

Group	Model Desc.	Model Param.	Time	G P U	Used_Images	Used_Demographic_Info	Used_Scanner_Info	Used_APOE_Info	VG1		VG2		VG3		VG4		VG5		Avg		CN		MCI		Dem.	
									M A E	R <sup>2</sup>	M A E	R <sup>2</sup>	M A E	R <sup>2</sup>	M A E	R <sup>2</sup>	M A E	R <sup>2</sup>	M A E	R <sup>2</sup>	M A E	R <sup>2</sup>	M A E	R <sup>2</sup>	M A E	R <sup>2</sup>

<b>Group</b>	indicates different labs participating in the challenge: Lab 1, 2, 3.
<b>Model Desc.</b>	Includes description of type of models used e.g. ML Model like XGBoost, Deep Learning Architecture (e.g., 2D or 3D CNN, Vision Transformer, etc.)
<b>Model Param.</b>	indicates the size of the model.
<b>Time</b>	indicates the amount of time it took to train the model. (Pretraining time for foundation models may not be known).
<b>GPU</b>	indicates the type of GPU used.
<b>Used_Images</b>	indicates whether the T1w scans were used to train the model (0 - NO, 1 - YES).
<b>Used_Demographic_Info</b>	indicates whether demographic data were used to train the model (0 - NO, 1- YES).
<b>Used_Scanner_Info</b>	Indicates whether the model used any scanner information for training (0 - NO, 1- YES).
<b>Used_APOE_Info</b>	indicates whether APOE genotype values were used to train the model (0 - NO, 1- YES).
<b>VG 1-5</b>	represents the Cross-Validation Groups 1-5. For each cohort, the MAE and R <sup>2</sup> score must be reported
<b>Avg</b>	indicates the average MAE and R <sup>2</sup> score for all the Cross-Validation Groups.
<b>CN, MCI and Dem.</b>	indicate the MAE and R <sup>2</sup> score for the diagnostic groups: Controls, MCI and dementia, respectively.