Supplemental Figure 1 (a-b). (a) Quantile-Quantile plots of the GWAS of hippocampal volume results for each individual study from the ENIGMA Consortium (split into two panels a and b). Lambda inflation factors are provided for each plot. The red line represents the expected null distribution.
Supplemental Figure 1 (a-b). (b) Quantile-Quantile plots of the GWAS of hippocampal volume results for each individual study from the ENIGMA Consortium (split into two panels a and b). Lambda inflation factors are provided for each plot. The red line represents the expected null distribution.
Supplemental Figure 2. (b) Quantile-Quantile plots of the GWAS of hippocampal volume results for each individual study from the CHARGE Consortium. Lambda inflation factors are provided for each plot. The red line represents the expected null distribution.
Supplemental Figure 3. LDSCORE regression analysis split by chromosome. Plotted values are the proportion of $h^2_g$ explained divided by the proportion of SNPs in a given chromosome. Values are significantly over- or under-represented if they differ significantly from 1. Values are plotted with a standard error calculated with a jackknife in LDSCORE. Chromosome 4 had a significant under-representation in its contribution to the overall heritability estimate (indicated by **).
Supplementary Note 1: Consortium Authors:

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ADNI Methods:

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The Principal Investigator of this initiative is Michael W. Weiner, MD, VA Medical Center and
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Supplementary Note: Author Contributions
