## **Description of Additional Supplementary Files**

File Name: Supplementary Data 1

Description: Detailed clinico-pathological and sequencing result descriptions of AD and non-

AD subjects

File Name: Supplementary Data 2

Description: Detailed information on deep whole exome sequencing results of non-SCZ and

SCZ subjects

File Name: Supplementary Data 3

Description: Targeted amplicon sequencing results for determining the cut-off value of

**EBscore** 

File Name: Supplementary Data 4

Description: Targeted amplicon sequencing results and related primer sequences for

randomly selected brain somatic mutations

File Name: Supplementary Data 5

Description: List of post-filtered brain and blood somatic single nucleotide variations in AD

and non-AD

File Name: Supplementary Data 6

Description: PCAWG signatures adjusted to human whole-exome trinucleotide frequencies

File Name: Supplementary Data 7

Description: Detailed summary report of mutation signature analyses of AD brain and AD

blood from Mutalisk tool

File Name: Supplementary Data 8

Description: List of putatively pathogenic brain and blood somatic single nucleotide

variations in AD and non-AD

File Name: Supplementary Data 9

Description: List of brain somatic mutations in AD with CADD Phred score and the number of

evidence sentences from the DigSee engine

File Name: Supplementary Data 10

Description: Targeted amplicon sequencing results of a PIN1 (c.477 C>T) mutation from bulk, sub-region, and AT8-positive/negative entorhinal cortical neurons of the hippocampal

formation

File Name: Supplementary Data 11

Description: List of pathogenic germline risk factors and Tau-pathology affecting putatively

pathogenic somatic mutations in AD and non-AD

File Name: Supplementary Data 12

Description: Variant allelic frequencies, mutation context, and predicted  $\Delta\Delta G$  of AD

brain somatic mutations enriched in PI3K-AKT, MAPK, and AMPK pathways

File Name: Supplementary Data 13

Description: List of read counts of 22 variants from Wei et. al<sup>1</sup> (Genetics in Medicine, 2018) and the nine variants from Nicolas et. al<sup>2</sup> (Alzheimer's & Dementia, 2018) in our cohorts

File Name: Supplementary Movie 1

Description: Demonstration of laser capture microdissection (LCM) to isolate Nissl-

stained sections of the dentate gyrus