



The ALFA Study and nested Sub-studies

Synopsis

The ALFA (ALzheimer and FAmilies) study¹ is a large-scale research initiative aimed at understanding the **early pathophysiological changes and risk factors associated with preclinical Alzheimer's disease (AD)**. The ALFA parent cohort comprises nearly **3,000 cognitively unimpaired individuals**, mostly first-degree descendants of AD patients, aged 45–75, thoroughly characterized through sociodemographic, clinical, blood biomarker, lifestyle, genetic, and cognitive assessments. ALFA is enriched for genetic risk of developing cognitive impairment, particularly a higher prevalence of the *APOE-\varepsilon4* allele, is found. ALFA is a highly valuable study to pursue AD prevention research.

Nested within ALFA are two major sub-studies: ALFA-MRI and ALFA+. The ALFA-MRI study (n \approx 1,600) expanded phenotyping via advanced magnetic resonance imaging (MRI), olfactory testing, and biological aging markers such as telomere length. The ALFA+ study (n \approx 420), a longitudinal cohort study, includes, aside from all the variables collected in ALFA, in-depth phenotyping with fluid (both cerebrospinal and plasma) and imaging biomarkers (MRI and several modalities of positron emission tomography), and repeated assessments over time (every 3 years). A significant portion of cognitively unimpaired participants (35-40%) already exhibit preclinical AD biomarker profiles (e.g., A+T-, A+T+). These efforts are pivotal in modelling AD's preclinical stages, identifying early biomarkers, and facilitating preventive clinical trials.

_

¹ The ALFA Study was established in 2013 by the Pasqual Maragall Foundation's research centre, the Barcelonaβeta Brain Research Centre (BBRC), and supported by "la Caixa Foundation".





The ALFA Study

The setup of preventive strategies requires the understanding, from a molecular perspective, of how risk factors generate the risk and the identification of individuals with an increased risk of developing cognitive impairment in the near future that are suitable to be recruited as asymptomatic subjects in prevention studies and trials. With this in mind, and aiming at increasing our knowledge of the pathophysiology and pathogenic factors emerging at early preclinical Alzheimer's disease (AD) stages, the Barcelonaßeta Brain Research Centre (BBRC), the research centre of the Pasqual Maragall Foundation, started the ALFA (for ALzheimer and FAmilies) study, supported by "la Caixa Foundation.

The ALFA study (also referred to as ALFA parent cohort) was set-up for the prospective followup of a cohort of cognitively unimpaired individuals that were recruited between 2013 and 2014. ALFA is composed of 2,743 cognitively unimpaired participants, most of them firstdegree descendants of AD patients, aged between 45 and 75 years, who were thoroughly characterised from a sociodemographic, clinical, lifestyle, genetic and cognitive point of view. Other variables of interest obtained during the baseline visit were those lifestyles and cardiovascular risk factors that had been previously suggested as modifiable risk factors that may increase or decrease the risk of cognitive impairment and dementia such as cardiovascular and endocrine-metabolic co-morbidities, the participants' level of physical activity and their smoking habits. In addition, participants' APOE (Apolipoprotein E) haplotype also has been determined as well as a whole **GWAS** is also available (Illumina Infinium Neuro Consortium [NeuroChip] Array). Finally, a subset of ~600 participants that were selected based on their APOE genotype (preferentially including APOE- $\varepsilon 4$ allele carriers), underwent cerebral magnetic resonance imaging (MRI, example). In addition, a number of plasma biomarkers have been more recently determined from samples that were obtained in the ALFA study visit. These include Aβ42, Aβ40, pTau181, pTau231, NfL and GFAP.

The family history of AD of ALFA participants was recorded during baseline visit. In particular, we registered who, their mother and/or father, had been diagnosed with cognitive impairment. In this regard, 86.3% of the study participants had at least one of their parents that had suffered AD. When considering a more strict family history encoding, it is remarkable that 47.4% of the ALFA study participants had at least one of their parents that had been diagnosed with AD before the age of 75. Family studies have shown that having a parental history of AD represents a risk factor for sporadic AD and the biggest genetic susceptibility factor is the *APOE-ε4* allele. In agreement with this, a higher frequency of the *APOE-ε4* allele was found in ALFA parent cohort participants than in the general population (38% and 14%, respectively; *P* <.001). In brief, of 2,714 ALFA members whose genotype could be determined, 9 were *APOE-ε2/ε2* homozygotes, 171 were *APOE-ε2/ε3* heterozygotes, 60 were *APOE-ε2/ε4* heterozygotes and, finally, 89 were *APOE-ε4/ε4* homozygotes.

Therefore, we have **established** a **research platform that is enriched in genetic risk factors for AD**. As a consequence, the proportion of individuals presenting altered biomarkers, neuroimaging changes and eventually the development of cognitive decline is also expected to be higher, which is being evaluated in longitudinal assessments. In summary, the ALFA parent cohort represents a valuable infrastructure of middle-aged participants representing the **whole spectrum of risk that will leverage with different studies and trials to prevent AD**.

The main characteristics of ALFA study's participants are: mean age 56.4 years, 63.2% females, 34,3% *APOE-ε4* carriers, mean years of education 13.4.





The ALFA-MRI Study

As a nested study to ALFA, the BBRC established the cross-sectional <u>ALFA-MRI study</u>, with the main objective of expanding the clinical, lifestyle, cognitive and brain characterisation of a subset of cognitively unimpaired ALFA study participants. Around **1,600 participants with no contraindications to MRI** that were selected based on their AD risk profile were recruited from 2016 to 2019.

On top of a review of the clinical and cognitive status, participants underwent a high-resolution MRI acquisition protocol in our centre's Philips Ingenia CX 3T including several sequences (T1, T2, FLAIR, DWI, IR and resting state fMRI). In addition, the study visits also included an odour identification test, a blood extraction to determine basic biochemical variables and be kept for future analyses (e.g. plasma samples to determine AD-related biomarkers). Finally, the telomere length (TL), as a proxy of biological age, has been also determined in ALFA-MRI study participants, which allows us to assess whether TL may be generating an age-related structural and functional vulnerability and mediating the effect of ageing on AD pathology.

The main characteristics of ALFA-MRI study's participants are: mean age 60.1 years, 61.2% females, 38,1% *APOE-ε4* carriers, mean years of education 13.5.

The ALFA+ Study

Alfa+ study in which a more detailed phenotyping is performed. ALFA+ is a prospective and observational cohort study for the early identification of biomarkers (both fluid and neuroimaging) of AD in ~420 cognitively unimpaired individuals. Participants with no contraindications to MRI or lumbar puncture (LP) were invited based on their risk profile (mainly APOE and family history status). The aim of the study is to describe the biological processes and identify factors that may precede the clinical phase of AD. Likewise, thanks to the extensive characterisation of its participants, ALFA+ aims to analyse the association between the biological, structural, functional and neurocognitive brain markers that characterize the preclinical phase of the disease and its natural history. The baseline visit (V1) of the ALFA+ cohort study took place between 2016 and 2019 and follow-up visits take place every 3 years: The first follow-up visit (V2) started in 2019 and finished by the end of 2022, and V3 started by 2023 and is currently underway (expected to finalise by 2026).

Each ALFA+ cohort study visit is organised in three core sessions: Session 1 (S1) includes a clinical, cognitive, nursing and lifestyle characterisation of study participants as well as a high-resolution MRI acquisition protocol in our centre's Philips Ingenia CX 3TMRI session (including T1, T2, FLAIR, DWI, IR and resting state fMRI sequences). In S2, participants undergo a LP to obtain cerebrospinal fluid (CSF) for their biomarker characterisation and obtaining several CSF aliquots that can be used for future biomarker studies. Various types of biological samples are also collected in S1 (non-fasting conditions) and S2 (fasting conditions) to determine biochemical variables and be kept for future analyses (e.g. plasma samples to determine AD-related biomarkers). Finally, S3 entails a further, more experimental and sensitive cognitive





testing session and another MRI scan entailing the acquisition of more experimental sequences (T1, ASL, Spectro, Swip, QFLOW_CSF and multi-b).

In addition to the ALFA+ cohort study core sessions, participants have also been invited to undergo ¹⁸F-Flutemetamol (amyloid) and ¹⁸F-Fludeoxyglucose Positron Emission Tomography (PET) in the context of the ALFA+ V1 visit. Similarly, a subset of 213 participants have, in the context of the ALFA+ V2 visit, undergone longitudinal ¹⁸F-Flutemetamol PET. Finally, a subset of 100 participants have undergone RO-948 (tau) PET in the context of the study's V2 and are currently undergoing longitudinal RO-948 PET. Please see figure 1 below:

	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025	2026
ALFA+ V1 (N=419)											
Amyloid PET (N=361)											
FDG PET (N=360)											
ALFA+ V2 (N=370)											
Amyloid PET (N=213)											
Tau PET (N=100)											
ALFA+ V3 (ongoing)											
Tau PET (ongoing)											

With regard to the <u>fluid</u> and <u>neuroimaging</u> biomarker characterization of ALFA+ participants at V1, 342 have both full CSF (with 25 biomarkers measured [e.g. A β 42, p-tau and t-tau, A β 40, GFAP, YKL-40, sTREM2, IL6, NfL, neurogranin, S100 and α -synuclein and various other forms of p-tau in different platforms]) and PET (amyloid and FDG) data, 58 have only CSF data and, finally, 19 have only PET data. As expected due to the selection strategy, ~35% of (cognitively unimpaired) study participants are already in the <u>earliest preclinical stages of AD</u>. Specifically, based on CSF core AD biomarkers (from 396 participants with available CSF biomarkers):

- A-T- 63,1%
- A+T- 26,3%
- A+T+ 7,6%
- A-T+ 3,0%

In addition, we have also expanded the characterization of ALFA+ study participants with regard to recently developed **blood-based biomarkers**. The following blood biomarkers, have been determined in plasma samples (baseline) of the 419 individuals included in the study: A β 42, A β 40, GFAP, NfL, APOE- ϵ 4 and various forms of p-tau using a variety of techniques and platforms (a total of 15 blood-based biomarkers).

At present, the **longitudinal determinations** (samples from V2) of the following CSF biomarkers (A β 42, p-tau and t-tau, A β 40, GFAP, YKL-40, sTREM2, IL6, NfL, neurogranin, S100 and α -synuclein) have also been performed. ~40% of (cognitively unimpaired) study participants are already in the **earliest preclinical stages of AD** Specific results are as follows (from 281 participants with available CSF biomarkers):

- A-T- 56,2%
- A+T- 26,0%
- A+T+ 13,5%
- A-T+ 4,3%

Finally, we have also expanded the characterization of ALFA+ study participants with regard to recently developed **blood-based biomarkers in V2**. The following blood biomarkers have been determined in plasma samples (longitudinal): A β 42, A β 40, GFAP, NfL, APOE- ϵ 4 and various forms of p-tau using a variety of techniques and platforms (a total of 15 blood-based biomarkers).





In brief, the ALFA+ study will serve to untangle the natural history of the disease and to model the preclinical stages in order to develop successful trials.

The main characteristics of ALFA-V1 participants are: mean age 61.1 years, 60.6% females, 55,1% *APOE-ε4* carriers, mean years of education 13.6.

The main characteristics of ALFA-V1 participants are: mean age 64.2 years, 60.1% females, 55.6% APOE- $\varepsilon 4$ carriers, mean years of education 13.7.

The **timelines** of the ALFA, ALFA-MRI and ALFA+ studies as well as a **detailed list of variables** can be found in the following pages of this document.





The ALFA Study and nested Sub-studies - TIMELINES

Q2 2013 Q3 2013 Q4 2013 Q1 2014 Q2 2014 Q3 2014 Q4 2014 Q4 2014 Q4 2014 Q1 2015 - Q1 2016 Q2 2016 Q3 2016 Q4 2016 Q1 2017 Q2 2017 Q3 2017 Q4 2017 Q1 2018 Q2 2018 Q3 2018 Q4 2019 Q1 2019 Q2 2019 Q3 2019 Q4 2020 Q2 2020 Q3 2020 Q4 2020 Q1 2021 Q2 2021 Q3 2021 Q4 2021 Q1 2022 Q2 2022 Q3 2020 Q4 2

ALFA-MRI (~1,600

ALFA+ V1 (~420)

ALFA+ V2 (~420)





The ALFA Study and nested Sub-studies – DETAILED LIST OF VARIABLES

STU	DY/SUB-STUDY	N
	Alfa Parent Cohort	~2,700
	Alfa-MRI	~1,600
	Alfa+	~420

SOCIODEMOGRAPHIC		
VARIABLE	TYPE	STUDY
Gender	Transversal	
Age at inclusion	Transversal	
Place of Birth	Transversal	
Years of education	Longitudinal	
Level of education	Longitudinal	
Ethnicity	Transversal	
Civil status	Longitudinal	
Employment status	Longitudinal	
Cohabitation status	Longitudinal	
Caregiver status	Longitudinal	
Professional Activity	Longitudinal	
Socioeconomic status	Longitudinal	
Laterality/handedness	Transversal	
Bilingualism / poliglotism	Transversal	

	FAMILY HISTORY
--	----------------

VARIABLE		TYPE	STUDY	1
	Date of birth	Transversal		
	Date of birth History of dementia	Longitudinal		
Parents -	Age of parents at onset of scognitive impairment and diagnosis	Longitudinal		
	Death (date and cause) Anamnesis	Longitudinal		
	Anamnesis	Longitudinal		
Siblings	Number of siblings	Transversal		
Other	Other family members with cognitive decline	Longitudinal		
Offspring	Number of children	Longitudinal		





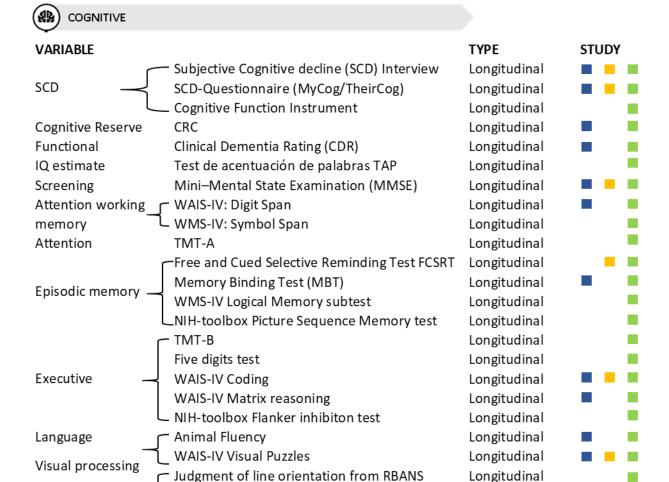


VARIABLE		TYPE	STUDY
	Psychomotor development	Longitudinal	
	Infancy diseases	Longitudinal	
	Cardiovascular disease	Longitudinal	
	Endocrine/metabolic diseases	Longitudinal	
	Digestive tract diseases	Longitudinal	
	Psychiatric diseases	Longitudinal	
	Neurological diseases	Longitudinal	
Anamnesis	Respiratory diseases	Longitudinal	
Allalillesis	Hematological diseases	Longitudinal	
	Immunological diseases	Longitudinal	
	Ophtalmological diseases	Longitudinal	
	Reproductive ans renal system diseases	Longitudinal	
	Infectious diseases	Longitudinal	
	Neoplastic diseases	Longitudinal	
	Muscle skeletal diseases	Longitudinal	
	Other diseases	Longitudinal	
	Climacteric / Reproductive history	Longitudinal	
	Medication	Longitudinal	
	GADS	Transversal	
Anxiety/depression ——	HADS	Longitudinal	
	- STAI	Longitudinal	
	PSS Perceived Stress Scale	Longitudinal	
	BRS Brief Resilience Scale	Longitudinal	
	Stressful Life Events (SNAC)	Longitudinal	
	Surgery procedures	Longitudinal	
	Practice of contact sports	Longitudinal	
	Weight	Longitudinal	
Anthropometric measures	Height	Longitudinal	
Antinopometric measures	Hip circumference	Longitudinal	
L	- Waist circumference	Longitudinal	
	Blood pressure	Longitudinal	
	Heart rate	Longitudinal	
	Hemogram	Longitudinal	
	Biochemistry	Longitudinal	
٢	- Tobacco	Longitudinal	
Substance use -	Alcohol	Longitudinal	
Ĺ	- Drugs	Longitudinal	





Transversal



LIFESTYLE HABITS

Personality

VARIABLE		TYPE	STUDY
Physical activity	MINNESOTA PHYSICAL ACTIVITY QUESTIONNAIRE	Longitudinal	
Quality of life	SF-12	Longitudinal	
	Pittsburgh Sleep Quality Index	Longitudinal	
	ESEMED	Transversal	
	SQQ Sleep Quality Questionaire	Transversal	
Sleep	ISI Insomnia Severity Index	Transversal	
Sieeh —	ESS Epworth Sleepiness Scale	Transversal	
	REM	Transversal	
	RLSYND Restless leg syndrom questionaire	Transversal	
	_ SLPDO	Transversal	
Diet	Adherence to mediterranean diet	Longitudinal	
	Exposure to pollutants	Transversal	
	Exposure to green and blue spaces	Transversal	
Cognitive activities	Leisure Activities Questionnaire (LAQ)	Longitudinal	
	Spiritual activities	Transversal	

Eysenck Personality Questionnaire (EPQ)







VARIABLETYPESTUDYAPOE genotypeTransversal■ ■ ■GWAS (Illumina Infinium Neuro Consortium (NeuroChip) Array)Transversal■ ■ ■Telomere lengthTransversal■ ■ ■



VARIABLE

Clinincal sequences (T1, T2, FLAIR)

fMRI sequences

Other sequences (ASL, Qflow, SWIP...)

Longitudinal

Longitudinal

Note: for ALFA, ~600 MRIs available (not for the whole sample)



VARIABLE	TYPE	STUDY
Incidental Findings	Longitudinal	
Fazekas scale	Longitudinal	
Global cortical atrophy scale	Longitudinal	
Left medial temporal atrophy scale	Longitudinal	
Right medial temporal atrophy scale	Longitudinal	
Koedam parietal atrophy scale	Longitudinal	
Changes in basal ganglia according to the Wahlund scale	Longitudinal	

Note: for ALFA, ~600 MRIs available (not for the whole sample)



VARIABLE	TYPE	STUDY
Amyloid (Flutemetamol) Whole Cerebellum SUVr	Longitudinal	
Amyloid (Flutemetamol) SUVr_GreyCerebellum	Longitudinal	
$Amyloid \ (Flutemetamol) \ SUVr_WholeCerebellumBrainStem$	Longitudinal	
Amyloid (Flutemetamol) Pons SUVr	Longitudinal	
Amyloid (Flutemetamol) Centiloids	Longitudinal	
Amyloid (Flutemetamol) Visual read	Longitudinal	
Amyloid (Flutemetamol) raw iamges	Longitudinal	
FDG PET result	Transversal	
FDG PET raw images	Transversal	



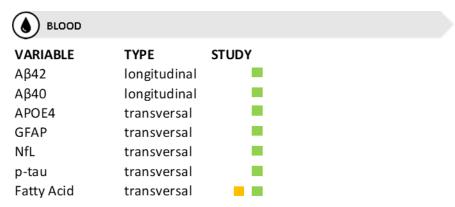




CSF

-		
VARIABLE	TYPE	STUDY
Αβ42	Longitudinal	
Αβ40	Longitudinal	
Aβ42/40 ratio	Longitudinal	
p-tau	Longitudinal	
t-tau	Longitudinal	
ATN classification	Longitudinal	
IL-6	Longitudinal	
α-Synuclein	Longitudinal	
GFAP	Longitudinal	
Neurogranin	Longitudinal	
NfL	Longitudinal	
S100	Longitudinal	
sTREM2	Longitudinal	
YKL40	Longitudinal	
N-p-tau217	Longitudinal	
Mid-p-tau231	Longitudinal	

Please note that this may not be an exhaustive, updated list of CSF biomarkers determined



Please note that this may not be an exhaustive, updated list of blood-based biomarkers determined

