



Supplemental Fig. S3 Boxplots of genotype error and pairwise SNV error for SCsnvcna without constraining SNV cells (dark blue), SCsnvcna with constraining SNV cells (light blue), COMPASS (dark red) and BiTSC² (yellow) when A. the number of SNV cells varies; B. the number of SNVs varies; and C. the number of subclones varies. The data was simulated by a simulator different from the one proposed in this study, and thus served as an orthogonal validation.