

Integration Setup								Bio Conservation			Batch Correction					
Dataset	Method	Output Type	Batch	Label	Lineage Specific	Lineage Key	Lineage	Overall Score	Bio Conservation	Nmi	Ari	Asw Label	Batch Correction	Asw Batch	Ilisi	Pcr
tegration_merged_data	scanvi	embed	batch	harmonized_anno	global	None	global	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>
tegration_merged_data	scvi	embed	batch	harmonized_anno	global	None	global	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>
tegration_merged_data	unintegrated	embed	batch	harmonized_anno	global	None	global	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>
tegration_merged_data	unintegrated	full	batch	harmonized_anno	global	None	global	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>
tegration_merged_data	unintegrated	knn	batch	harmonized_anno	global	None	global	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>
tegration_merged_data	scanorama	embed	batch	harmonized_anno	global	None	global	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>
tegration_merged_data	harmony	embed	batch	harmonized_anno	global	None	global	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>
tegration_merged_data	scanorama	full	batch	harmonized_anno	global	None	global	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>	<div></div>

Score

00.20.40.8