Batch Adjustment

- ● None
- ○ Combat
- ○ SVA

No. of Genes to include in the table

50		22,283

1 2,230 4,459 6,688 9,177 11,146 13,375 15,604 17,833 20,062 22,283
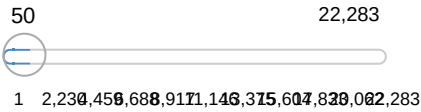
Confounding   Variation Analysis   P-value Analysis

Number of samples in each Batch and Condition

	Batch 1	Batch 2	Batch 3	Batch 4	Batch 5
Condition Biopsy	0	0	0	5	4
Condition Cancer	11	14	0	0	15
Condition Normal	0	4	4	0	0

Measures of confounding between Batch and Condition

	Standardized Pearson Correlation Coefficient	Cramer's V
Confounding Coefficients (0=no confounding, 1=complete confounding)	0.89	0.74