Supplemental Figure 1. A) Expression of NPC and proliferation markers, B) astrocyte marker, C) glial markers D) neuronal dendrite marker at weeks (W) 2, 4, 8, 10 and 12 of organoid differentiation. A-D) Data is represented as  $\log_2$  of the difference in expression, normalised to NPCs. *ACTB* was used as a reference gene. Data represents 2-3 independent experiments with 3 technical replicates each. E) Estimated brain region proportions of W8 organoids using multiple deconvolution methods (ordinary least squares, non-negative least squares regression, quadratic programming without constraints, quadratic programming non-negative and sum-to-one constraints, dtangle, robust linear regression, support vector regression) on bulk RNAseq data (4 biological replicates – T14-T17).

## Figure S1









