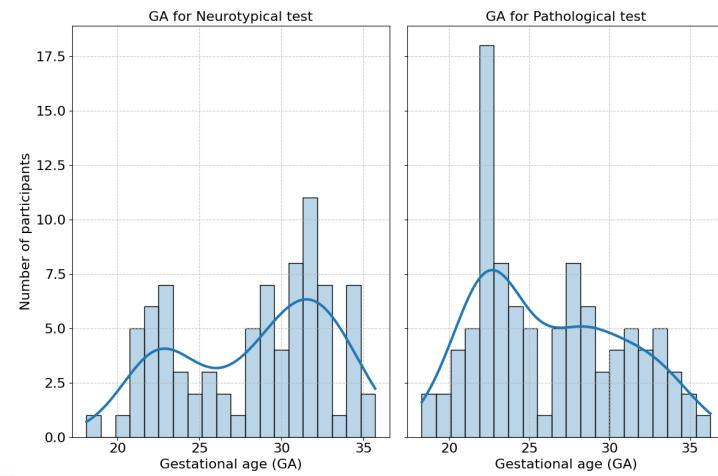
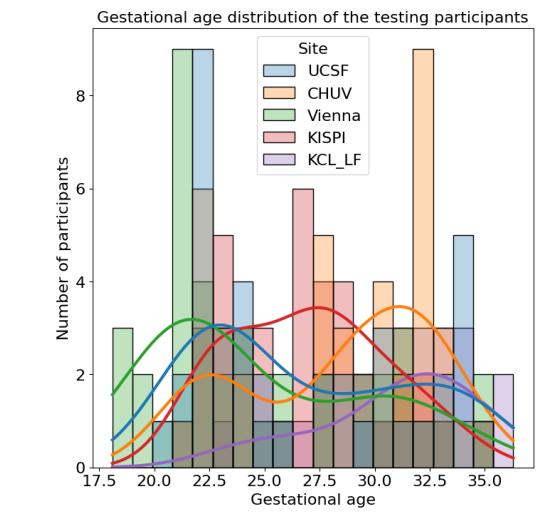
## Our goal with FeTA Challenge

Encourage the development of effective, domain-generalizable and reproducible methods for analyzing high resolution reconstructed MR images of the developing fetal brain from gestational week 21-36

# COHORTS

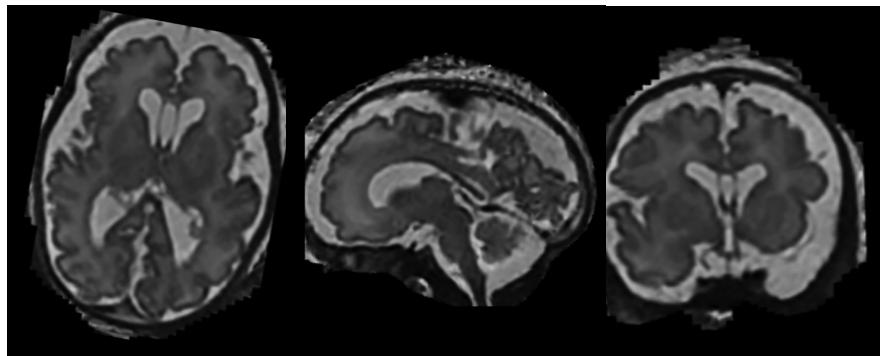


Payette et al 2021 Nature Scientific Data, Payette et al 2023 Media, FeTA Challenge 2022  
<https://arxiv.org/pdf/2402.09463.pdf>, Low-field data: Aviles-Verdera et al 2023 Radiology.

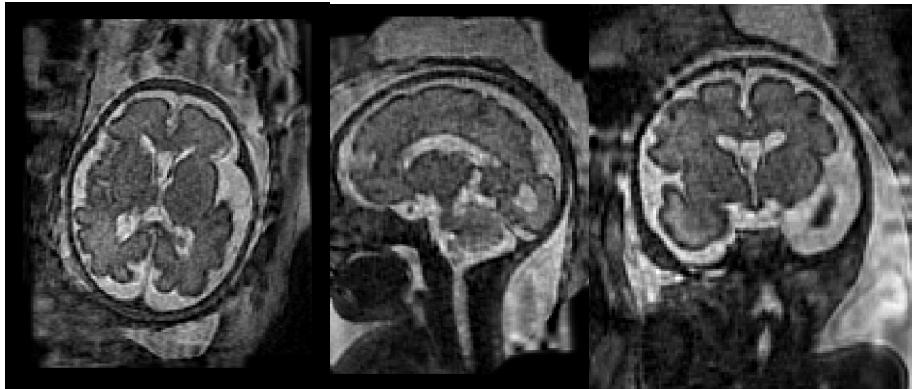


## COHORTS

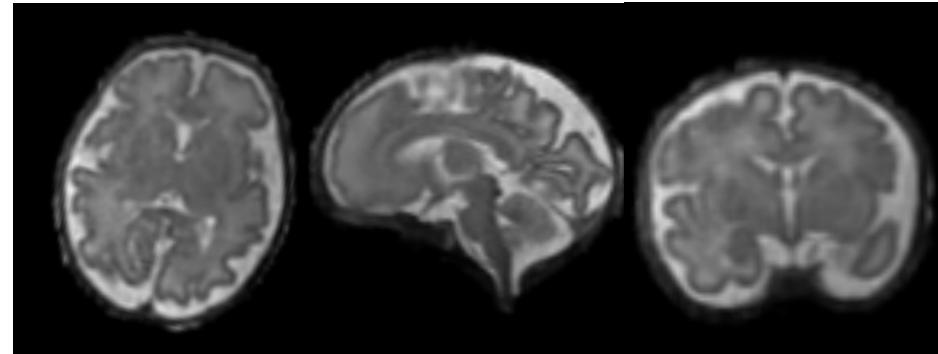
 & 3T IRTK



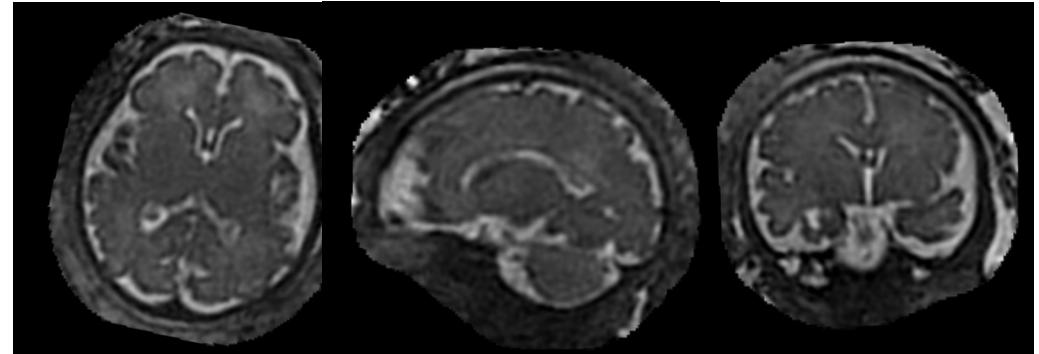
 & 1.5T Niftymic



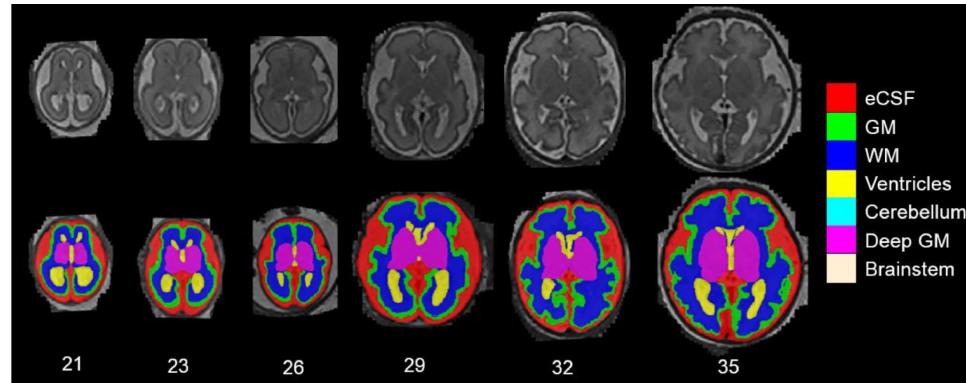
 & 1.5 T MIALSRTK



 0.55T & SVRTK



# SEMANTIC SEGMENTATION (TASK 1)



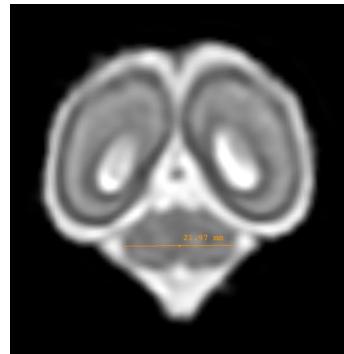
- Manual annotations of voxel-wise brain tissue class for LF MRI : 20 cases

- Same protocol than previous years
- Each person – 1 structure
- Fusion of structures
- Revisions by 2 expert raters
- Annotation time: from 8h to around 12h per case

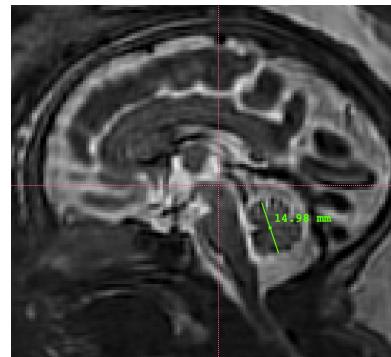
1. Margaux Roulet
2. Maurice Aschwanden
3. Yvan Gomez
4. Vladyslav Zalevskyi
5. Raphael Gaubert-Rachmühl
6. Céline Steger
7. Andras Jakab
8. Kelly Payette



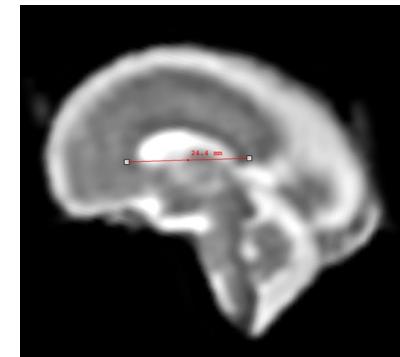
Biometry  
Measurements



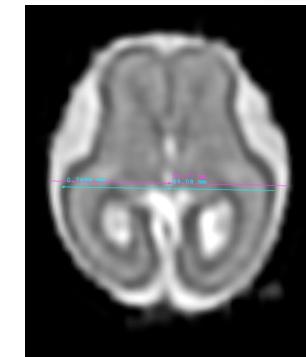
**TCD<sub>cor</sub>**, maximum transverse cerebellar diameter



**HV**, height of the vermis, measured in the mid-sagittal plane.



**LCC**, length of the corpus callosum, measured in the mid-sagittal plane.



**bBIP<sub>ax</sub>**, brain biparietal diameter : maximal brain diameter in the transverse plane through the atra.

**sBIP<sub>ax</sub>**, skull biparietal diameter: the inner-to-inner table maximal skull diameter in the transverse plane through the atra.

## ■ Manual annotations of 5 selected biometry measurements for all 300 cases

- Same protocol as [1,2,3,4,5]
- Each person – 1 case,
- Splits by centers
- Revisions by 1 expert rater

1. Mériam Koob
2. Yvan Gómez
3. Maurice Aschwanden
4. Raphael Gaubert-Rachmühl
5. Andras Jakab

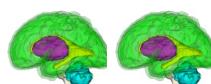


Tissue  
Segmentation

1. Overlap Dice similarity coefficient (DSC) ↑
2. Distance 95th percentile of Hausdorff distance (HD95) ↓
3. Volume similarity (VS) ↑
4. Topology Betti Number Error k (BNEk) ↓

$$\frac{2x}{\text{---} + \text{---}}$$

The diagram shows two sets of points, one blue and one red, with a green line connecting them. The formula  $\sup_{x \in X} \inf_{y \in Y} d(x,y)$  is written above the line.



k-dimensional Betti numbers in a 3D object:

BN0: connected components

BN1: holes

BN2 cavities in a 3D object

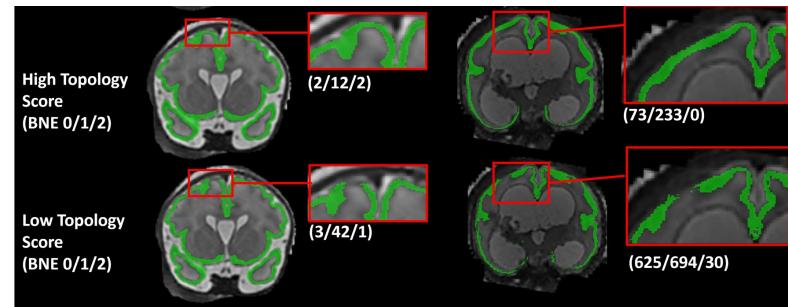


Figure extracted from XX

## EVALUATION



Biometry  
Measurements

1. Measurement Error in percentage (ME)

- Automated measurement
- Expert measurement



**MICCAI**

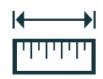
- Independent ranking per task
- Ranking per label and metric
- For each label, addition of 4 rankings
- All testing cases together

# PARTICIPANT TEAMS



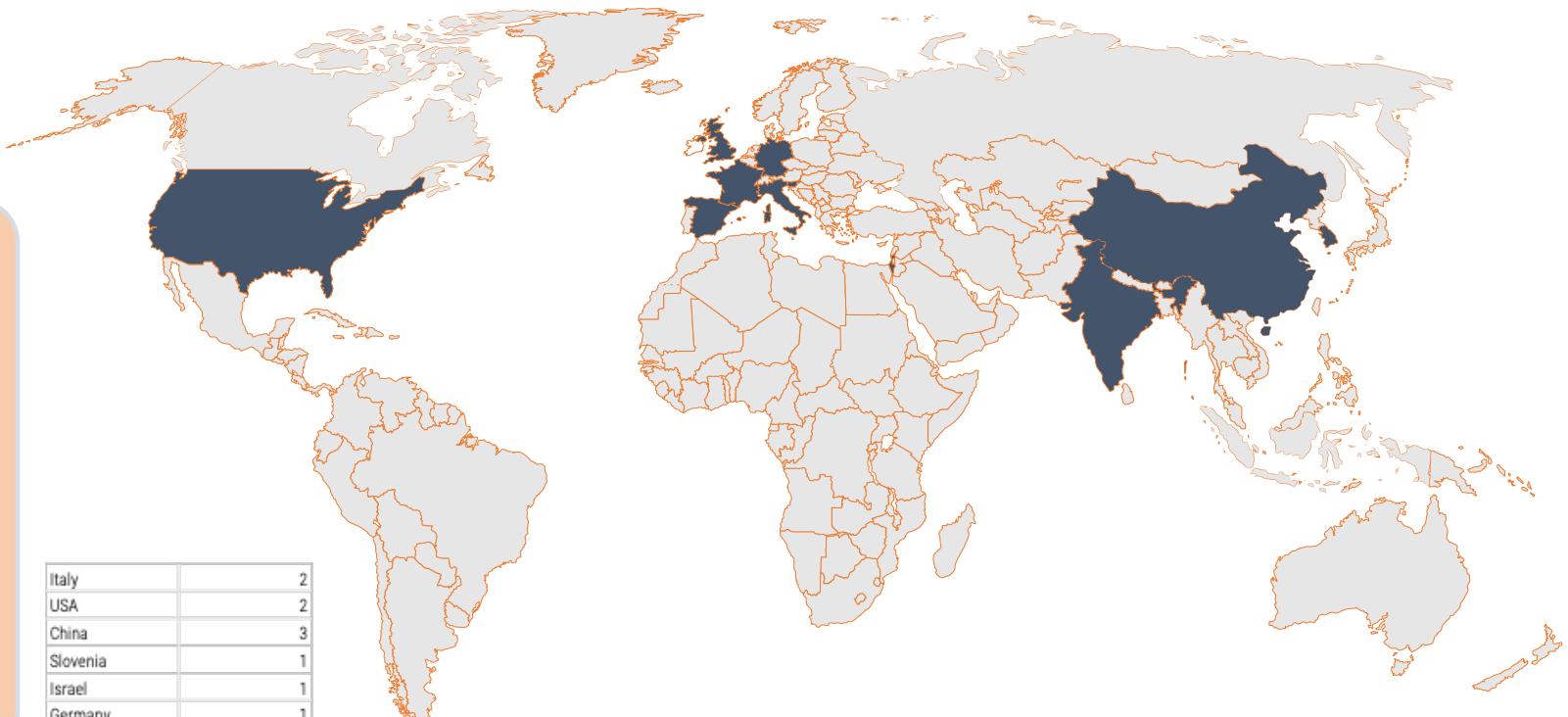
Tissue  
Segmentation

1. cemrg\_feta
2. cesne-digair
3. falcons
4. feta-sigma
5. hilab
6. jwcrad
7. lit
8. lmrcmc
9. mayanko
10. Mic-dkfz-feta24
11. paramahir\_2023
12. pasteurdbc
13. qd-neuroincyte
14. unipd-sum-aug
15. upfetal
16. vicorob



Biometry  
Measurements

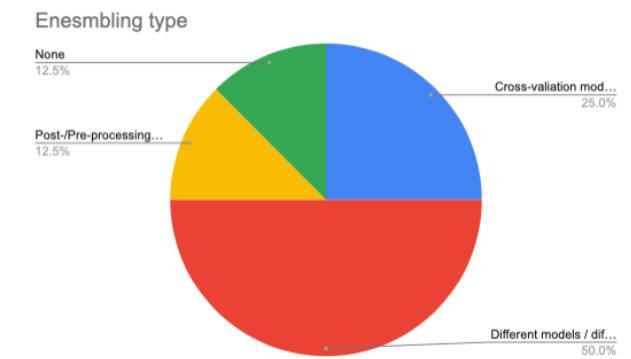
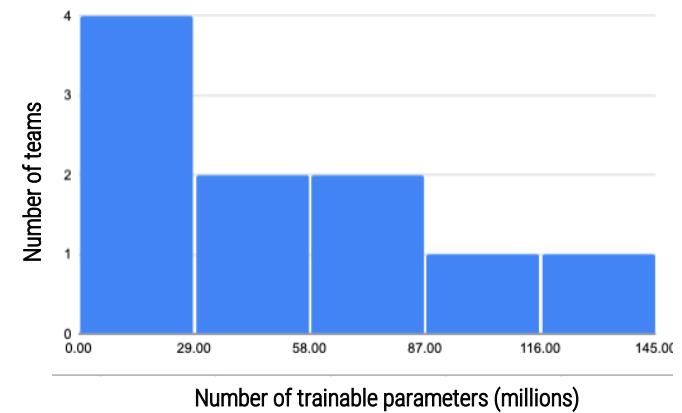
1. cesne-digair
2. falcons
3. feta\_sigma
4. jwcrad
5. paramahir\_2023
6. pasteurdbc
7. qd-neuroincyte



# SUBMITTED METHODS

## Segmentation

- Total Algorithms: 16
  - nnUnet: 9
  - U-Net + adaptations : 6
  - Swin\*(Mamba, Unetr): 3
  - MedNeXt\_L, SegVol, SegResNet: 3
- 3D / 2D: 14 / 2
- External datasets/atlasses: 5 (dHCP, Atlases or pre-trained foundation)
- All teams used Pytorch (except 1 TF)
- Ensemble learning: 9



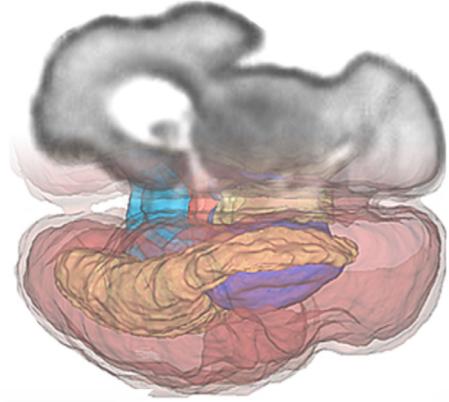


# SUBMITTED METHODS

## Biometry

- Total Algorithms: 7
  - nnUnet: 2
  - U-Net + adaptations : 4
  - CNN: 1
- 3D / 2D: 6/1
- External datasets/atlasses: 3 (dHCP, Atlases or pre-trained foundation)
- All teams used Pytorch (except 1 TF)
- All relied on segmentation map, some also input image
- Landmark heatmap: 2
- Landmark regression: 2
- Biometry regression: 2
- Missing description: 1





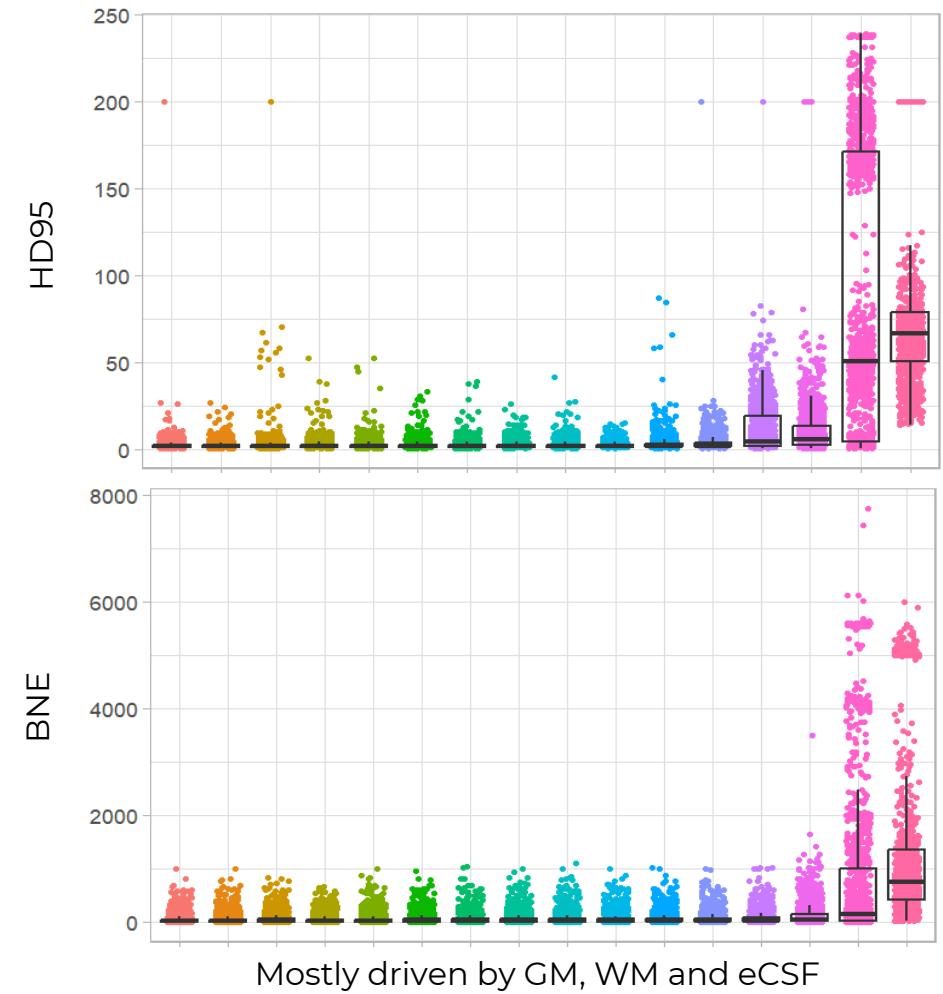
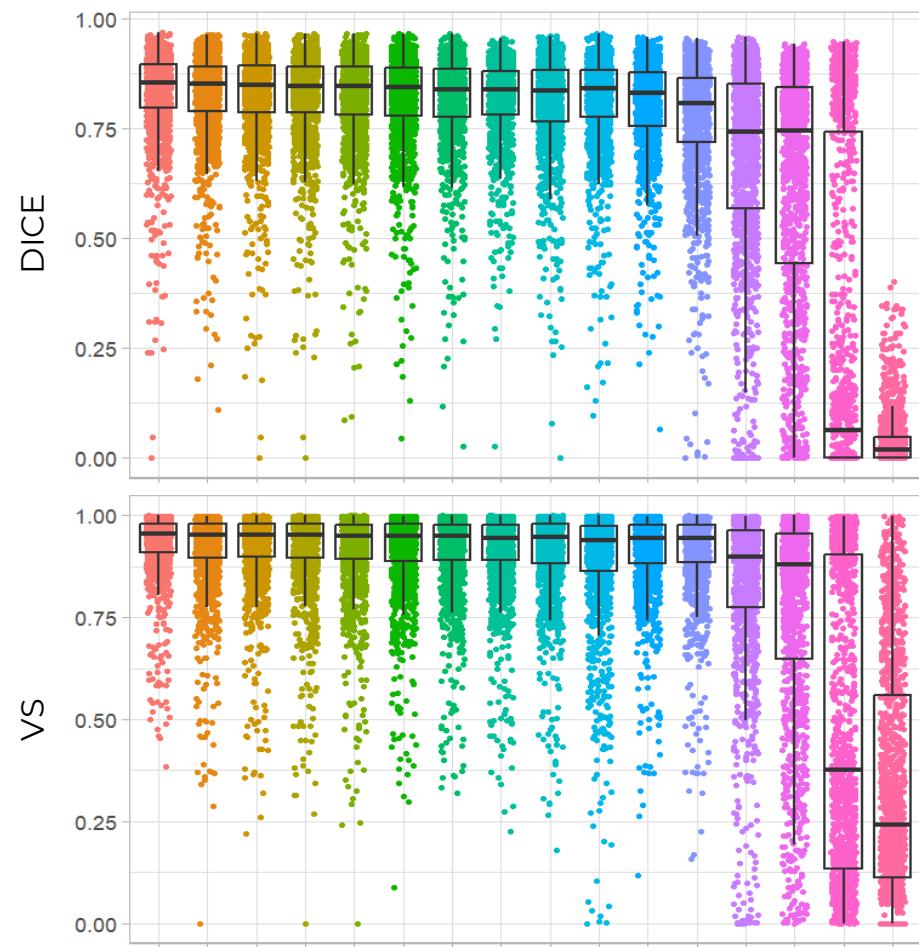
# *FeTA*

## SEGMENTATION TASK

RESULTS

# SEGMENTATION RESULTS: global test set

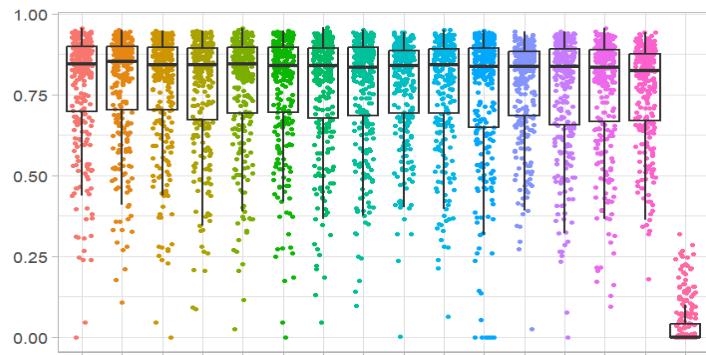
(Team colors are not consistent across plots)



# SEGMENTATION RESULTS: DICE per-site results

(Team colors are not consistent across plots)

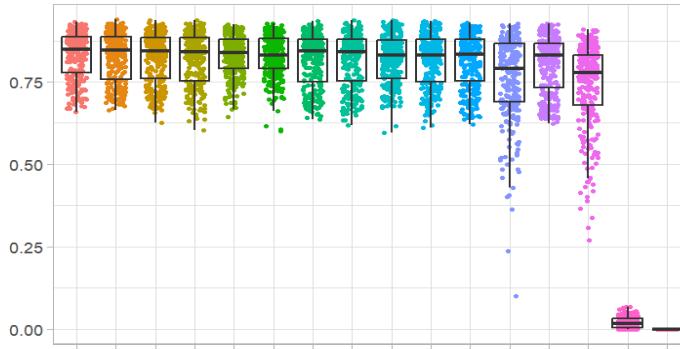
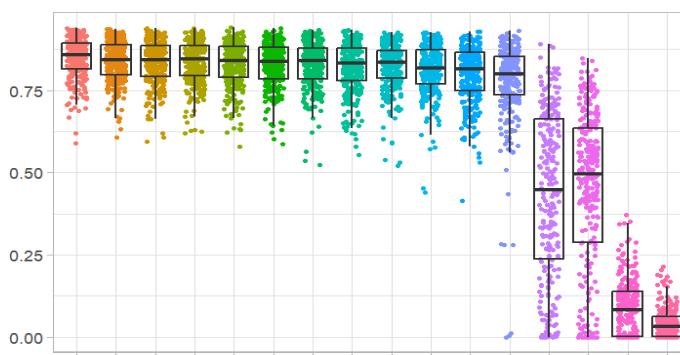
KISPI



Vienna

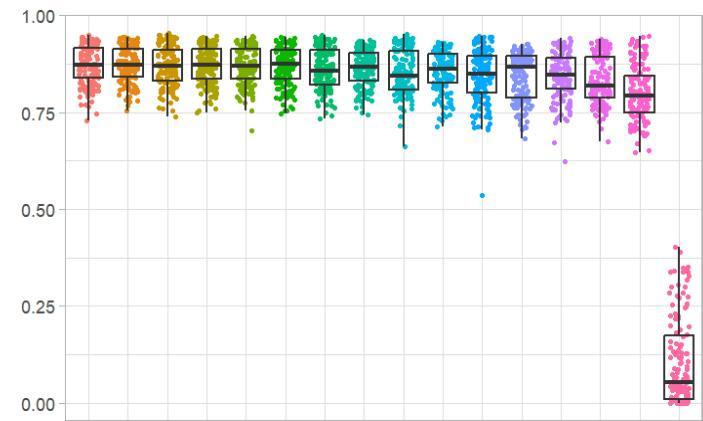
In-domain

UCSF



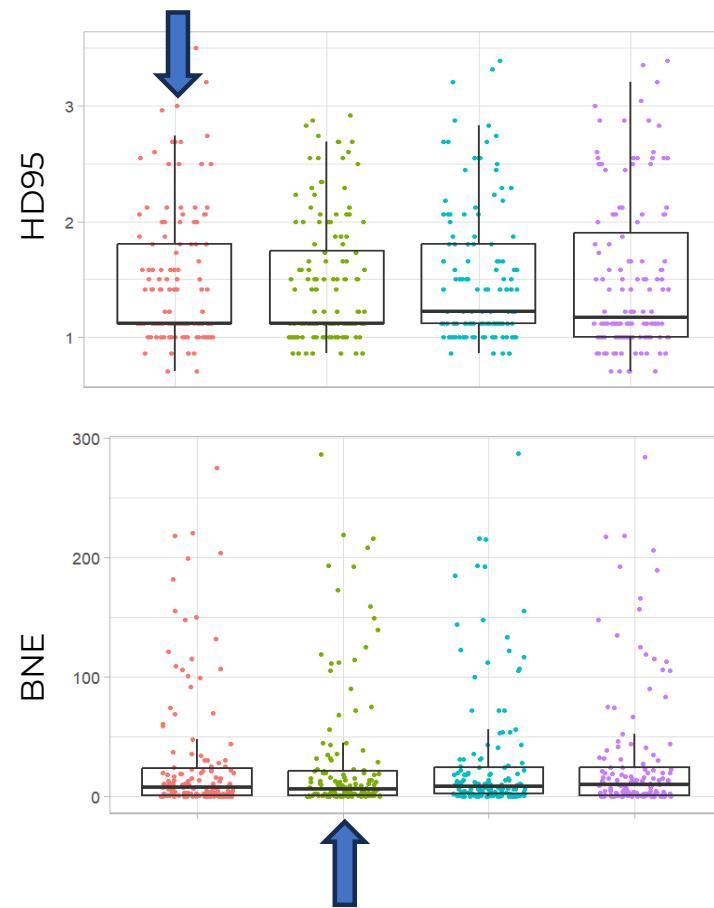
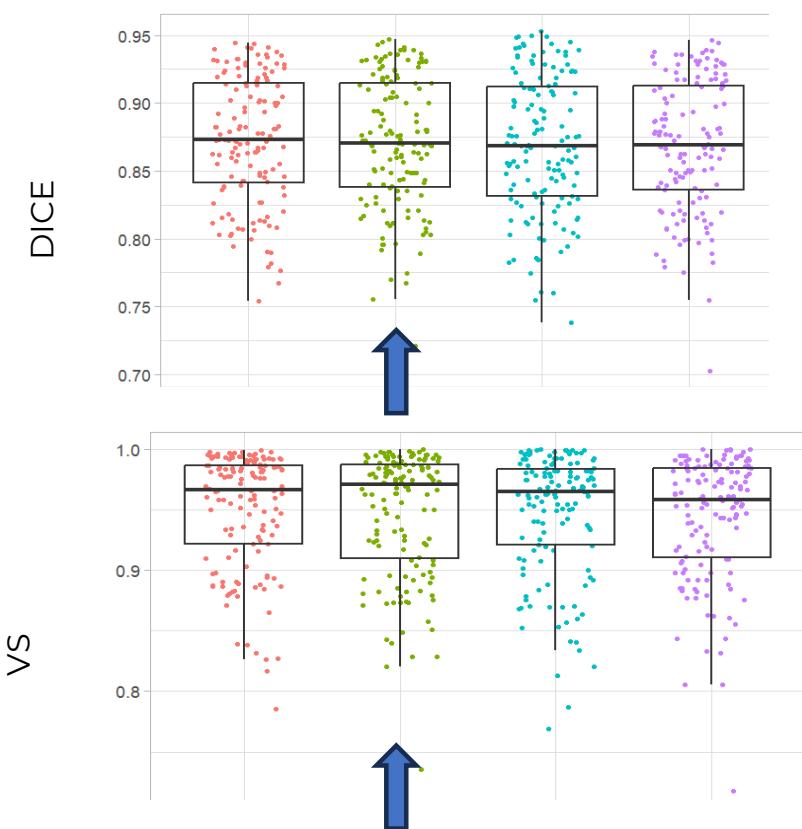
CHUV

KCL

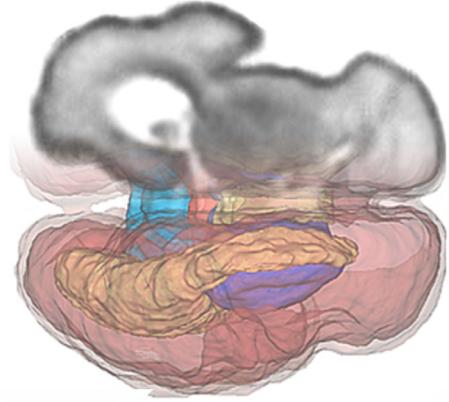


Out-of-domain

# Comparison of 3 top teams on Low-Field 0.55T scanner with FeTA 2022 winner\*



\*FIT - nnUNet, <https://arxiv.org/pdf/2402.09463>

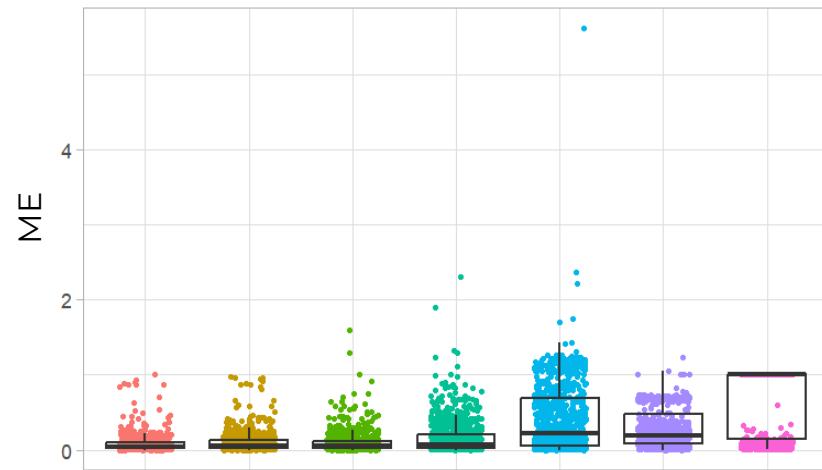


# *FeTA*

## BIOMETRY TASK

RESULTS

# BIOMETRY RESULTS: all sites



Algorithm	HV	LCC	TCD	bBIP	sBIP	ME
Algorithm	0.103	<b>0.112</b>	0.072	0.054	0.048	0.077
Algorithm	0.098	<b>0.177</b>	0.123	0.04	0.047	0.096
Algorithm	0.116	<b>0.126</b>	0.137	0.057	0.055	0.098
Algorithm	0.435	<b>0.205</b>	0.054	0.065	0.037	0.158
Algorithm	0.428	<b>0.328</b>	0.479	0.384	0.378	0.401
Algorithm	0.293	<b>0.285</b>	0.308	0.263	0.256	0.281
Algorithm	0.107	<b>0.088</b>	0.027	0.035	0.097	0.071

# BIOMETRY RESULTS: baseline comparisons

- We setup two regression model baselines:
  1. Uses only GA values
  2. Combine GA and total brain volume

Algorithm	HV	LCC	TCD	bBIP	sBIP	Total ME
Baseline (GA + GA <sup>2</sup> + Brain Vol.)	0.106	0.103	0.101	0.042	0.034	0.077
Algorithm	0.103	0.112	0.072	0.054	0.048	0.077
Baseline (GA + GA <sup>2</sup> )	0.108	0.122	0.108	0.068	0.065	0.094
Algorithm	0.098	0.177	0.123	0.04	0.047	0.096
Algorithm	0.116	0.126	0.137	0.057	0.055	0.098
Algorithm	0.435	0.205	0.054	0.065	0.037	0.158
Algorithm	0.428	0.328	0.479	0.384	0.378	0.401
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Algorithm	0.107	0.088	0.027	0.035	0.097	0.071

## NEXT STEPS

### Further analysis of the results & paper writing

- Annotation of SR image quality <sup>[1,2]</sup> : link between quality and generalization
- Comparing results from normal and pathological cases
  
- Segmentation task:
  - Exploring *normalized* Dice <sup>[3]</sup> and GA effects
  - Deeper insights from topology errors
  - Comparison with previous top teams (from FeTA2022) <sup>[4]</sup>
  
- Biometry task:
  - Inter-rater variability <sup>[5]</sup>
  - Baseline models comparison
  - Moving to landmark error estimation

<sup>1</sup>T. Sanchez et al. MEDIA (2024); <sup>2</sup>T. Sanchez et al PIPPI 2024; <sup>3</sup>V. Raina et al ISBI 2023; <sup>4</sup>K. Payette et al arxiv 2402.09463 ; <sup>5</sup>T. Sanchez et al Medrxiv 10.1101/2024.09.23.24313965v1



# THANKS!

- Volunteers accepting to participate in studies
- Funding institutions
- All participating teams !!!



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University of  
Zurich UZH



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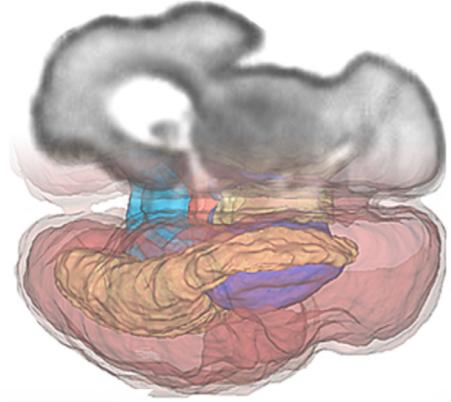
Q/A

- Robin Cremese: “Mednext adoption for fetal brain MRI segmentation”
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Q/A

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**12:00 - 13:30** Joint PIPPI & FeTA Poster Session



*FeTA*

SEGMENTATION TASK

WINNERS



# FeTA Segmentation Rankings



Ranking	Team Name
1	....
2	....
3	....
4	Feta_sigma
5	Pasteur DBC
6	UPFetal24
7	LIT <span style="color:red">Ranked 1st for Topology metric</span>
8	cesne-digair <span style="color:red">Ranked 1st for Volume Similarity</span>
9	Hilab

Ranking	Team Name
10	unipd-sum-aug
11	lmrcmc
12	jwcrad
13	qd_neuroincyte
14	falcons
15	ichilov-tau-maya
16	paramahir_2023



# FeTA Segmentation Rankings



## ViCOROB

Rachika Elhassna Hamadache, Amina Bouzid, Ricardo Montoya del Ángel, Marawan Elbatel, Cansu Yalçın, Hadeel Awwad, Adrià Casamitjana, Arnau Oliver, Robert Martí Marly, Xavier Lladó Bardera

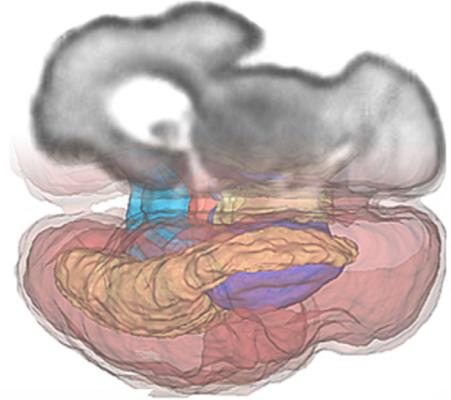


## mic-dkfz-feta2024

Maximilian Zenk, Michael Baumgartner, Klaus Maier-Hein

## cemrg\_feta

Abdul Qayyum, Moona Mazher, Steven A Niederer



*FeTA*

BIOMETRY TASK

WINNERS



# FeTA Biometry Rankings



Ranking	Team Name
1	....
2	....
3	....
4	PasteurDBC
5	qd_neuroincyte
6	Paramahir_2024
7	falcons

# FeTA Segmentation Rankings

## **cesne-digair**

Tommaso Ciceri, Marina Di Stefano, Giulia Frigerio, Giorgio Longari, Francesca Maccarone, Simone Melzi, Denis Peruzzo, Rocco Prudentino, Gloria Rizzato



## **jwcrad**

Jae Won Choi

## **feta\_sigma**

Jiang Jingwen, Zhang Chengsheng, Wang Hanling, Zhang Xuezhi, Cao Jiarui, Tong Lyuyang, Du Bo



# Challenge Pictures





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