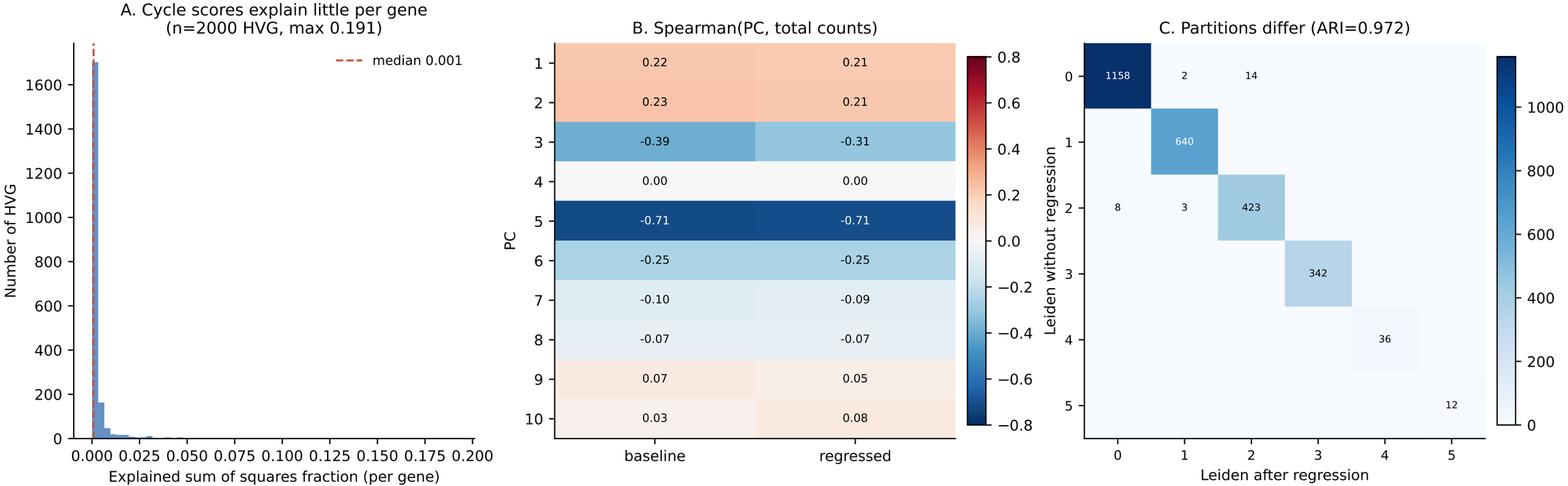
A. Cycle scores explain little per gene
(n=2000 HVG, max 0.191)
median 0.001
Number of HVG
Explained sum of squares fraction (per gene)
B. Spearman(PC, total counts)
PC
baseline
regressed
C. Partitions differ (ARI=0.972)
Leiden without regression
Leiden after regression