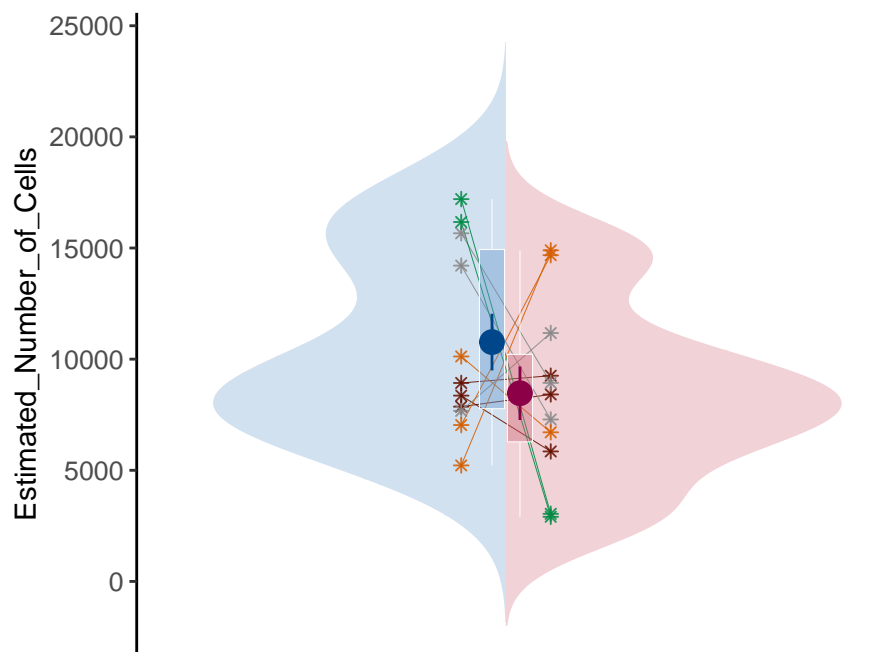


**Estimated\_Number\_of\_Cells**

Wilcoxon signed-rank test; p-value: 0.5195



Legends

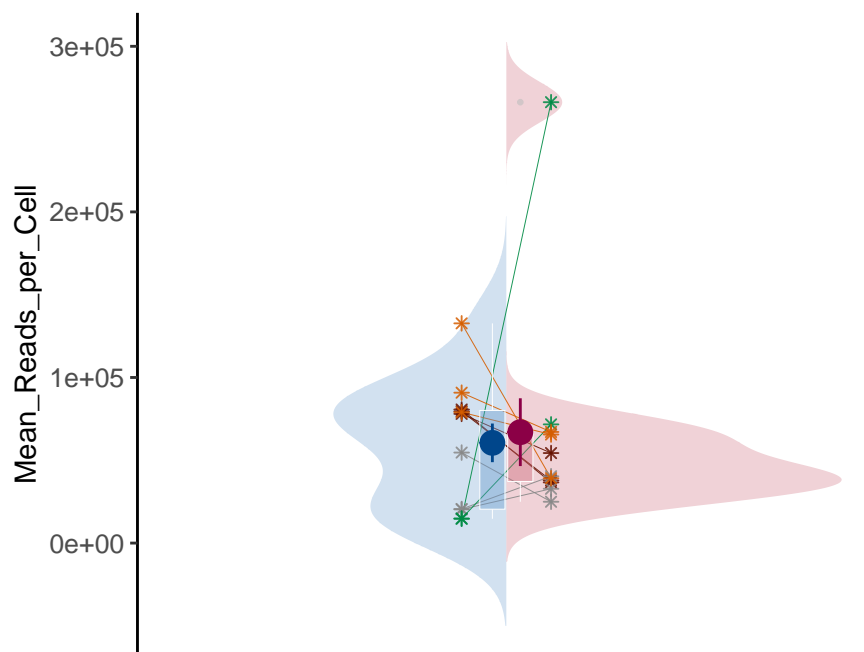
- Lab1
- Lab2
- Lab3
- Lab4
- scRNA-seq
- snRNA-seq

Platform

- Sc
- Sn

**Mean\_Reads\_per\_Cell**

Wilcoxon signed-rank test; p-value: 0.4648



Legends

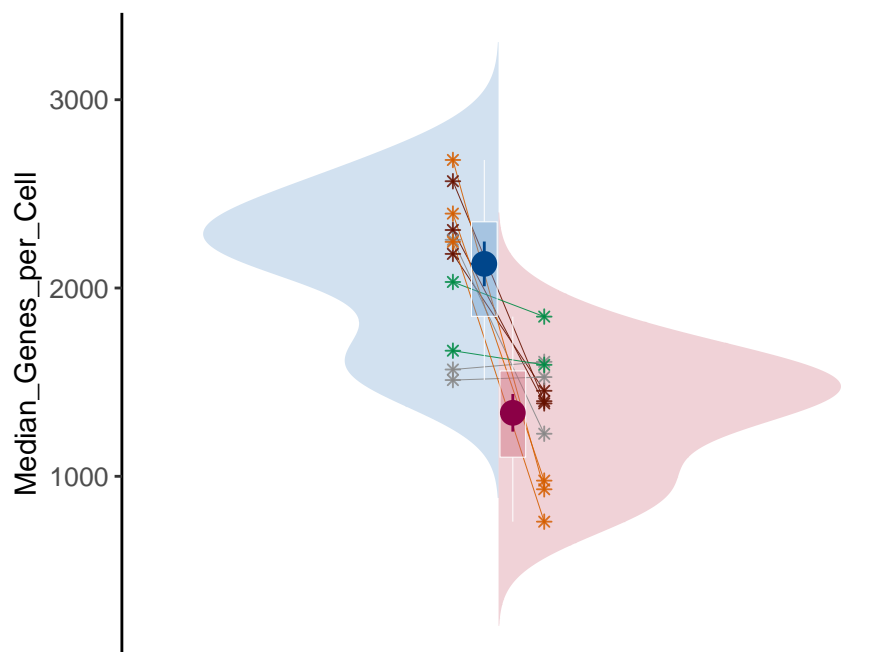
- Lab1
- Lab2
- Lab3
- Lab4
- scRNA-seq
- snRNA-seq

Platform

- Sc
- Sn

**Median\_Genes\_per\_Cell**

Wilcoxon signed-rank test; p-value: 0.0049



Legends

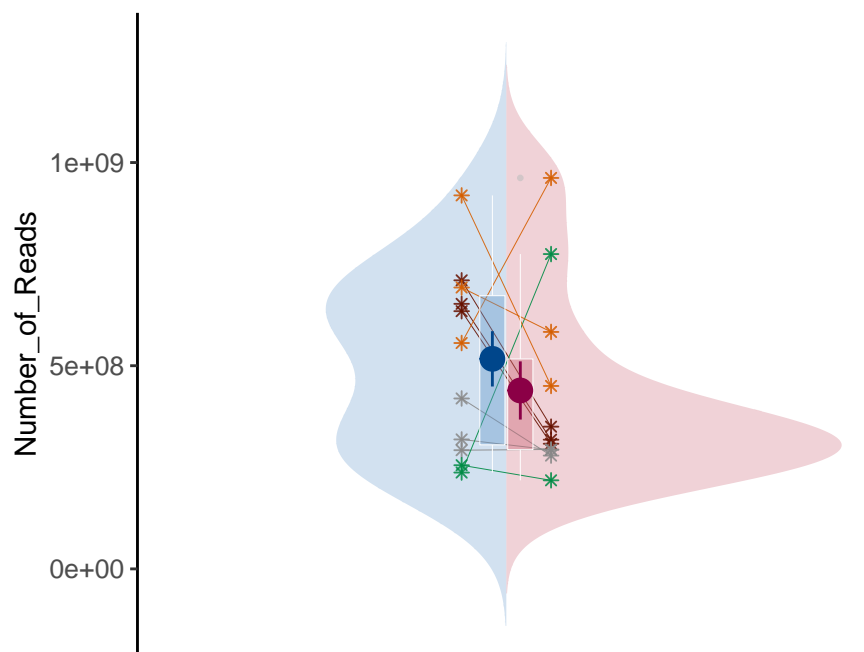
- Lab1
- Lab2
- Lab3
- Lab4
- scRNA-seq
- snRNA-seq

Platform

- Sc
- Sn

**Number\_of\_Reads**

Wilcoxon signed-rank test; p-value: 0.3203



Legends

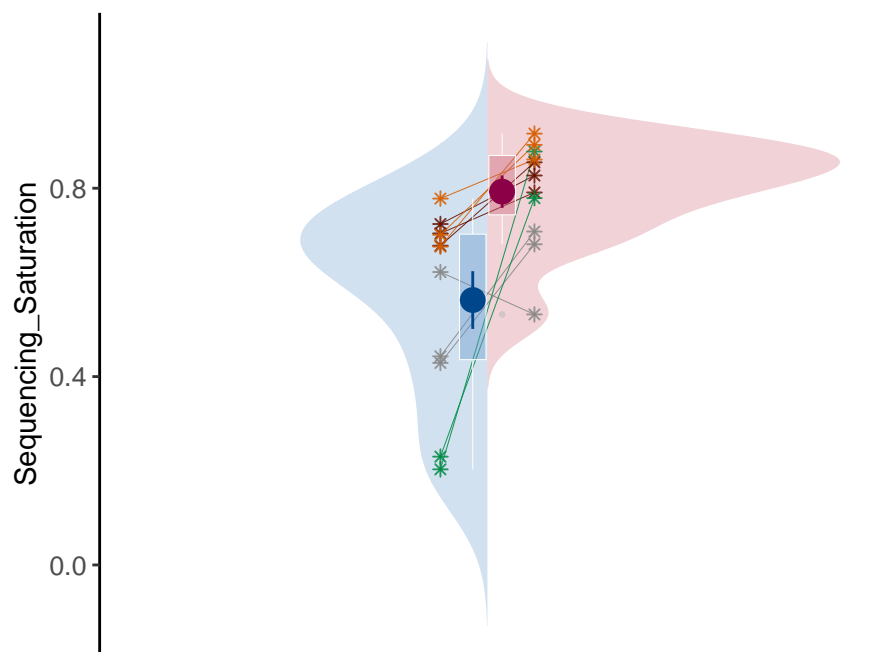
- Lab1
- Lab2
- Lab3
- Lab4
- scRNA-seq
- snRNA-seq

Platform

- Sc
- Sn

**Sequencing\_Saturation**

Wilcoxon signed-rank test; p-value: 0.0049



Legends

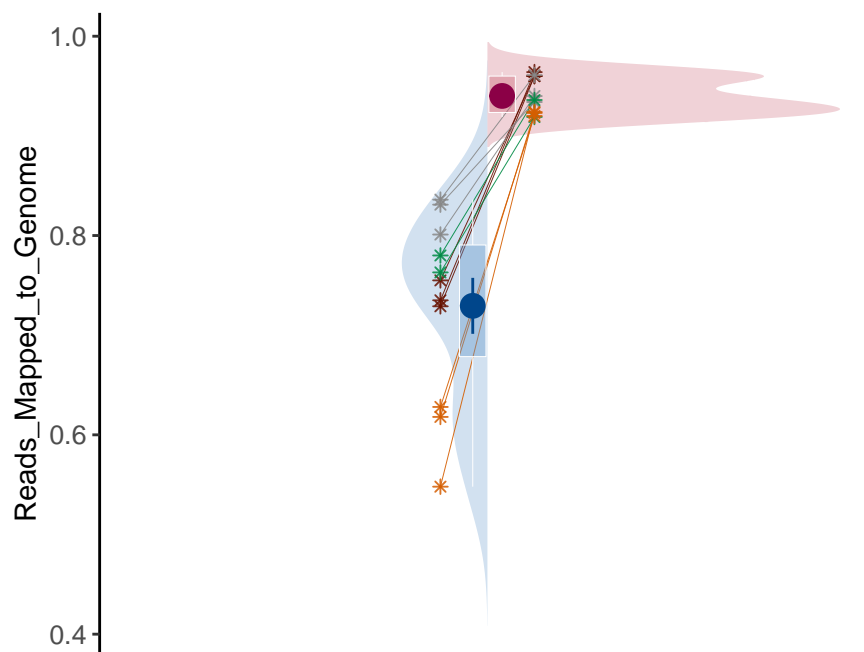
- Lab1
- Lab2
- Lab3
- Lab4
- scRNA-seq
- snRNA-seq

Platform

- Sc
- Sn

**Reads\_Mapped\_to\_Genome**

Wilcoxon signed-rank test; p-value: 0.0038



Legends

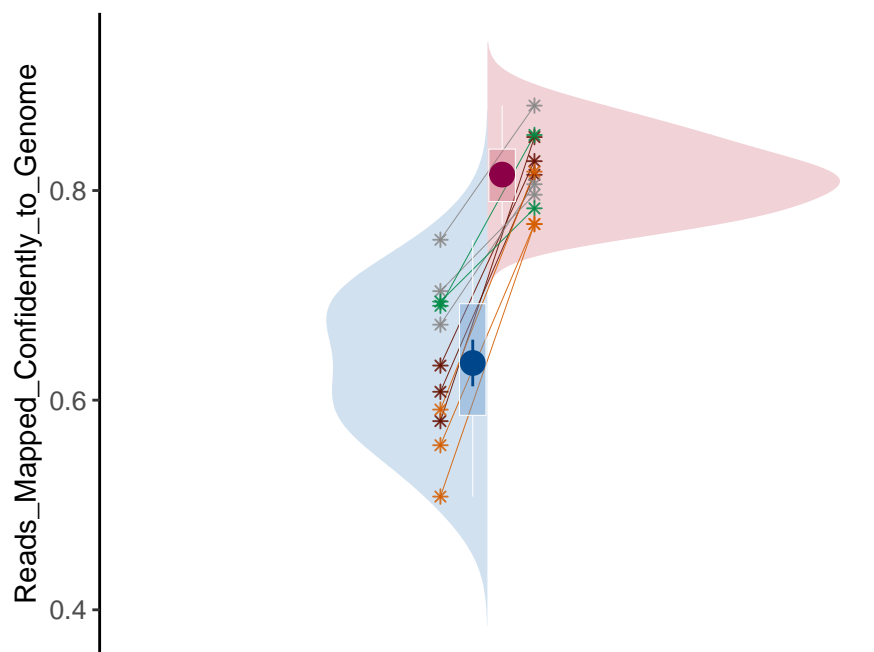
- Lab1
- Lab2
- Lab3
- Lab4
- scRNA-seq
- snRNA-seq

Platform

- Sc
- Sn

**Reads\_Mapped\_Confidently\_to\_Genome**

Wilcoxon signed-rank test; p-value: 0.001



Legends

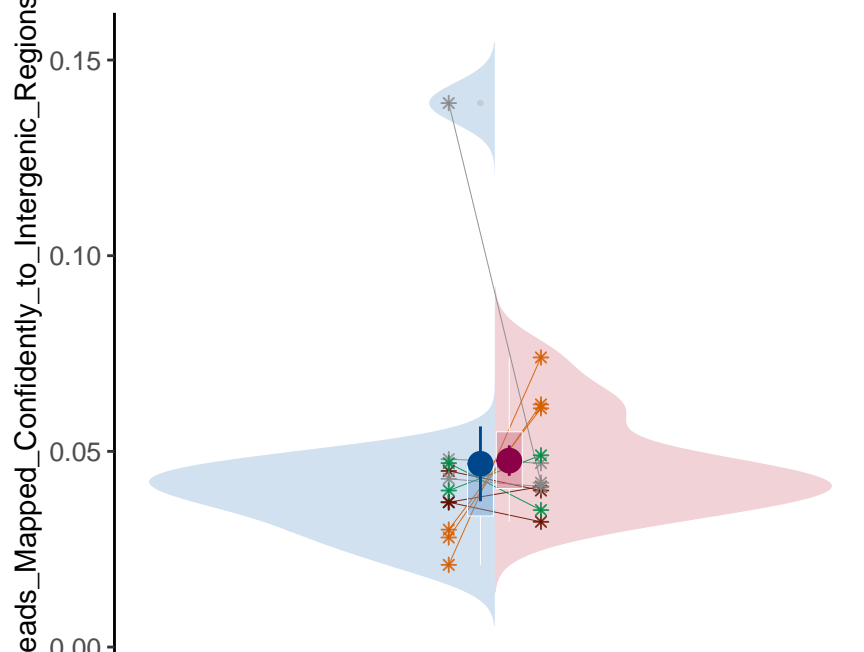
- Lab1
- Lab2
- Lab3
- Lab4
- scRNA-seq
- snRNA-seq

Platform

- Sc
- Sn

**Reads\_Mapped\_Confidently\_to\_Intergenic\_Regions**

Wilcoxon signed-rank test; p-value: 0.824



Legends

- Lab1
- Lab2
- Lab3
- Lab4
- scRNA-seq
- snRNA-seq

Platform

- Sc
- Sn

