

Supporting information

Transcriptomic Mapping of Neural Diversity, Differentiation and Functional Trajectory in iPSC-Derived 3D Brain Organoid Models

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Table S1. Gene subsets governing specific neural functions.

| Axon Guidance | Axon Development | Neuronal Action Potential | Neuron Synaptic Transmission | NOTCH Signaling | WNT Signaling | Neuron Differentiation | BMP Signaling |
|-----------------|------------------|---------------------------|------------------------------|-----------------|---------------|------------------------|---------------|
| n | | | | | | | |
| ABL1 | ABL1 | ANK3 | DLGAP2 | ADAM17 | APC | ABI2 | ACVR2A |
| ABLIM1 | ACTB | CACNA1 | DRD2 | APH1A | APC2 | ABITRAM | ACVR2B |
| ABLIM2 | ACTBL2 | G | KIF1B | CIR1 | AXIN1 | ABL1 | ACVRL1 |
| ABLIM3 | ADARB1 | CACNA1 | PTEN | CREBBP | AXIN2 | ABL2 | AMH |
| ARHGEF12 | ADCY1 | H | TMOD2 | CTBP1 | BTRC | ACAP3 | AMHR2 |
| CDC42 | ADGRB1 | CACNA1I | VDAC1 | CTBP2 | CACYBP | ACP4 | BMP10 |
| CDK5 | AFG3L2 | CHRNA1 | | DLL1 | CAMK2A | ACSL3 | BMP2 |
| CFL1 | ALCAM | FGF12 | | DLL3 | CAMK2B | ACSL4 | BMPR1A |
| CFL2 | AMIGO1 | FMR1 | | DLL4 | CAMK2D | ACTB | BMPR1B |
| CFL2 | ANAPC2 | GBA | | DTX1 | CAMK2G | ACTBL2 | BMPR2 |
| CHP1 | ANK3 | GPER1 | | DTX2 | CCND1 | ACTL6B | CER1 |
| CHP2 | ANOS1 | GPR35 | | DTX3 | CCND2 | ACTR2 | CHRD1 |
| CXCL12 | APBB1 | GPR88 | | DTX3L | CCND3 | ADAM10 | FSTL1 |
| CXCR4 | APOA1 | GRIK2 | | DTX4 | CER1 | ADARB1 | GDF2 |
| DCC | APOD | KCNA1 | | DVL1 | CHD8 | ADCY1 | GREM2 |
| DPYSL2 | APOE | KCNA2 | | DVL2 | CHP1 | ADCY6 | NOG |
| DPYSL5 | APP | KCNND2 | | DVL3 | CHP2 | ADCYAP1 | SKI |
| EFNA1 | ARHGAP35 | KCNMB2 | | EP300 | CREBBP | ADGRB1 | SMAD1 |
| EFNA2 | ARHGAP4 | KCNMB3 | | HDAC1 | CSNK1A1 | ADGRB3 | SMAD4 |
| EFNA3 | ARHGDIA | KCNMB4 | | HDAC2 | CSNK1A1 | ADGRF1 | SMAD5 |
| EFNA4 | ARTN | MTNR1B | | HES1 | L | ADGRV1 | SMAD6 |
| EFNA5 | ARX | MYH14 | | HES5 | CSNK1E | ADM | SMAD7 |
| EFNB1 | ATL1 | P2RX1 | | JAG1 | CSNK2A1 | ADNP2 | SMAD9 |
| EFNB2 | ATOH1 | P2RX3 | | JAG2 | CSNK2A2 | ADORA2A | SMURF1 |
| EFNB3 | ATP8A2 | P2RX4 | | KAT2A | CSNK2B | ADRA2B | SMURF2 |
| EPHA1 | AUTS2 | SCN10A | | KAT2B | CTBP1 | ADRA2C | UBE2D1 |
| EPHA2 | B3GNT2 | SCN11A | | LFNG | CTBP2 | AFG3L2 | UBE2D3 |
| EPHA3 | B4GALT5 | SCN1A | | MAML1 | CTNNB1 | AGBL4 | ZFYVE16 |
| EPHA4 | B4GALT6 | SCN2A | | MAML2 | CTNNBIP | AGER | |
| EPHA5 | BAIAP2 | SCN3A | | MAML3 | 1 | AGRN | |

| | | | | | |
|-------|----------|-------|--------|--------|---------|
| EPHA6 | BARHL2 | SCN4A | MFNG | CUL1 | AGTPBP1 |
| EPHA7 | BCL11B | SCN5A | NCOR2 | CXXC4 | AHI1 |
| EPHA8 | BCL2 | SCN7A | NCSTN | DAAM1 | AKT1 |
| EPHB1 | BDNF | SCN8A | NOTCH1 | DAAM2 | ALCAM |
| EPHB2 | BMP7 | SCN9A | NOTCH2 | DKK1 | ALDH1A2 |
| EPHB3 | BMPR1B | | NOTCH3 | DKK2 | ALK |
| EPHB4 | BMPR2 | | NOTCH4 | DKK4 | ALKAL1 |
| EPHB6 | BOC | | NUMB | DVL1 | ALKAL2 |
| FES | BRSK1 | | NUMBL | DVL2 | ALKBH1 |
| FYN | BRSK2 | | PSEN1 | DVL3 | ALS2 |
| GNAI1 | BSG | | PSEN2 | EP300 | AMIGO1 |
| GNAI2 | C12orf57 | | PSENEN | FBXW11 | ANAPC2 |
| GNAI3 | C9orf72 | | PTCRA | FOSL1 | ANK3 |
| GSK3B | CAMSAP2 | | RBPJ | FRAT1 | ANKRD1 |
| HRAS | CASP3 | | | FRAT2 | ANKRD27 |
| ITGB1 | CCK | | | FZD1 | ANKS1A |
| KRAS | CCKAR | | | FZD10 | ANOS1 |
| L1CAM | CDH11 | | | FZD2 | AP2A1 |
| LIMK1 | CDH2 | | | FZD3 | APBB1 |
| LIMK2 | | | | FZD4 | |
| | | | | FZD5 | |

Table S2. Comparison between the sample properties of the datasets used in this study.

| scRNA-seq data | Reference | Identifier | No. of cells | No. of Batches | Differentiation induction period (days) | Neural maturation start day | Notable steps/signaling pathways |
|--------------------|-----------------------|------------|--------------|----------------|---|-----------------------------|----------------------------------|
| Human Brain | Trujillo et al., 2019 | GSE130238 | 3,480 | 1 | 10 | 17 | Inhibition of BMP, TGFβ |
| Organoids | Velasco et al. 2019 | GSE129519 | 99,760 | 15 | 18 | 35 | Inhibition of WNT, TGFβ |

| | | | | | | | |
|------------------------------|-----------------------|------------|---------------------------------|-------------------|------------|------|---|
| | | | | | | | Matrigel coating |
| Giandomenico et al., 2019 | GSE124174 | 12,824 | 1 | 11 | 20 | | Matrigel coating |
| Fiddes et al., 2018 | GSE106245 | 13,747 | 2 | 18 | 24 | | Inhibition of SHH, WNT, BMP, TGFβ |
| Madhavan et al., 2018 | GSE110006 | 3,743 | 1 | 6 | 25 | | Inhibition of BMP, TGFβ |
| Quadrato et al., 2017 | GSE86153 | 43,706 | 26 | 4 | 8 | | Matrigel coating |
| Birey et al., 2017 | GSE93811, GSE96045 | 4,953 | 1 | 6 | 25 | | Inhibition of BMP, TGFβ |
| Xiang et al., 2017 | GSE97882 | 23,707 | 2 | 10 | 18 | | Inhibition of WNT, BMP, TGFβ |
| Fetal Brain | Reference | Identifier | No. of cells | No. of Batches | Cell count | | |
| | Zhong et al. 2020 | GSE104276 | 2,394 | 101 | Female | Male | |
| | | | Early stage (week 8-13) | | 278 | 136 | |
| | | | Mid-early stage (week 16-19) | | 889 | 0 | |
| | | | Middle stage (week 23-26) | | 928 | 143 | |

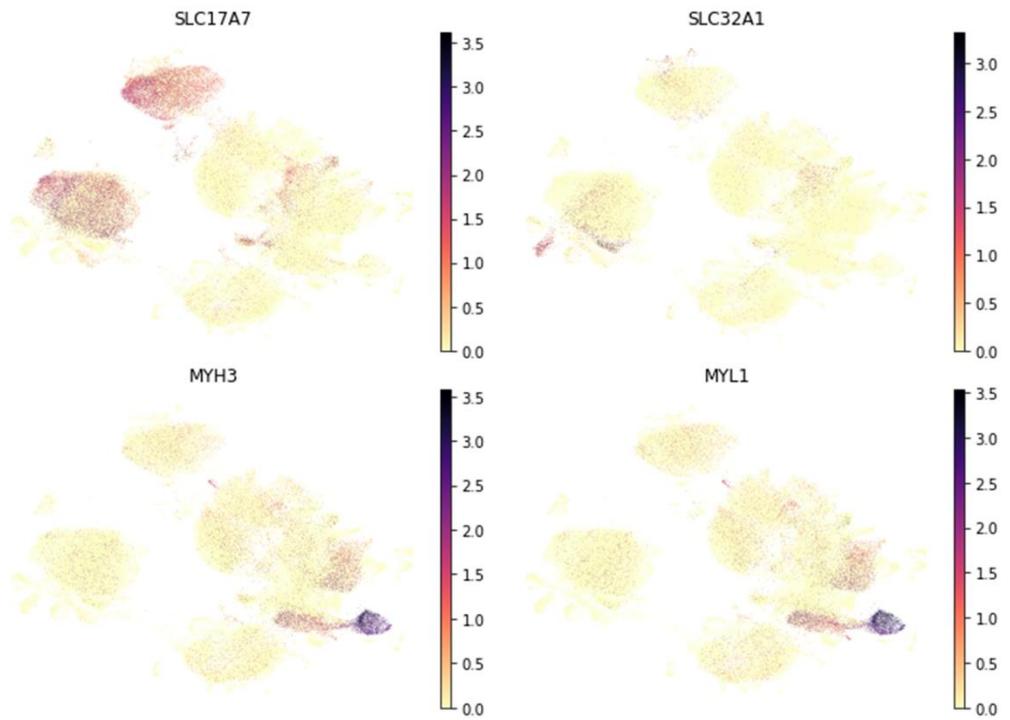


Figure S1. Visualization of various genes (top: Glutamatergic, bottom: GABAergic) on the integrated UMAP from organoid and fetal brain samples.

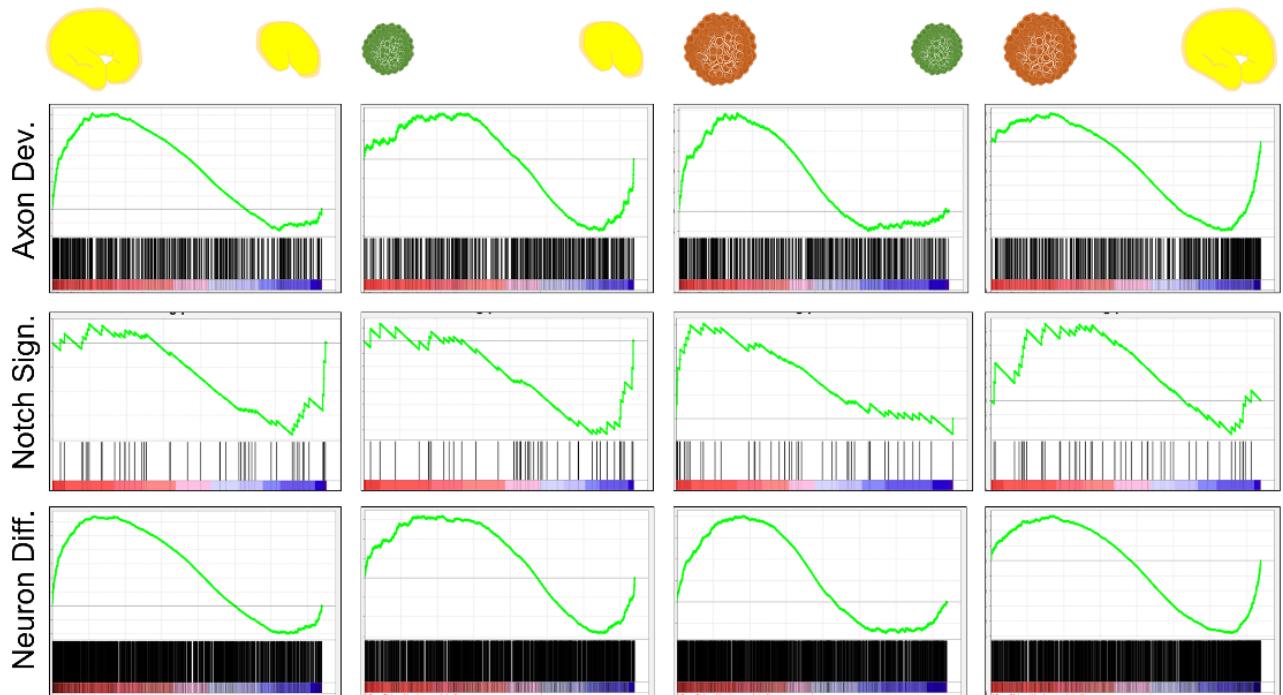


Figure S2. GSEA analysis of developmental genesets (axon development, notch signaling and neuron differentiation) between organoid and fetal groups. From left to right: fetal 6-month vs 3-month, organoid 3-month vs fetal 3-month, organoid 6-month vs organoid 3-month, organoid 6-month vs fetal 6-month.

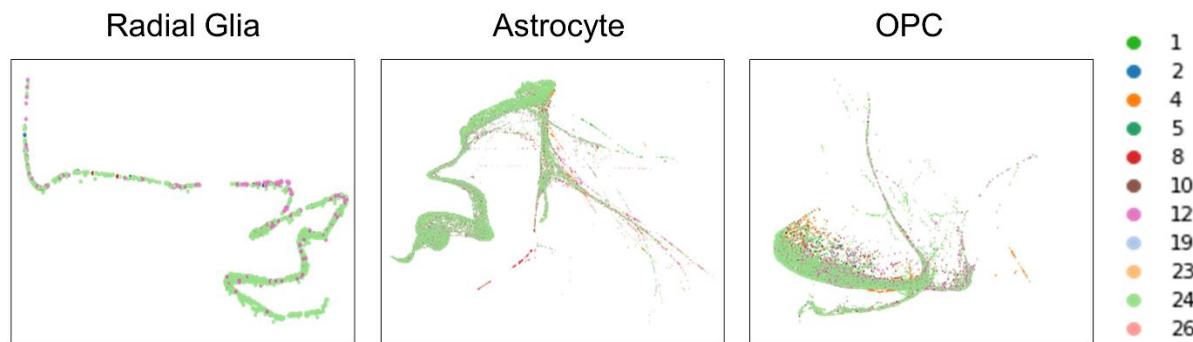


Figure S3. Cell type specific developmental trajectory in radial glia, astrocytes and OPCs across timepoints of 1-26 weeks.

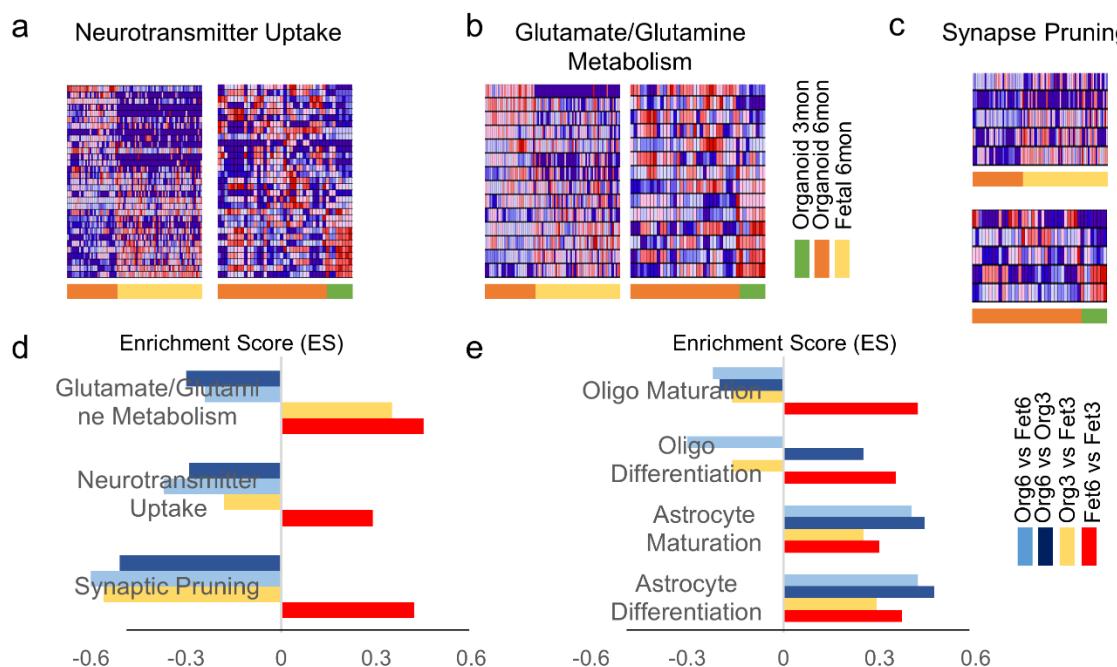


Figure S4. Gene set enrichment analysis of functional pathways (a-d), differentiation and maturation (e) relevant to astrocytes in organoid cultures and fetal brain at 3-month and 6-month timepoints. The differentiation gene set in (e) refers to the transformation process of relatively unspecialized cells to astrocytes [oligodendrocytes]. The maturation

gene set refers to the progression of astrocytes [oligodendrocytes] from initial commitment to the astrocyte [oligodendrocyte] fate to a fully functional differentiated cell.