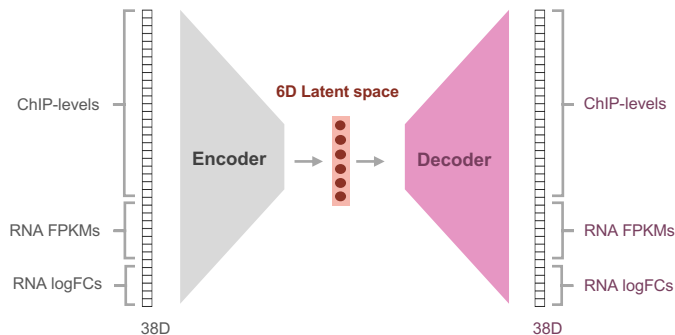


Reconstruction metrics:
Root Mean Squared Error
Cosine Similarity

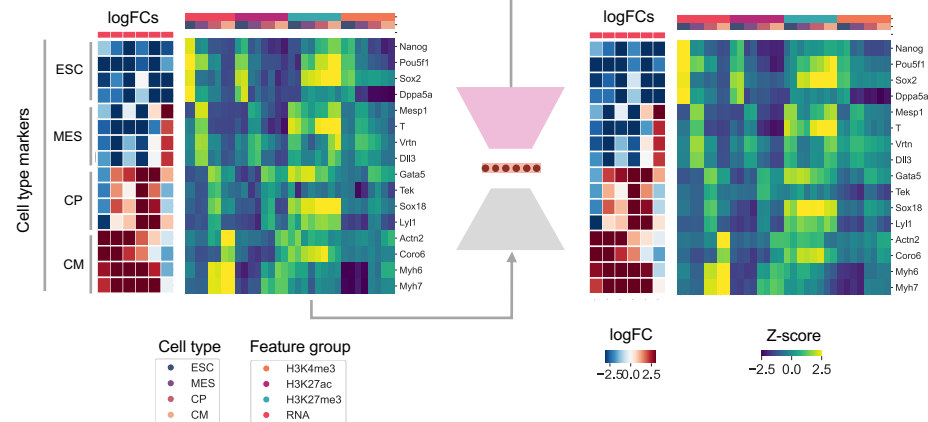
Input gene

Reconstructed gene



Original values

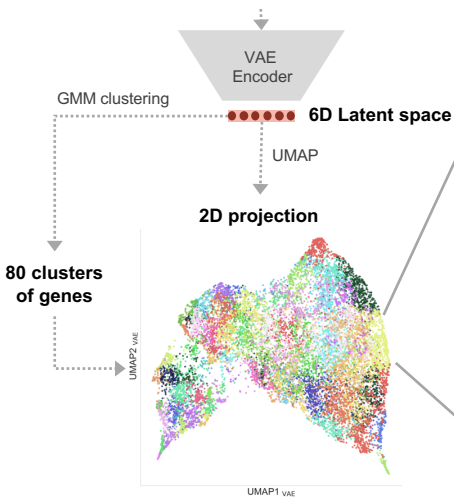
Reconstructed values



Dataset

Cluster n

Genome navigator interface (Web App)



Feature distributions

TSS plots

