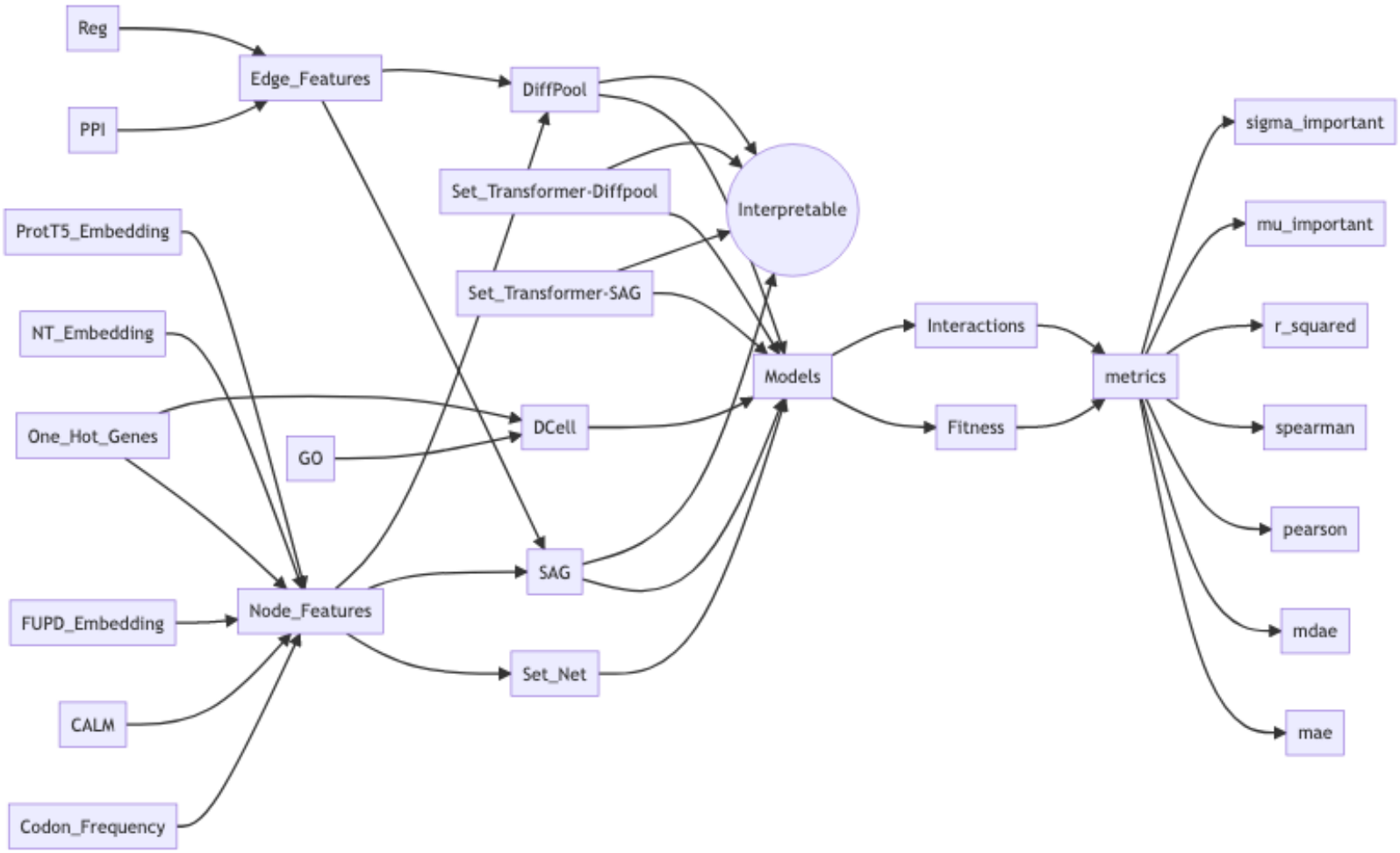
model	nodes features ($\mathcal{N}$)	edge features ($\mathcal{E}$)	mae	mdae	pearson	spearman	r^2	μ(1.0-1.1)	σ(1.0-1.1)	μ(1.1-1.2)	σ(1.1-1.2)
DCell	One Hot Genes	-									
Set Net	One Hot Genes	-									
Set Net	Codon Frequency	-									
Set Net	CALM	-									
Set Net	Fungal-UTR-Transformer Embeddings	-									
Set Net	Nucleotide-Transformer Embeddings	-									
Set Net	Prot T5	-									
Set Transformer - SAG	One Hot Genes	-									
Set Transformer - SAG	Codon Frequency	-									
Set Transformer - SAG	CALM	-									
Set Transformer - SAG	Fungal-UTR-Transformer Embeddings	-									
Set Transformer - SAG	Nucleotide-Transformer Embeddings	-									
Set Transformer - SAG	Prot T5	-									
Set Transformer - DiffPool	One Hot Genes	-									
Set Transformer - DiffPool	Codon Frequency	-									
Set Transformer - DiffPool	CALM	-									
Set Transformer - DiffPool	Fungal-UTR-Transformer Embeddings	-									
Set Transformer - DiffPool	Nucleotide-Transformer Embeddings	-									
Set Transformer - DiffPool	Prot T5	-									
SAG	One Hot Genes	PPI									

model	nodes features ($\mathcal{N}$)	edge features ($\mathcal{E}$)	mae	mdae	pearson	spearman	r^2	μ(1.0-1.1)	σ(1.0-1.1)	μ(1.1-1.2)	σ(1.1-1.2)
SAG	Codon Frequency	PPI									
SAG	CALM	PPI									
SAG	Fungal-UTR-Transformer Embeddings	PPI									
SAG	Nucleotide-Transformer Embeddings	PPI									
SAG	Prot T5	PPI									
DiffPool	One Hot Genes	PPI									
DiffPool	Codon Frequency	PPI									
DiffPool	Fungal-UTR-Transformer Embeddings	PPI									
DiffPool	Nucleotide-Transformer Embeddings	PPI									
DiffPool	Prot T5	PPI									
SAG	One Hot Genes	Reg									
SAG	Codon Frequency	Reg									
SAG	CALM	Reg									
SAG	Fungal-UTR-Transformer Embeddings	Reg									
SAG	Nucleotide-Transformer Embeddings	Reg									
SAG	Prot T5	Reg									
DiffPool	One Hot Genes	Reg									
DiffPool	Codon Frequency	Reg									
DiffPool	Fungal-UTR-Transformer Embeddings	Reg									
DiffPool	Nucleotide-Transformer Embeddings	Reg									
DiffPool	Prot T5	Reg									


Reg
PPI
Edge_Features
DiffPool
Set_Transformer-Diffpool
Interpretable
ProtT5_Embedding
Set_Transformer-SAG
NT_Embedding
Models
Interactions
Fitness
metrics
sigma_important
mu_important
r_squared
spearman
pearson
mdae
mae
One_Hot_Genes
GO
DCell
SAG
FUPD_Embedding
Node_Features
Set_Net
CALM
Codon_Frequency