

Dmf Fitness Table

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