

Effect of Age and APOE on Hippocampal Subfields in Cognitively Normal Subjects (N=1168, single site)

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INTRODUCTION

- **Hippocampal atrophy** is associated with memory deficits and is used to monitor disease progression such as **Alzheimer’s Disease (AD)** [1].
- **Age** and **APOE ε4 allele** (major genetic risk for sporadic AD) have a strong impact on hippocampal volume in cognitively normal subjects [2].
- **Aim::** in this study, we estimated **hippocampal subfield volumes** in **cognitively normal participants** and looked for associations with age, sex, education and APOE.

METHODS

- **N=1168** cognitively healthy participants aged between 45 and 74 years were selected from the ALFA study, which is a large cohort program for neuroimaging biomarkers of preclinical AD in the general population [3]. Acquired data covers **multimodal MRI, DNA extraction, CSF and blood sampling**, full cognitive and lifestyle habits evaluation.

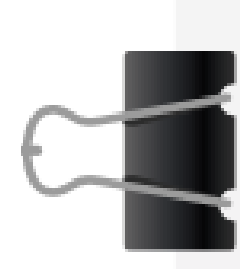
	Total sample (N=1168)	Non-carriers (N=576)	ε4-heterozygous (N=417)	ε4-homozygous (N=75)	Inferential Statistics
Age (in years) (SD)	59.41 (6.82)	59.23 (7.00)	60.06 (6.55)	57.57 (5.98)	F=4.84 p=0.008
Education (in years) (SD)	13.43 (3.49)	13.33 (3.47)	13.82 (3.51)	13.19 (3.44)	F=1.02 p=0.36
Males / Females	412 / 756	240 / 438	149 / 268	23 / 50	χ²=0.5 p=0.78

- **Single MR scanner:** Philips Ingenta CX 3T (3DT1, 0.75-mm isotropic voxels, TR=9.9ms, TE=4.6ms, TI=600ms, Flip Angle=8°, sagittal plane, FOV 320x320 mm, imaging matrix 240x240x180).

- Segmentation of subfields was achieved using **FreeSurfer 6.0** [4].

- **Statistical analysis:** we performed a group ROI-based analysis and assessed the effect of APOE on each subfield and its interaction with age by fitting a general linear model. Age was modeled as a quadratic term centered at age 55. We partitioned genetic variance using 3 dummy regressors coding for the number of ε4 alleles carried. Sex, education and total intracranial volume were included as confounding variables. Results shown at p<0.05 corrected for multiple comparisons.

- Both raw and processed data are stored on an **XNAT platform**. Processing workflows were implemented and run on that platform. Statistical analysis was performed using Python (*statsmodels* 0.9).



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RESULTS

- **APOE ε4 carriers** showed **reduced volumes** as compared to non-carriers (**Figure 1**).
 - Bilateral effect of APOE found in CA3, CA4, dentate gyrus and hippocampal tail and whole hippocampus. Effects found in both recessive and additive contrasts(*).
 - Only fimbria showed **larger volumes** in ε4 carriers, also in both recessive and additive contrasts(*).
 - Left hippocampal fissure is found **larger** in **ε4 carriers** in the dominant contrast(*).
 - All subfields **decrease in volume with age**, except right hippocampal fissure (p=0.8).
 - **Strongest effect** of age was found, in decreasing order of magnitude, in whole hippocampus, hippocampal tail, molecular layer, CA1, presubiculum, subiculum, dentate gyrus, CA4, fimbria, CA3, HATA, left hippocampal fissure and parasubiculum (**Figure 2**).
- Pairwise comparisons revealed **stronger age-related atrophy** in:
- hippocampal tail > CA1, CA3, CA4, fimbria, parasubiculum, hippocampal fissure, HATA and subiculum
 - molecular layer > CA3, fimbria, parasubiculum, hippocampal fissure and HATA
 - presubiculum and CA1 > parasubiculum

(*) Contrasts: **recessive** (ε4-homozygotes vs others), **dominance** (ε4 carriers vs others), **additive** (the more ε4 alleles the stronger)

CONCLUSIONS

- Highlights the impact of both **age** and **APOE ε4 status on hippocampal subfields in cognitively normal participants**. Both factors show a differential effect across subfields.
- With a uniquely **high number of homozygotes**, this study shows that carrying **two copies of the ε4 allele** is also associated with a significantly higher impact.

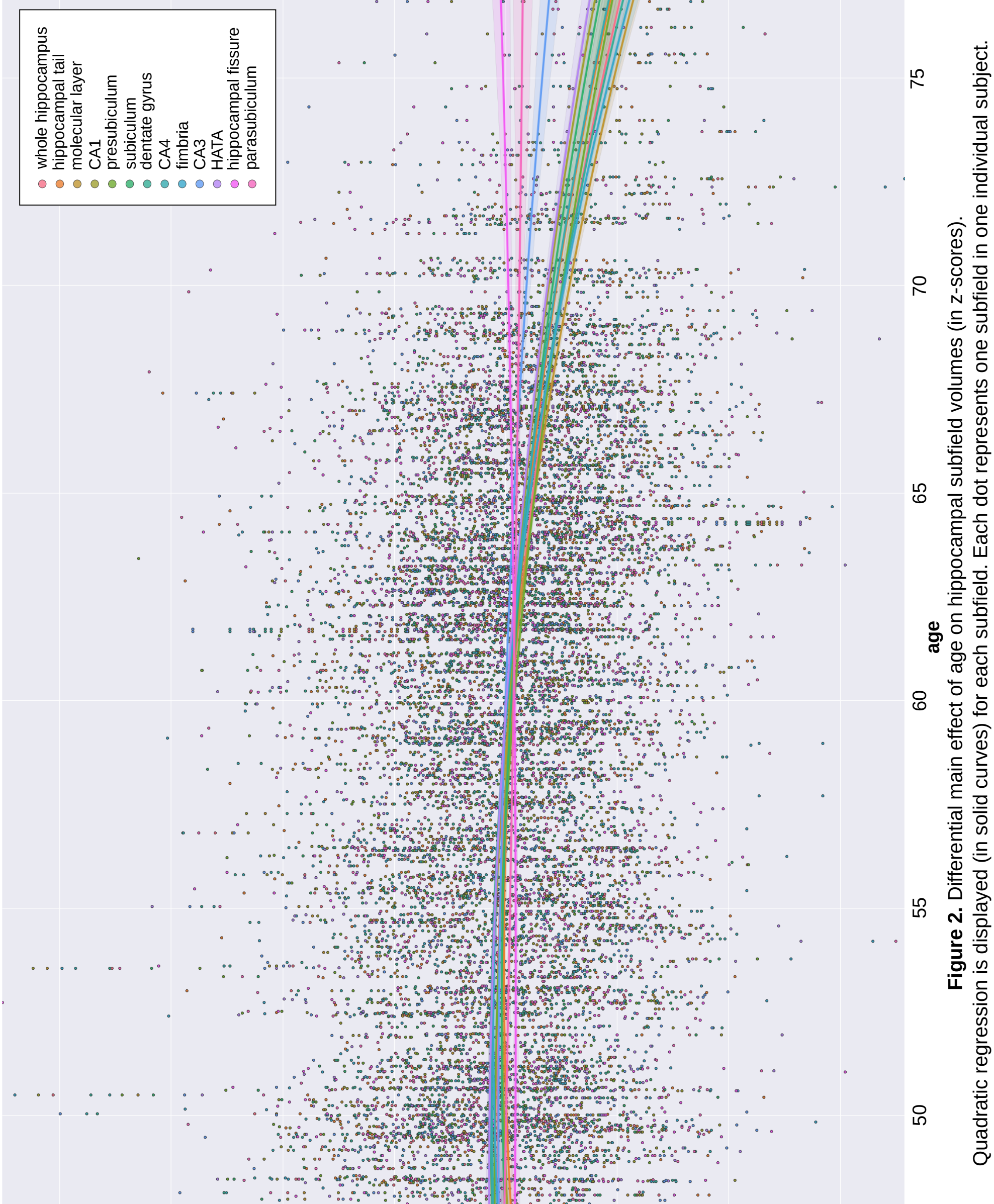
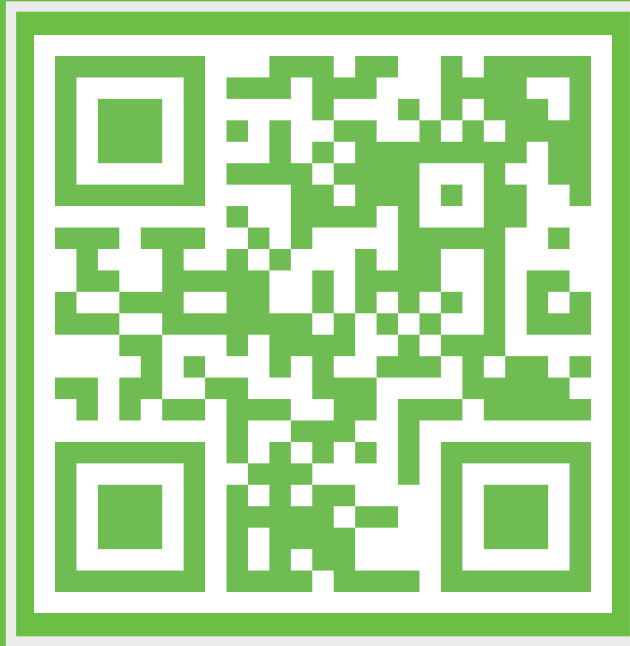


Figure 2. Differential main effect of age on hippocampal subfield volumes (in z-scores). Quadratic regression is displayed (in solid curves) for each subfield. Each dot represents one subfield in one individual subject.

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References: [1] Chételat, G. (2018). Multimodal Neuroimaging in Alzheimer’s Disease: Early Diagnosis, Physiopathological Mechanisms, and Impact of Lifestyle. Journal of Alzheimer’s Disease. [2] Cacciaglia, R. (2018) Effects of APOE-ε4 allele load on brain morphology in a cohort of middle-aged healthy individuals with enriched genetic risk for Alzheimer’s disease. Alzheimer’s & Dementia. [3] Molinuevo, J. L. et al. (2016) The ALFA project: A research platform to identify early pathophysiological features of Alzheimer’s disease. Alzheimer’s & Dementia. [4] Iglecias, J. E., et al. (2015) A computational atlas of the hippocampal formation using ex vivo, ultra-high resolution MRI: Application to adaptive segmentation of in vivo MRI. NeuroImage

Both age and the APOE ε4 allele have a negative differential impact on hippocampal subfields in cognitively normal subjects. ε4-homozygotes show heavier burden than heterozygotes and non-carriers.



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