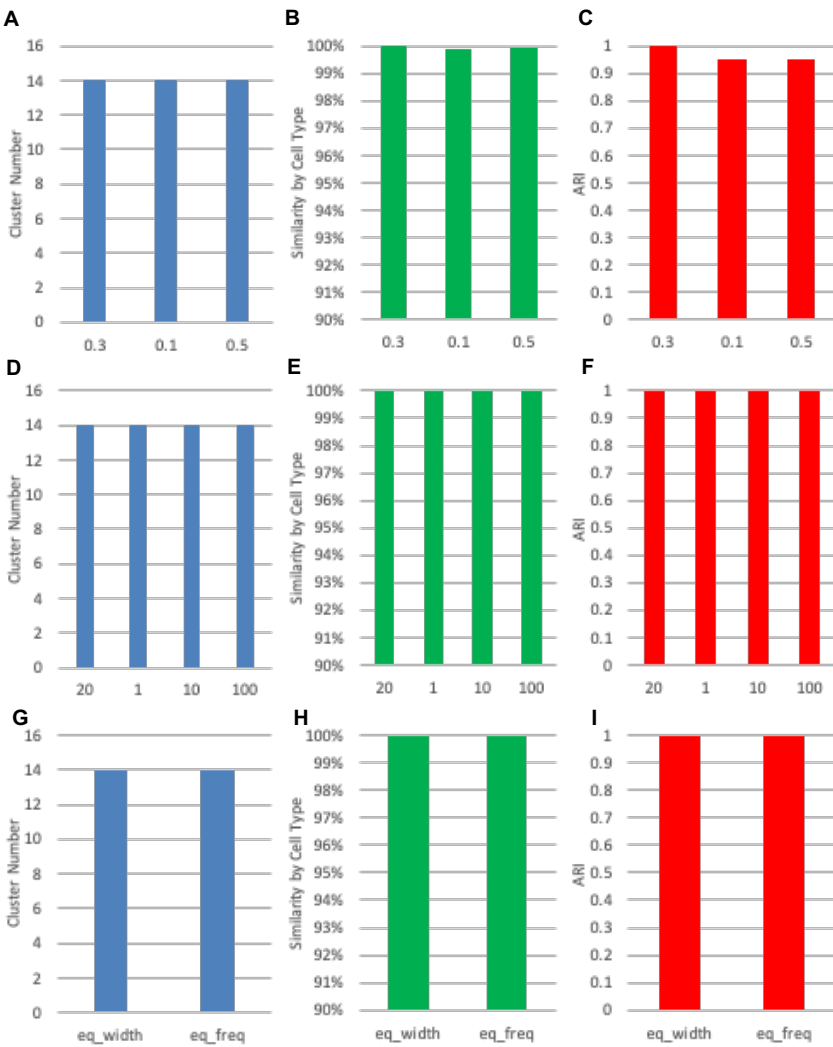


Figure S3



**Figure S3.** Cluster numbers (A, D, G), cell annotation similarity (B, E, H), and ARI values (C, F, I) when comparing loess spans of 0.3 (baseline) to 0.1 and 0.5 (A, B, C), or bin numbers of 20 (baseline) to 1, 10 and 100 (D, E, F), or using a binning method of equal width (baseline) to equal frequency (G, H, I).