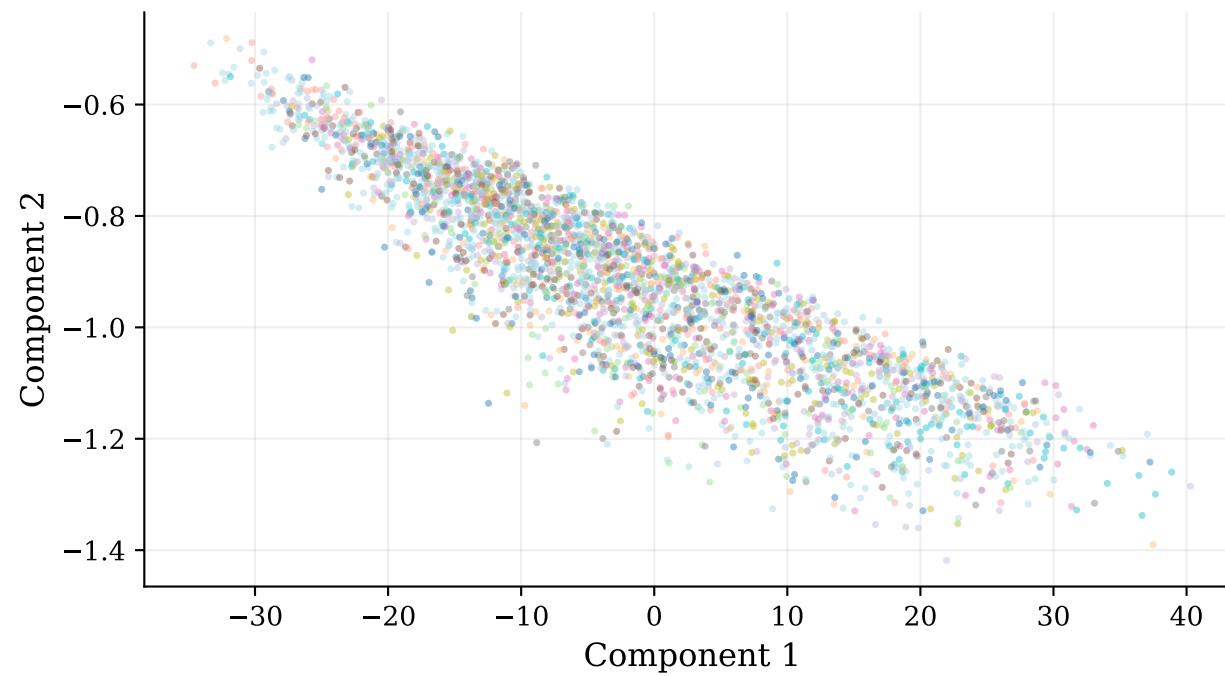


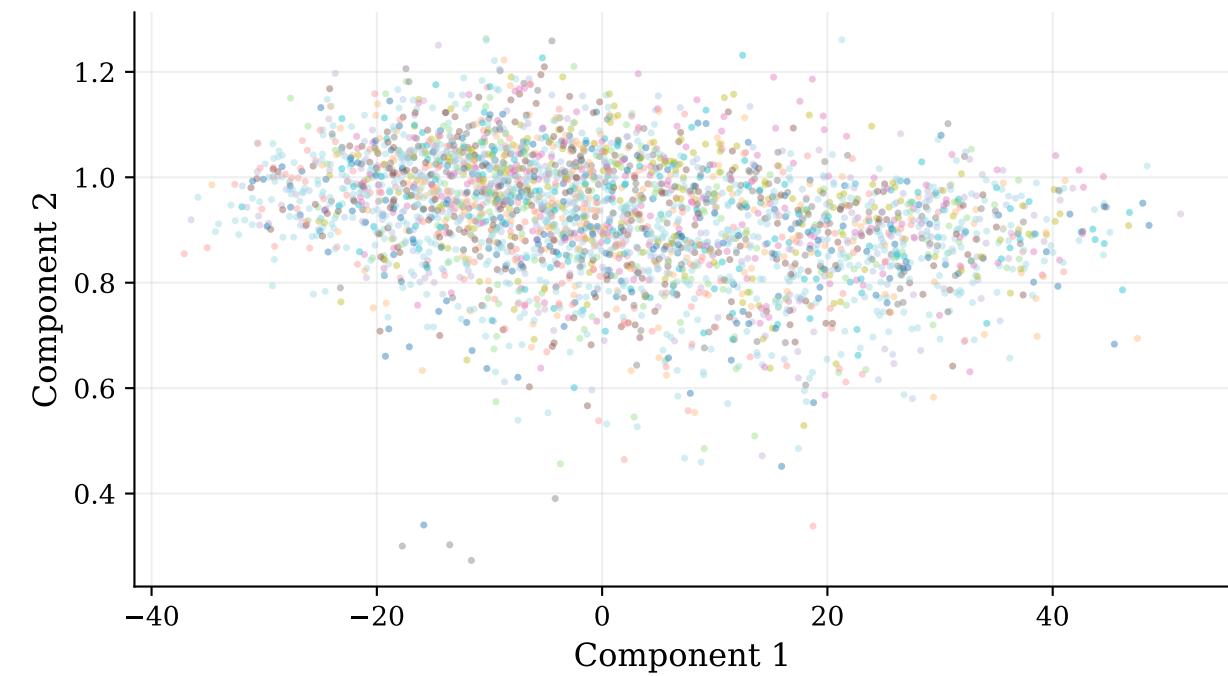
RQ4 Representation Structure: PCA vs UMAP vs t-SNE

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

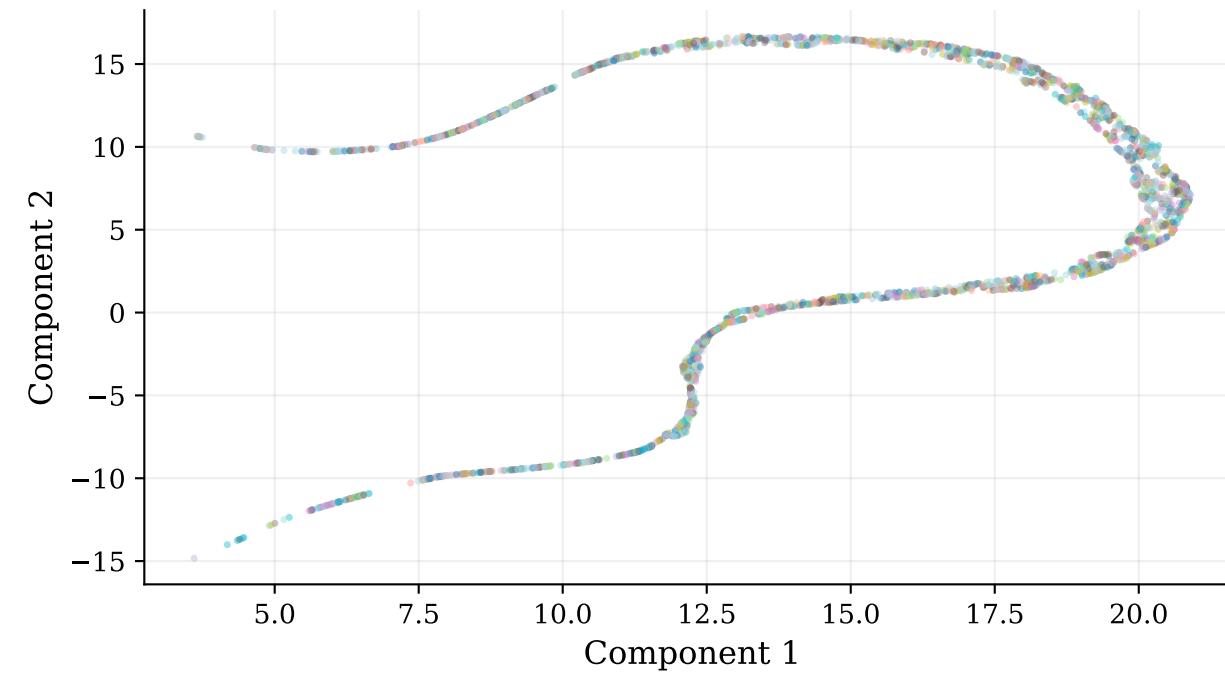
PCA: ERM



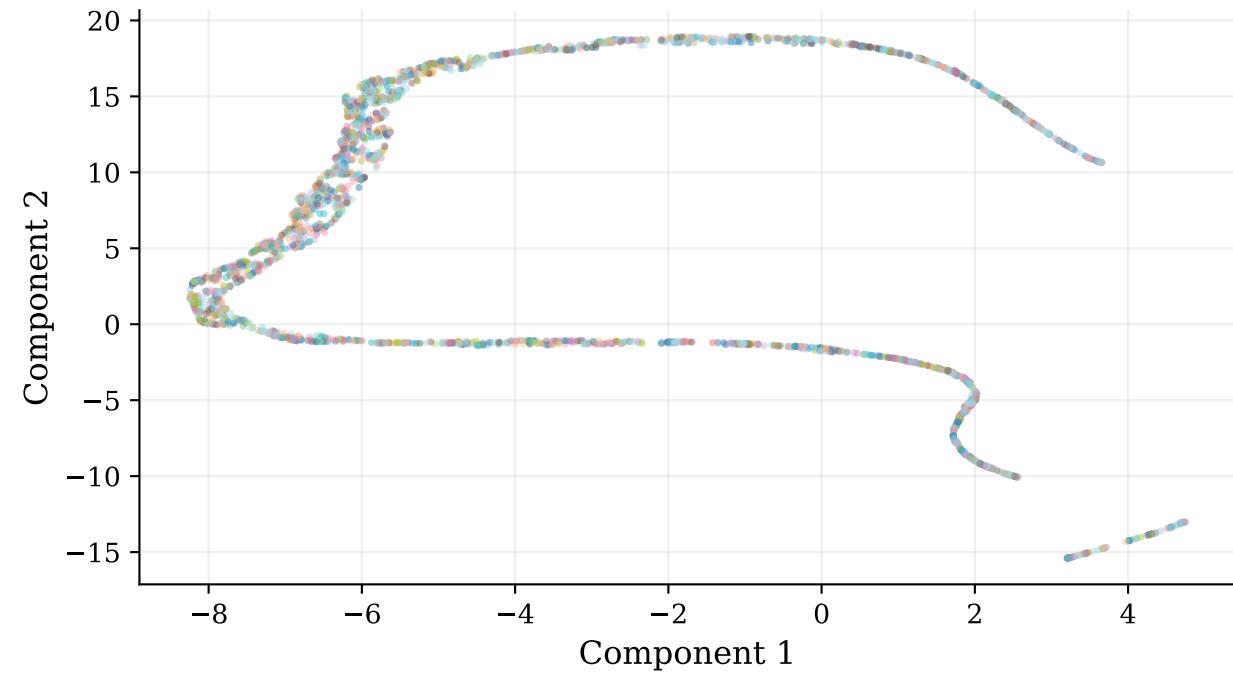
PCA: DANN



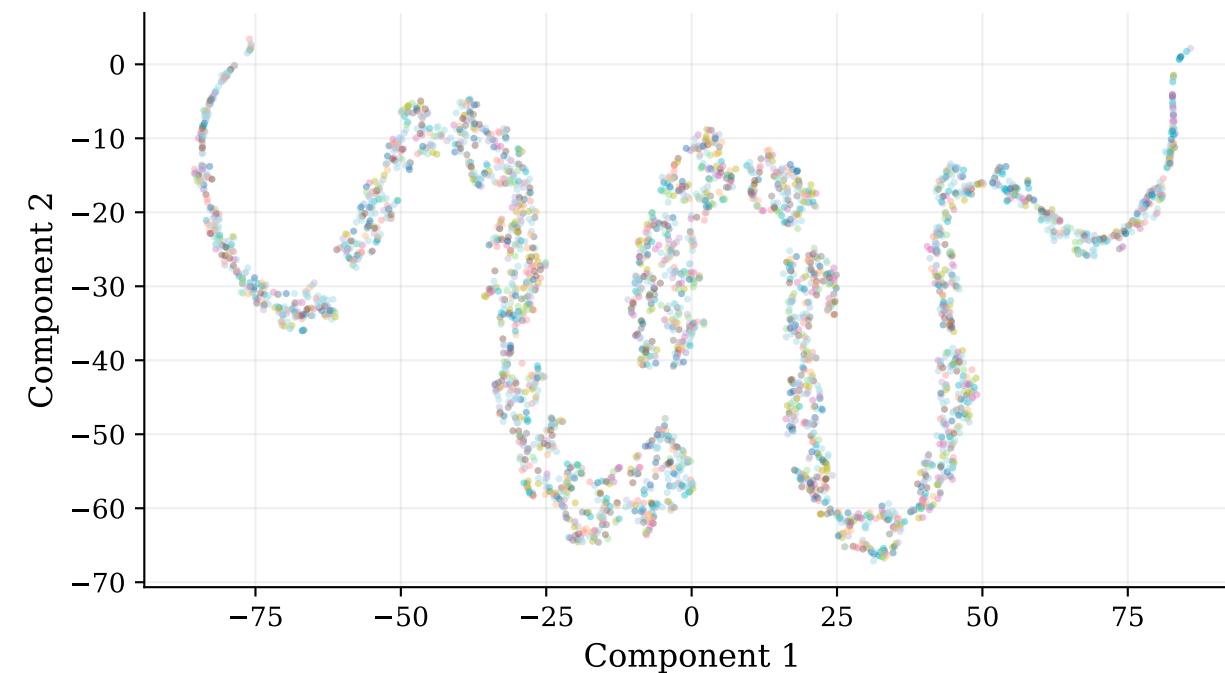
UMAP: ERM



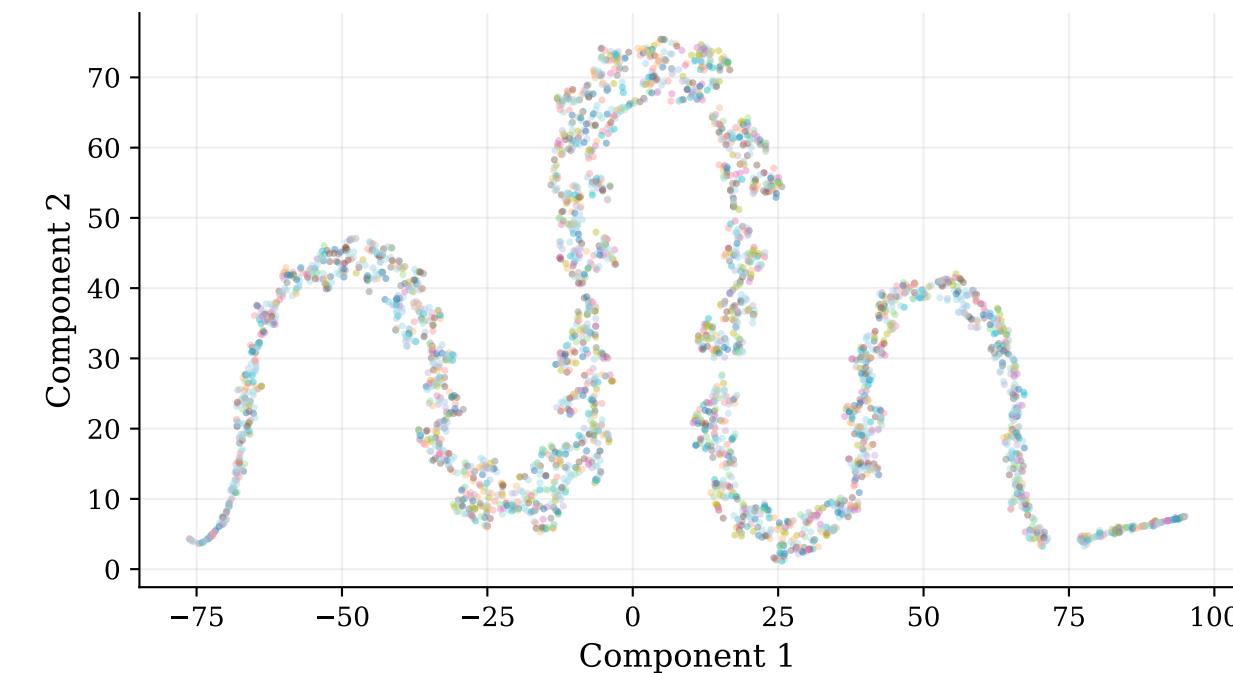
UMAP: DANN



t-SNE: ERM



t-SNE: DANN



CODEC=0	CODEC=2	CODEC=4	CODEC=6	CODEC=8	CODEC=10
CODEC=1	CODEC=3	CODEC=5	CODEC=7	CODEC=9	CODEC=11