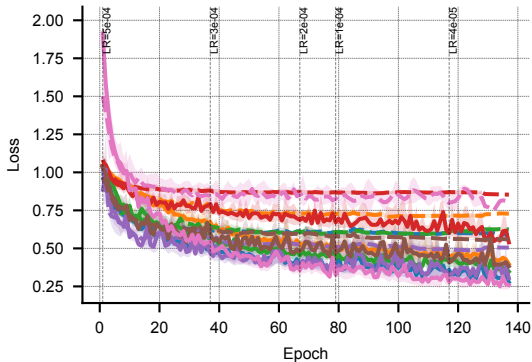


Total loss



Losses

variable

- mse_proteomics
- mse_metabolomics
- mse_drugresponse
- mse_crisprcas9
- mse_methylation
- mse_transcriptomics
- mse_copynumber

type

- train
- val