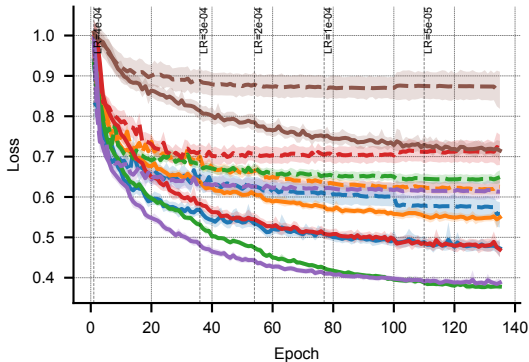


Total loss



Losses  
variable

- `mse_methylation`
  - `mse_transcriptomics`
  - `mse_proteomics`
  - `mse_metabolomics`
  - `mse_drugresponse`
  - `mse_crispcas9`
- type
- train
  - val