

# **Functional Description**

adVAE aims to:

- 1. Aid Alzheimer's research by creating high-quality multimodal synthetic data (gene expression, EEG, MRI)
- 2. Easily integrate with existing biomedical workflows

#### User Interface

Command line interface (CLI)

### Functionality

- Preprocess input data used to train the VAE
- 2. Latent space representation of various data modalities
- Generation of synthetic data using latent space variables

#### Task Prioritization

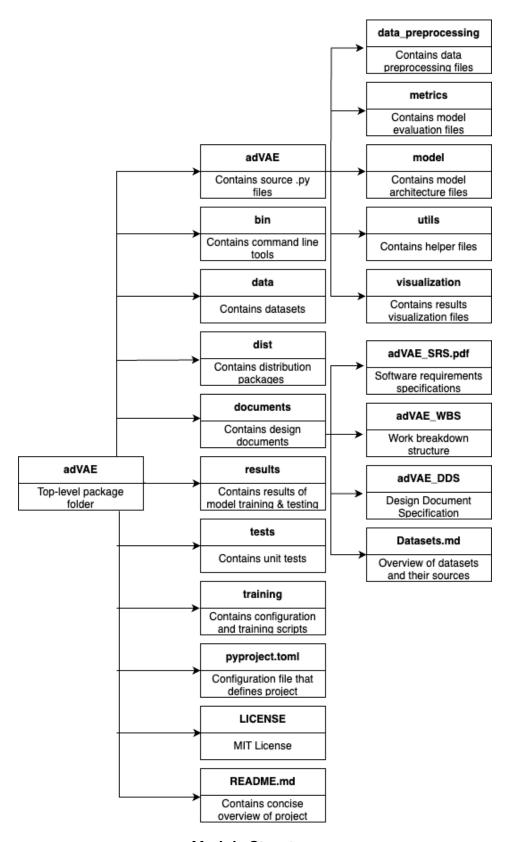
- 1. Identify example data (gene expression, EEG and MRI)
- 2. Create VAE model architecture for one modality (gene expression)
- 3. Train the model using the example gene expression dataset
- 4. Perform latent space optimization
- 5. Validate the results by calculating KL loss and reconstruction error of generated dataset
- 6. Extend the model to include other modalities
- 7. Train, validate, and test the model using the bulk data

### **Goals and Milestones**

- 1. Finish all design documentation
- 2. Set up GitHub repository
- 3. Create modules
- 4. Start implementing code
- 5. Create model for one modality
- 6. Extend model for other modalities
- 7. Peer review
- 8. Refine model and update design documentation

# Solutions

- 1. Tensorize all input data
- 2. Create separate modules/classes for handling different data modalities
- 3. Validate by calculating KL loss and reconstruction loss



**Module Structure** 

	Week 1 (02.03 - 02.09)				We	ek2	(02.1	0 - 0	2.016	)	Week 3 (02.17			- 02	23)		Week 4					Week 5						Week 6						Wee						Veek I				
Tasks	М	T۱	N TI	h F	S	Su	м т	W	/ Th	F	S S	u M	Т	W	Th F	- s	Su	М	T۱	N TI	ı F	S	Su	м т	W	Th F	s	Su	М 1	r w	Th	FS	Su	М	T١	N Th	ı F	S	Su I	ч т	W	Th I	F S	St
Create SRS																							$\Box$																П					
Create Datasets.md																																												
Create WBS																																												
Create Timeline																																												
Create DDS																																												
Identify data sources																																												
Validate data																																												
Implement first function																							$\Box$																					
Split dataset into example-training-validation-test data																																												
Perform data preprocessing for gene expression data																							П																T					
(normalization, scaling, handling missing values)																																												
Perform exploratory statistical analysis on the example data		ш										$\perp$							ш				_			ш								ш				ш	_			ш	_	
Visualize the example data																																												
Construct the VAE architecture																							$\Box$																					
Set up hyperparameter tuning, backpropagation and training																																												
functions												$\perp$											_																_				_	_
Set up loss functions and other validation metrices		ш		_				_				$\perp$				_			ш			_	_			ш	_			_		_		$\perp$	_			ш	_			_	_	_
Train and test model on example dataset																							$\Box$																					
Evaluate reconstruction quality																																												
Optimize latent space for gene expression VAE model																																												
Adapt model for MRI data												$\perp$											$\perp$															Ш	$\perp$				_	
Adapt model for EEG data																																												
Validate using full training, validation and test data																																												
Develop Command Line Interface for adVAE																							$\Box$																					
Update all design documentation																																												
Summarize key findings																																												
Prepare for peer review																																												
Present findings																							$\Box$																7					I

**Timeline**