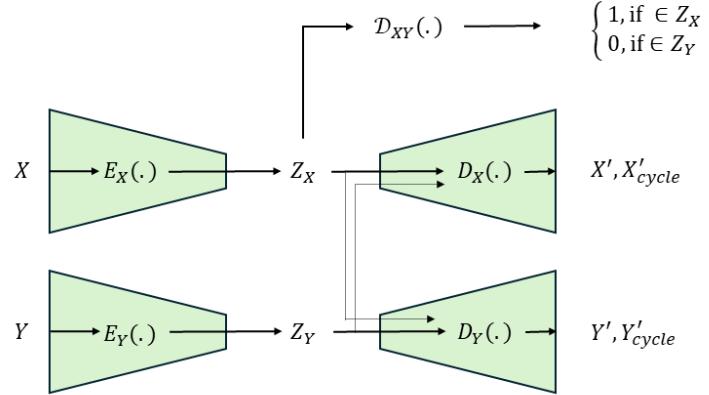


Cyclic multimodal autoencoder

We introduce a cycle-consistent autoencoder framework for integrating unpaired, multimodal single-cell RNA and ATAC data using a multiome-based bridge. We take inspiration from Seurat’s bridge integration strategy, along with the cycleGAN framework (Zhu et al 2017). Our approach outperforms bridge integration based on benchmarking using a ground truth multiomic dataset.

The model first learns a shared latent representation for each assay contained in paired multiomic data, jointly embedding RNA (X_0) and ATAC (Y_0) into a unified space. By enforcing cycle consistency, the we encourage cross-modal translations to be mutually coherent, producing denoised, biologically meaningful reconstructions in both modalities.

After training on multiomic data, our model can map unpaired single-cell RNA or ATAC datasets (X, Y) into the shared latent space for downstream integrative analysis by feeding the assays through the pretrained encoders. This approach provides a strategy for leveraging limited paired data to integrate large unpaired datasets across modalities.



$$\begin{aligned} \mathcal{L}(E_X, E_Y, D_X, D_Y; X, Y) = & \lambda_{recon} \cdot (MSE(D_X \circ E_X(X), X) + MSE(D_Y \circ E_Y(Y), Y)) \\ & + \lambda_{cycle} (MSE(D_X \circ E_Y(Y), X) + MSE(D_Y \circ E_X(X), Y)) \\ & + \lambda_{adv} (-\log D_{XY}(E_Y(Y))) \end{aligned}$$

Figure 1 – graphical representation of the underlying model. X and Y are mapped to a joint latent space, Z . Each set of embeddings, Z_X and Z_Y , are fed through both decoders $D_X(\cdot)$, $D_Y(\cdot)$. This ensures the learned latent representation encodes the joint structure of the data.

Model details

We begin with two paired noisy datasets, X_0 and Y_0 . We would like to learn two pseudo invertible mappings, $f: \mathcal{X} \rightarrow \mathcal{Z}$, and $g: \mathcal{Y} \rightarrow \mathcal{Z}$, to a latent space that encodes the joint structure of the data. By

pseudo-invertible, we mean that each function f, g , has a reverse mapping $\hat{f}^{-1}, \hat{g}^{-1}$ that maps the latent embeddings in Z , back to the original spaces, \mathcal{X}, \mathcal{Y} . To ensure that the latent embedding encodes both information about X_0 and Y_0 , we employ a multimodal autoencoder framework that utilizes cycle consistent loss.

First, two encoders, $E_X(X)$, $E_Y(Y)$, map X and Y to their latent embeddings. We define the reconstruction loss as follows:

$$\mathcal{L}_{recon}(E_X, E_Y, D_X, D_Y; X, Y) = MSE(D_X(E_X(X))) + MSE(D_Y(E_Y(Y)))$$

This alone is not enough to ensure that the learned embeddings, $E_X(X)$, $E_Y(Y)$, encode the joint structure of X and Y . We use cycle consistency loss to ensure that the latent space is representative of the joint structure $[X, Y]$:

$$\mathcal{L}_{cycle}(E_X, E_Y, D_X, D_Y; X, Y) = MSE(D_Y(E_X(X)), Y) + MSE(D_X(E_Y(Y)), X)$$

Note that during training, the gradient from \mathcal{L}_{cycle} only corresponds to the parameters of the encoders E_X and E_Y . The decoders D_X and D_Y are only trained using their corresponding modalities. This encourages the encoders to mix X and Y in the latent space.

Additionally, we employ adversarial training to aid in the mixing of the latent space. We use a discriminator, $\mathcal{D}_{XY}(Z)$, that classifies the modality of an embedding as $X (= 1)$ or $Y (= 0)$. The discriminator is trained using the following loss:

$$\mathcal{L}_{scrim}(\mathcal{D}_{XY}; X, Y) = -\log(\mathcal{D}_{XY}(E_X(X))) - \log(1 - \mathcal{D}_{XY}(E_Y(Y)))$$

And the adversarial loss is defined as

$$\mathcal{L}_{adv}(E_Y; Y) = -\log(\mathcal{D}_{XY}(E_Y(Y)))$$

Finally, we define our overall loss function as follows:

$$\begin{aligned} \mathcal{L}(E_X, E_Y, D_X, D_Y; X, Y) = & \lambda_{recon} \cdot \mathcal{L}_{recon}(E_X, E_Y, D_X, D_Y; X, Y) \\ & + \lambda_{cycle} \cdot \mathcal{L}_{cycle}(E_X, E_Y, D_X, D_Y; X, Y) \\ & + \lambda_{adv} \cdot \mathcal{L}_{adv}(E_Y; Y) \end{aligned}$$

Where λ_{recon} , λ_{cycle} , and λ_{adv} are user-specified weights, with default values of 1. A graphical overview of the model is shown in figure 1.