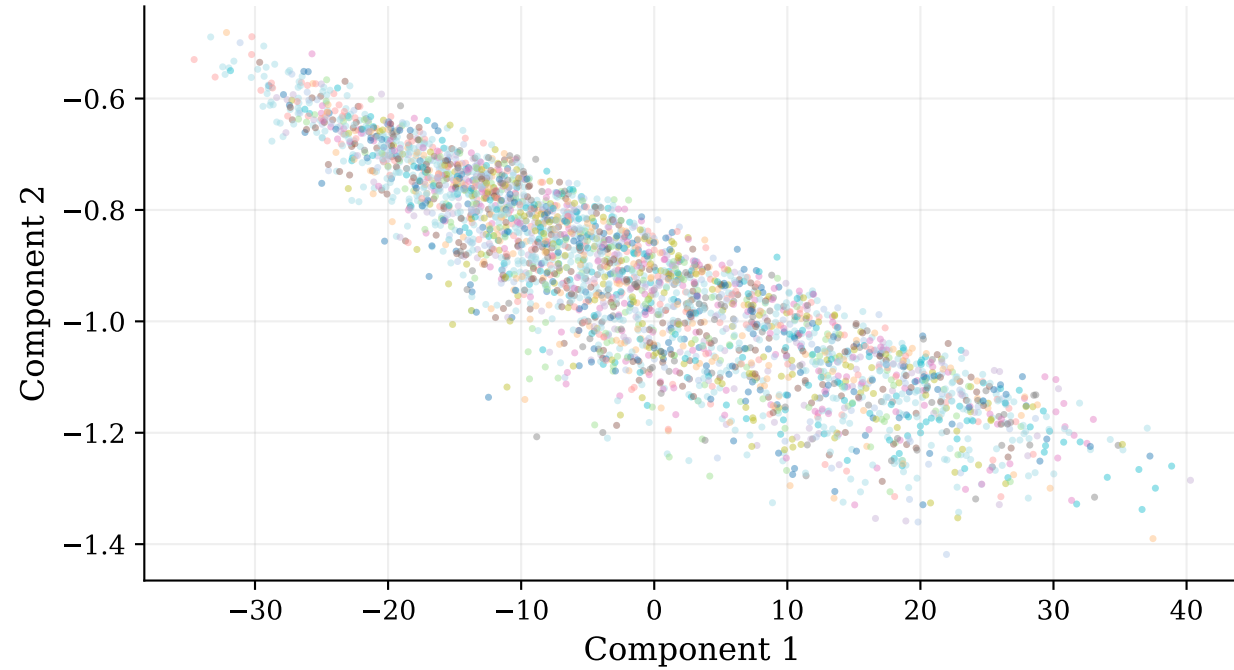


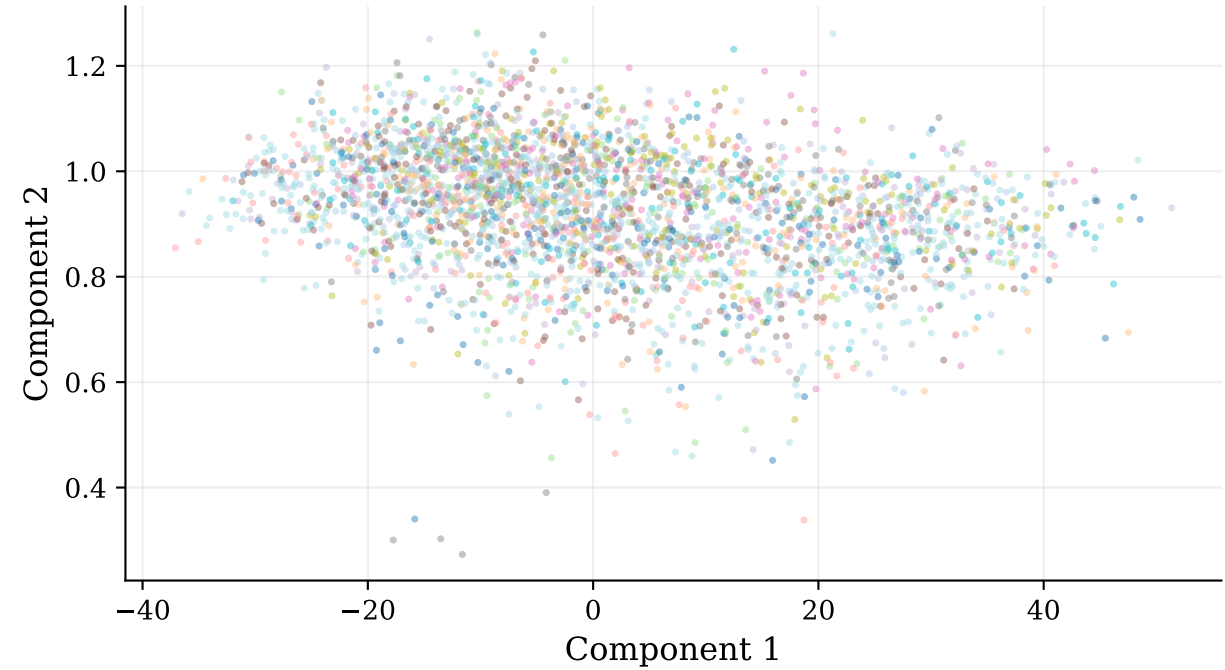
# RQ4 Representation Structure: PCA vs UMAP vs t-SNE

Eval split=eval | Probe split=eval | Samples: ERM=3000 DANN=3000

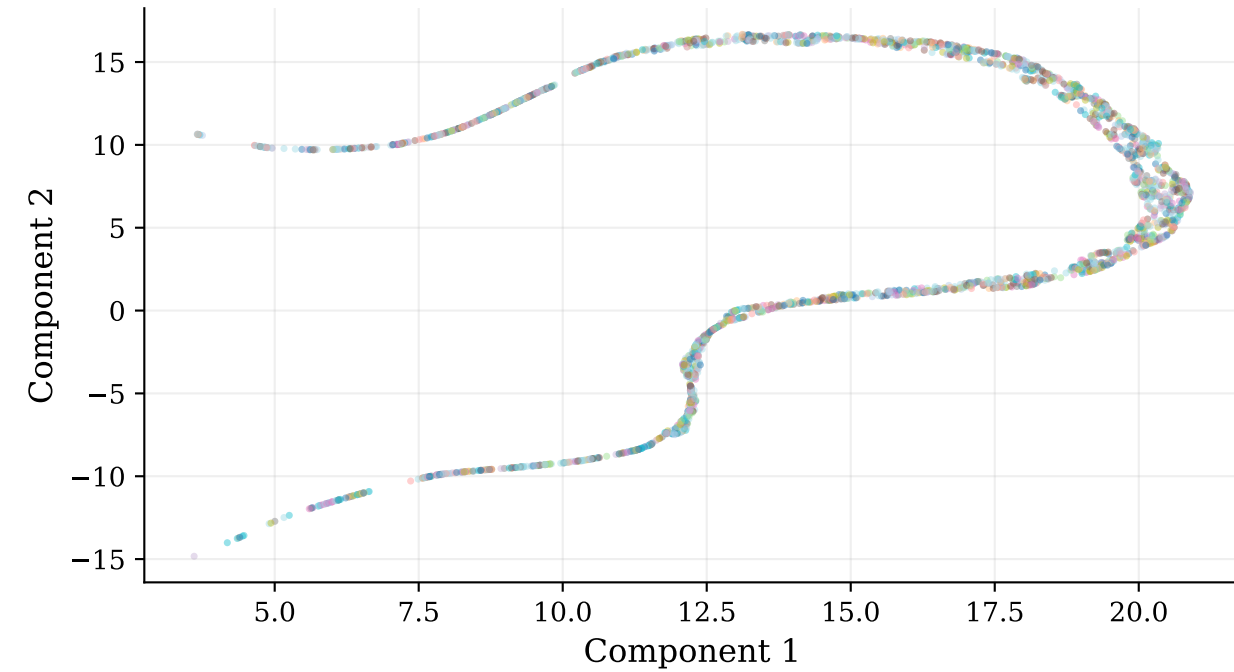
**PCA: ERM**



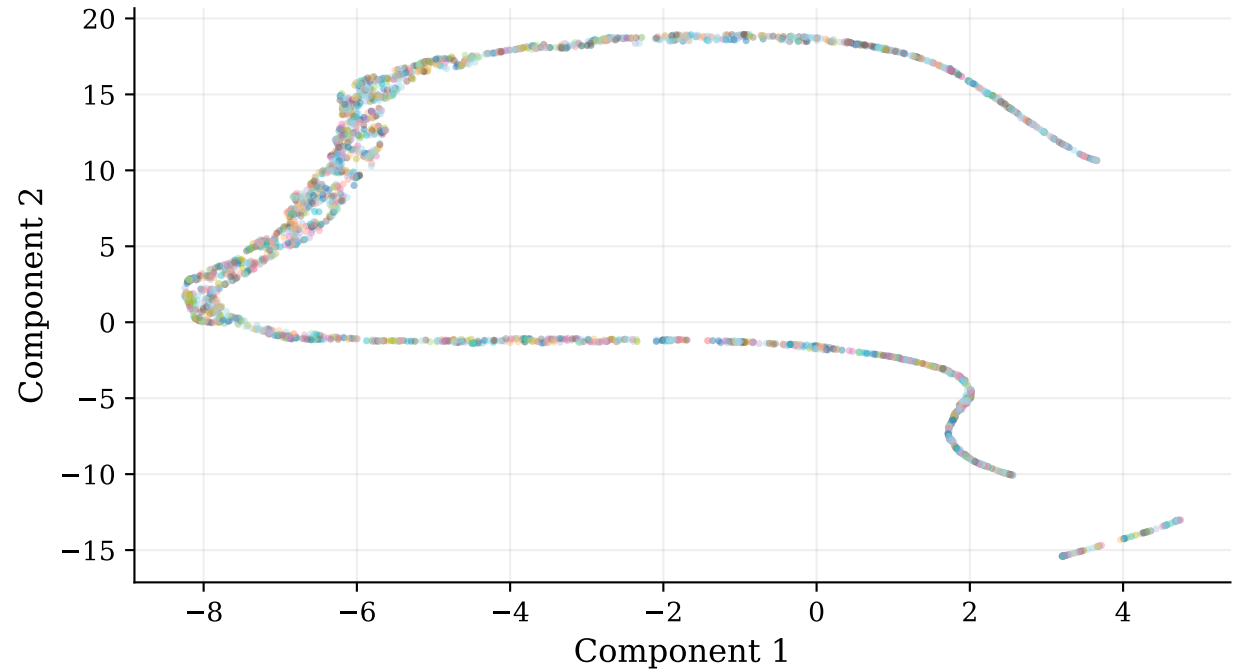
**PCA: DANN**



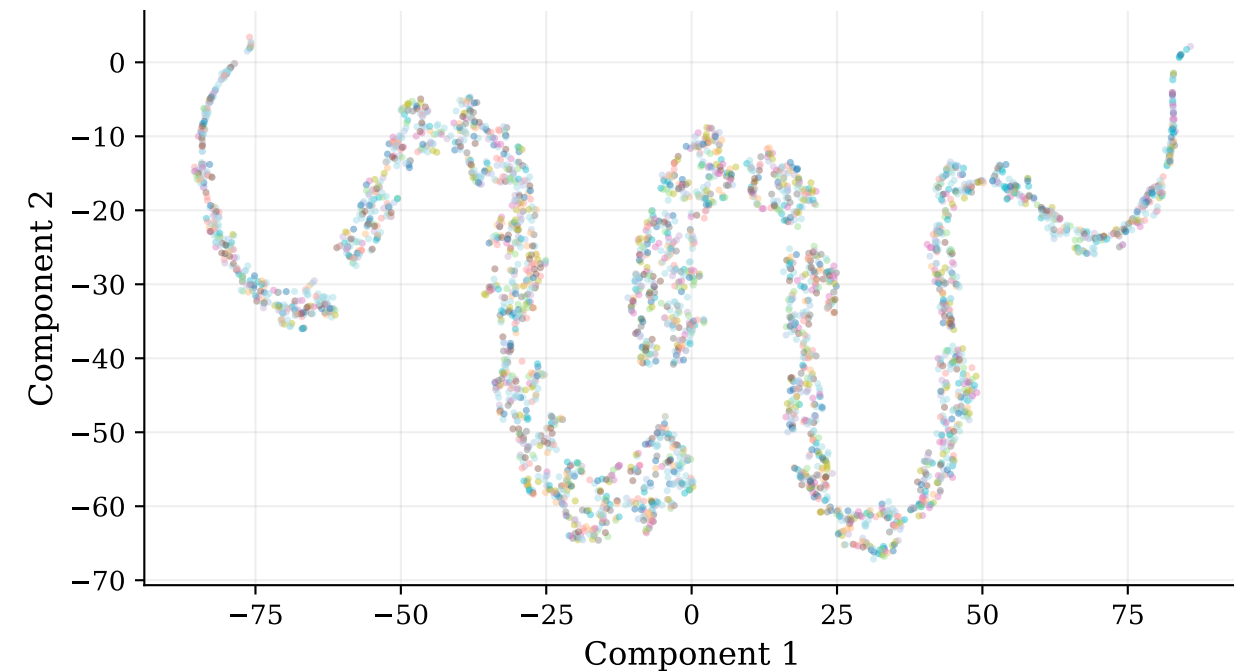
**UMAP: ERM**



**UMAP: DANN**



**t-SNE: ERM**



**t-SNE: DANN**

