



Additional file 6. Overlap between differentially expressed genes from our study on human blood RNA-seq of 15q13.3 microdeletion subjects (Human), mouse (Df[h15q13]/+) cerebral cortex reported by Gordon *et al.* 2021 (Gordon's Mouse), mouse (Df[h15q13]/+) medial prefrontal cortex reported by Al-Absi *et al.* 2021 (Al-Absi's Mouse), and induced pluripotent stem cells (Zhang's iPSCs), as well as induced neurons (Zhang's iNs) from Zhang *et al.* 2021. There is a significant overlap between the human blood RNA-seq and differentially expressed mouse genes data (p -value < 0.001) and the human blood RNA-seq data and induced neurons (p -value < 0.001). However, the overlap between the iPSCs and iNs and the two mouse models is not significant.