

MESTRADO INTEGRADO EM MEDICINA

Clinical phenotypes in autosomal dominant Alzheimer`s disease

Jorge Manuel Gouveia Ferreira

M

2020



Clinical phenotypes in autosomal dominant Alzheimer`s disease

Dissertação de candidatura ao grau de Mestre em Medicina, submetida ao Instituto de Ciências Biomédicas Abel Salazar – Universidade do Porto

Jorge Manuel Gouveia Ferreira

Aluno do 6º ano profissionalizante de Mestrado Integrado em Medicina

Afiliação: Instituto de Ciências Biomédicas Abel Salazar – Universidade do Porto

Endereço: Rua de Jorge Viterbo Ferreira nº228, 4050-313 Porto

Endereço eletrónico: jorgegouveiaferreira@gmail.com

Orientador: Ricardo Jorge Ferreira Taipa

Médico Assistente Hospitalar de Neurologia e Neuropatologia - Unidade de Neuropatologia e Serviço de Neurologia do Departamento de Neurociências- Centro Hospitalar do Porto

Afiliação: Prof. Auxiliar Convidado da Unidade de Anatomia e Fisiologia do Sistema Nervoso do ICBAS/UP

Endereço: rtaipa.neuropat@chporto.min-saude.pt

Porto, Data: __/__/__

Jorge Manuel Gouveia Ferreira

Resumo

Introdução

A Doença de Alzheimer (AD) é a principal doença neurodegenerativa do sistema nervoso central causadora de demência, cuja etiologia é multifatorial, englobando fatores ambientais e genéticos. Diversos processos celulares e metabólicos (incluindo deposição amilóide, hiperfosforilação *tau* e ativação da micróglia) levam à morte neuronal e diminuição do fluxo sinático.

A AD divide-se na forma esporádica (sAD) e familiar (fAD), sendo a primeira a mais prevalente (99% dos casos de AD) e fAD correspondendo a 1% dos casos de AD.

Objetivos

Realizar uma revisão acerca da relação entre as diferentes mutações genéticas conhecidas e fenótipos clínicos da fAD, tendo em conta a idade de início da doença, morbimortalidade, prognóstico e variantes clínicas em cada uma das mutações genéticas supramencionadas. Pretende-se ainda rever os processos fisiopatológicos associados aos genes mutados associados à fAD.

Metodologia

A presente revisão pretende resumir as diferenças entre os genótipos e fenótipos relacionados na doença de Alzheimer's familiar autossômica dominante (ADfAD). Deste modo, para a realização desta tarefa foi realizada uma pesquisa bibliográfica na base de dados Pubmed, selecionando artigos publicados nos últimos 20 anos, em inglês, ao pesquisar por termos MeSh (Medicals Subject Headings) relacionados com ADfAD, as suas mutações, fisiopatologia e sintomas clínicos.

Resultados

A doença de Alzheimer familiar é uma forma rara de AD, que tem manifestações clínicas, para além da perda progressiva da memória. A maioria dos casos de AD apresentam-se tipicamente com défice amnésico progressivo, com início acima dos 65 anos (AD de início tardio), correspondendo na sua larga maioria às formas esporádicas de AD. Por oposição, a fAD está associada a um conjunto de manifestações cognitivas mais heterogêneas, geralmente tem uma idade de início mais precoce e um curso da doença mais agressivo. Existem 3 genes descobertos até à data cujas mutações estão relacionadas com o metabolismo de β -amilóide, e, consequentemente na fisiopatologia da fAD: presenilina-1 (PSEN-1), presenilina-2 (PSEN-2) e proteína precursora da amilóide (APP).

A fAD difere da sAD, quanto à progressão fisiopatológica, atingimento topográfico do córtex cerebral e, consequentemente, apresenta um amplo espectro de manifestações clínicas. Entre os 3

diferentes genótipos da fAD, PSEN-1 é o mais comum e existem algumas diferenças fenotípicas entre os mesmos, como por exemplo, a PSEN-2 apresentar uma idade de início mais tardio.

Conclusões

Até à data, a fisiopatologia da AD permanece pouco clara, explorar a base genética e patológica da fAD pode iluminar aspectos do mecanismo da doença subjacente e provavelmente servirá de ajuda à compreensão e tratamento da AD no futuro.

Palavras-chave: Doença de Alzheimer autossómica dominante; Doença de Alzheimer de início precoce; PSEN1; PSEN2; APP.

Abstract

Background

Alzheimer's disease (AD) is the main neurodegenerative disorder of the central nervous system that cause dementia, whose etiology is multifactorial, encompassing environmental and genetic factors. Several cellular and metabolic processes lead to neuronal death and synaptic flow decrease. Neuropathological studies showed amyloid deposits, neuritic plaques and neurofibrillary tangles at the microscopic level and, macroscopically, cerebral tissue atrophy.

AD is divided into sporadic Alzheimer's Disease (sAD) and familial Alzheimer's Disease (fAD), the first being the most prevalent (99% of AD cases) and fAD corresponding to less than 1% of AD total cases.

Objectives

To carry out a review of the relationship between the different known genetic mutations and clinical phenotypes of fAD, taking to account the age of onset of the disease, morbidity and mortality, prognosis and clinical variants in each of the aforementioned genetic mutations. Additionally, to review pathophysiological processes associated to the disease associated mutated genes.

Methods

The present review will summarize the differences between the 3 genotypes and the resulting phenotypes in autosomal dominant familial Alzheimer's Disease (ADfAD). Therefore, in order to carry out this task a bibliographic research was made using Pubmed's database, selecting articles published in the last 20 years, in English, searching for MeSh (Medical Subjects Headings) terms related with ADfAD, the mutations involved in the pathophysiology and the clinical symptoms.

Results

Familial Alzheimer's disease is a rare form of AD with clinical phenotypes beyond progressive memory impairment. The majority of AD cases has a clinical presentation limited to progressive amnesic dementia, classically aged > 65 years old, corresponding to late onset Alzheimer's disease (LOAD), almost entirely embracing cases of sAD. Contrary, fAD is associated with a wider range of clinical manifestations, often begins on earlier ages at onset (AAO) and has a more aggressive course compared to the sporadic form. There are 3 genes discovered to date whose mutations influence the β -amyloid metabolism, being responsible for the etiology of fAD: presenilin-1 (PSEN-1), presenilin-2 (PSEN-2) and amyloid precursor protein (APP).

The fAD differs from the sporadic counterpart regarding physiopathological progression and preferential topographical of brain involvement, consequently, having a wider range of clinical manifestations. Among the 3 different genotypes of fAD, PSEN-1 is the most common and there are some phenotypic differences between them, for example, PSEN-2 has a later age of onset.

Conclusions

To date, the pathophysiology of AD remains unclear, exploring the genetic and pathological basis of fAD can illuminate aspects of the underlying disease mechanism, and is likely to inform our understanding and treatment of AD in the future.

Keywords:autosomal dominant Alzheimer's disease; early-onset Alzheimer's disease; PSEN1; PSEN2; APP

Abbreviations list

AChI – Acetylcholinesterase inhibitors

AAO – age at onset

A β – β -amyloid

ABCA1 – ATP-binding cassette transporter 1

ABCA7 – ATP-binding cassette transporter 7

AD – Alzheimer's disease

ADEOFAD – Autosomal dominant early-onset familial Alzheimer's disease

ADfAD – Autosomal dominant familial Alzheimer's disease

APP – amyloid precursor protein

BIN1 – Bridging Integrator 1

BPSD – Behavioral and psychiatric symptoms of dementia

CAA – cerebral amyloid angiopathy

CNS – Central Nervous System

EOAD – Early-onset Alzheimer's disease

F-FDG-PET – Fluorodesoxyglucose Pet scan

fAD – Familial Alzheimer's disease

LOAD – Late-onset Alzheimer's disease

PIB PET – Pittsburgh compound B PET scan

PCR – Polymerase Chain Reaction

PSEN1 – Presenilin 1

PSEN2 – Presenilin 2

sAD – Sporadic Alzheimer's disease

TREM2 – triggering receptor expressed on myeloid cells 2

Index

Resumo.....	i
Abstract.....	iii
Abbreviations list.....	v
Table List.....	vii
Background.....	1
Familial Alzheimer's disease.....	3
Presenilin 1.....	6
Presenilin 2.....	8
Amyloid precursor protein.....	9
Early-onset Alzheimer's Disease.....	11
Conclusions.....	12
References.....	13

Table List

Table 1 – Examples of PSEN1 mutations and its clinic variants.

Table 2 – Example of PSEN2 mutations and its clinic variants.

Table 3 – Examples of APP mutations and its clinic variants.

Background

Alzheimer's disease, the commonest cause of dementia, is a progressive neurodegenerative disease recognised by the World Health Organisation as a global public health priority, with huge implications for individuals and society. As the population across the globe ages, the incidence and prevalence of age-related diseases is rising¹, such as dementia, defined as an acquired progressive cognitive impairment sufficient to impact on activities of daily living, resulting as a syndrome including memory and intellectual loss, behavioral modifications and incapacity to conclude complex tasks². Despite large gains in the understanding of AD pathogenesis there are no disease-modifying treatments, only symptomatic ones.

AD is divided into sporadic Alzheimer's Disease (sAD) and familial Alzheimer's Disease (fAD), the first being the most prevalent (99% of AD cases) and fAD corresponding to <1% of AD total cases.

Similar to sAD, there is no disease-modifying treatments for familial Alzheimer's Disease. Anti-AD pharmacotherapies (AChE i.e. donepezil, galantamine, rivastigmine, memantine) provide modest but meaningful benefits by delaying disability and reducing clinical progression³. Similar to sAD, some fAD cases have insidious onset of episodic memory symptoms but it often differs clinically due to its distinctive features, namely early age at onset (AAO), often in between 35-55 years old, faster development of the disease and more aggressive course with increased morbidity and mortality⁴. Therefore, a multidisciplinary approach and symptomatic targeted treatment should be implemented.

Regarding genetics, and in opposition to sAD that has no evidence to show an increased risk for the offspring to develop the disease, fAD is commonly transmitted to the offspring (with 50% chance) because the mutations in the genes responsible for fAD are transmitted in an autosomal dominant way. It also differs from the first in genotypes, in pathophysiological progression, cerebral cortex topography and, consequently, clinical characteristics, demonstrating a wider spectrum of manifestations. There are three genes discovered to date whose mutations are involved in the pathophysiology of fAD: presenilin-1 (PSEN-1); presenilin-2 (PSEN-2) and amyloid precursor protein (APP)⁵. All these mutations are related with β -amyloid metabolism. The amyloid cascade is the most accepted theory in the scientific community at the origin of the pathophysiology of AD, resulting from the imbalance in the production and clearance of β -amyloid⁶.

The present review will summarize the differences between the 3 genotypes and the resulting phenotypes in autosomal dominant familial Alzheimer's Disease (ADfAD). Therefore, in order to carry out this task we selected experimental and review studies from Pubmed database, searching for MeSh

(Medicals Subjects Headings) terms related with ADfAD, the mutations involved in the pathophysiology and the clinical symptoms.

Familial Alzheimer's Disease

AD is divided into sporadic Alzheimer's Disease (sAD) and familial Alzheimer's Disease, the first being the most prevalent (99% of AD cases) and fAD corresponding to <1% of AD total cases¹. The majority of AD cases has a clinical presentation limited to progressive amnesic dementia, classically aged > 65 years old, corresponding to late onset Alzheimer's disease (LOAD), almost entirely embracing cases of sAD. In rarer situations, the disease can manifest at earlier onset ages (<65 years), corresponding to early onset Alzheimer's disease (EOAD), which comprises about 5% to 6% of cases of AD, where about 20% of cases correspond to familial Alzheimer's disease (fAD), representing a much higher genetic predisposition than LOAD⁷.

FAD is transmitted to the offspring in an autosomal dominant way⁸, resulting from pathogenic mutations in one of the three identified genes responsible for the disease: PSEN-1; PSEN-2 and APP. The APOε4 allele is related with the sporadic form of AD, but not causative⁹. Currently it is believed that APOε4 is a risk factor for both sAD and late-onset fAD¹⁰. In both sporadic and familial AD, the possession of APOε4 allele is associated with decreased AAO.¹¹ Many other molecular factors (i. e. immunological cause (TREM2) with lipid disorder (ABCA1, ABCA7), metabolites transport (BIN1)) can be related to the pathophysiology of AD, including β-amyloid deposition, *tau* hyperphosphorylation, microglia activation and neuroinflammation¹², being the amyloid cascade theory the most accepted in the scientific community to explain the pathophysiology of AD¹³. Still, it remains uncertain if the neuropathology and molecular processes are the same for sAD and fAD cases. It is also believed that epigenetic has a role in fAD because the heterogeneity of clinical manifestations. Despite the neurodegenerative processes of both forms of AD appears to be similar, it is thought that the physiopathology of fAD differs from sAD by the excess production of β-amyloid (Aβ42, Aβ40), whereas in sAD it is thought to be related to the lack of β-amyloid clearance. Aβ42/Aβ40 ratio may also vary amongst these different fAD gene mutations with different clinical repercussions⁵, but, overall, it is commonly increased. Pathological studies have demonstrated that the proportion of brain tissue occupied by Aβ42-containing plaques is significantly higher in fAD than sAD.

Macroscopically, in both forms, cerebral atrophy is observed, with neuronal loss and diminished synaptic flow. Similarly, studies have shown extracellular β-amyloid deposition in the brain parenchyma, neurofibrillary tangles and neuritic plaques in both AD forms. It is believed that β-amyloid deposition starts around 20-25 years before onset of symptoms in cases of fAD¹. In sAD the time between β-amyloid accumulation and clinical symptoms remains to be quantified, but current theories suggest that this time may be for more than a decade¹⁴ Primitive β-amyloid deposits were

more frequently distributed in regular clusters and less frequently distributed in large clusters in fAD compared with sAD.¹⁵

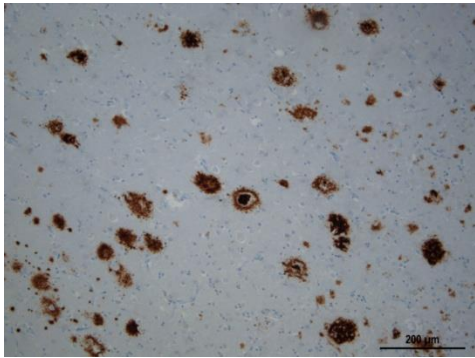


Figure 1 – Parenchymal beta-amyloid deposits (globose diffuse and mature plaques) in neocortex (Portuguese Brain Bank archive). Scale bar: 200µm

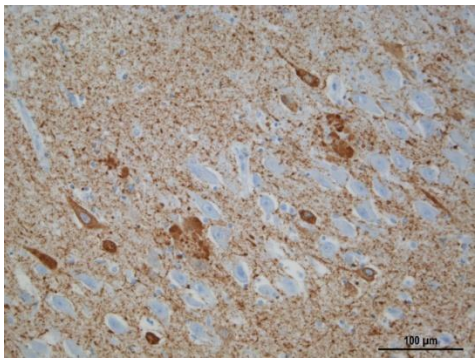


Figure 2 – Tau pathology (neurofibrillary tangles, neuritic plaques and dystrophic neurites) in the hippocampus (Portuguese Brain Bank archive). Scale bar: 200µm

Classically, in sAD patients the medial temporal lobe structures are most affected, with progressive memory loss as main clinical manifestation. In fAD cases many clinical manifestations, called variants (behavioral EOAD, disexecutive EOAD, aphasic variant, posterior cortical atrophy) occur according to the cerebral topography preferential involvement of the pathology. The hippocampus may be spared in an early stage of some of these variants⁵.

Compared to sAD the course of fAD variants seems to be more aggressive, and presents an age at onset (AAO) approximately 20 years earlier (AAO=35-55years). The commonest variants of fAD in descending order are logopenic variant primary progressive aphasia, posterior cortical atrophy, behavioral/disexecutive variant and the acalculic variant⁶. Most of fAD cases result from PSEN1 mutations (>75%), APP mutations are responsible for nearly 15-20% of autosomal dominant early

onset familial Alzheimer's disease (AEOFAD) and the rarest PSEN2 mutations related correspond to <5%.¹⁶

The management should be multidisciplinary and directed to the symptoms accordingly to each variant and genetics testing and counseling should be done in all fAD cases⁴. Recently, for diagnosis of fAD, a genetic testing has been advocate in the presence of an EOAD patient with family history of one relative with early onset dementia¹⁷.

Presenilin-1

PSEN1 gene is located in chromosome 14q24.3¹⁸, and it is a vital component of the γ -secretase complex, which cleaves APP into A β fragments.¹⁹ Mutations in PSEN1 account for up to 75% of fAD, with complete penetrance and early age of onset.²⁰ Mutant γ -secretase increases A β 42 level while it decreases A β 40 level, leading to an increased A β 42/A β 40 ratio. Morphologic variants in A β plaques due to PSEN1 mutations may result in cotton wool plaques¹⁹. Cotton wool plaque is formed by large rounded A β deposits, and it tends to be immune-positive for A β 42, rather than for A β 40²¹. PSEN1 mutations are often missense mutations, but deletions and insertions have also been detected. To date, around 300 pathogenic mutations have been identified in PSEN1, of which 70% mutations occur in exons 5, 6, 7 and 8⁵. The mutations are present in gnomAD database (<https://gnomad.broadinstitute.org>) or jMorp (<https://jmorp.megabank.tohoku.ac.jp>).

Regarding the cerebral topography, EOFAD with PSEN1 mutations involve multiple regions, mainly the neocortex, beyond the hippocampus and with subcortical affection, marked depigmentation of the substantia nigra and locus coeruleus²². Atrophy of the pyramidal tract portion at medulla oblongata and A β deposits in the cerebellum have also been identified¹⁹. Cerebral amyloid angiopathy was also noted in some cases. The cognitive profile has so far predominantly shown amnesic and cognitive symptoms typical of dementia, but may present with some unusual clinical features, in particular spastic paraparesis²³ and epileptic seizures⁵. EOAD with PSEN1 mutations can present extrapyramidal signs, behavioral features suggestive of frontotemporal dementia²⁴, language impairment and cerebellar ataxia, but also behavioral and psychiatric symptoms of dementia (BPSD) like anxiety, hallucinations and delusions²⁵. However, almost all these symptoms having been also reported in sAD cases.

EOFAD with PSEN1 mutations typically have an AAO in the early 40s, ranging from 35 to 65 years. The course of the disease and the clinical manifestations may change accordingly to specific PSEN1 mutations (see table 1), but there is no evidence to support a specific mutation-related phenotype. The disease duration varies from 5 to 15 years⁵.

Mutation	Age at onset	Age at death	Disease duration	Symptoms
Val97Leu	36.5	52.4	15.9	Memory disturbance, dyscalculia, apathy and personality changes
Ile167del	38	55	15.5	Memory decline, disorientation, spastic paraparesis and behavior variants
Leu226Arg	60	-	-	Language disability, memory decline and personality change, irritability
Met233Leu	40.6	44.8	4.2	Progressive memory decline, seizures and paralysis
Phe388Leu	43	44.4	1.41	Progressive memory decline, psychosis and paranoid delusions
Ala434Thr	35	55	8	Progressive memory decline, hallucinations and delusions
Table 1 – Examples of PSEN1 mutations and its clinic variants. Adapted from Quin <i>et al</i> ⁵ .				

Presenilin-2

PSEN2 gene is located on chromosome 1q31-q42²⁶, and its mutations are very rare, to date 33 pathogenic PSEN2 mutations have been detected worldwide⁵. PSEN2 is a main component of the γ -secretase complex along with PSEN1, nicastrin, Aph-1, and PEN-2²⁷. Pathophysiologically, PSEN2 mutations change the γ -secretase activity and leads to elevation of A β 42/A β 40 ratio, similarly to the PSEN1 mutation. Although, in PSEN2 mutations less amyloid peptide is produced then in PSEN1 mutations, besides the close homology. Neuritic plaque formation and neurofibrillary tangle accumulation may be seen as neuropathological changes in some people with PSEN2 mutations¹⁷.

Regarding EOFAD with PSEN2 mutations cerebral topography of involvement, imaging studies in these patients using PIB PET and F-FDG-PET showed amyloid deposits and hypometabolism²⁸, respectively, in frontal, temporal, parietal and insular lobes, cingulate cortex, precuneus, caudate nucleus, striatum and thalamus, alongside with general cortical atrophy (hippocampus included).

PSEN2 mutations are the least common of the three genes known to cause fAD, and therefore literature is relatively scant. The prominent symptoms of these patients is memory impairment, commonly associated with language impairment, depression, personality changes, parkinsonism and myoclonus, paranoia, visual hallucinations and agitation²⁹ (see table 2).

Compared to PSEN1 mutation carriers, PSEN2 mutation carriers have a later AAO, commonly in the early 50s, ranging from 35 to 65 years and the course of the disease seems to be more indolent³⁰, the duration of the disease appears to range from 10 to 20 years.

Mutation	Age at onset	Disease duration	Symptoms
Lys82Arg	50	-	Speech difficulties and depressions
Prol123Leu	57	-	Parkinsonism, myoclonus and personality changes
ASN141tYR	46	16	Speech difficulties, paranoia, visual hallucinations and agitation

Table 2 – Example of PSEN2 mutations and its clinic variants. Adapted from Quin *et al.*⁵

Amyloid precursor protein

The gene encoding APP is located on chromosome 21q21.3²³. It is the second most common mutation of the three, and to the date 63 pathogenic mutations in APP gene were detected worldwide. APP is a transmembrane protein, which plays a role in synaptic plasticity³¹. Three enzymes have cleavage sites in APP: α -, β - and γ -secretases. Proteolysis of APP by α - and γ -secretases results in nonpathogenic fragments (sAPP α and α -C-terminal fragment) in nonamyloidogenic pathway. In amyloidogenic pathway, abnormal cleavage of APP by β - and γ -secretases could result in two major A β precursors that include sAPP β and β -CTF. The action of γ -secretases on β -CTF generates A β extracellularly that deposits in the neuron in AD patients. Physiopathologically, different mechanisms occur from mutation to mutation, since the destruction of A β 40 cleavage site to increased γ -secretase action in A β 42 location, all leading to an increased A β 42/A β 40 ratio. A β 42, A β 40 and A β 42/A β 40 ratio have been established as the most important biomarkers for AD⁵.

In terms of cortical topography of hypometabolism and cerebral atrophy, APP-related fAD does not differ much from the others fAD mutations.³² Hippocampal sparing and greater posterior neocortical involvement is reported. Involvement of white matter tracts in posterior association areas and frontoparietal networks are characteristics that differ mainly from sAD areas (mainly neocortex).³³ Cerebral amyloid angiopathy (CAA) with microhemorrhages is reported often in studies with APP-related fAD brains, consequently present focal neurological signs, such as aphasia.³⁴

The commonest clinical manifestation is progressive memory impairment, followed by rapid progression to severe dementia within 5-10 years. The AAO is around 46-55 years and it varies according to specific mutations (see table 3).³⁵

Mutation	Age at onset	Age at death	Diese duration	Symptoms
Asp678His	50.6	58.5	7.9	Progressive memory decline, restlessness, persecutory delusions, self-talking and slurred speech
Val715Met	45	48	3	Memory decline, bradykinesia and irritability
Ile716Thr	35-40	-	-	Progressive memory decline, aphasia and increased muscular tension
Val717Ile	54.7	67.1	11.7	Executive dysfunction, disorientation, spastic paraparesis and cerebellar ataxia
Met722Lys	42.8	58.9	16.1	Memory decline, executive dysfunction, disorientation, aphasia and depression
Lys724Met	46.6	53.4	6.8	Verbalization, executive dysfunction, sluggishness and apathy
Table 3 – Examples of APP mutations and its clinic variants. Adapted from Quin <i>et al.</i> ⁵				

Early-Onset Alzheimer's Disease

Early Onset Alzheimer's Disease (EOAD) is defined as having an AAO younger than 65 years. The overall clinical profile of EOAD differs from that of LOAD. A higher proportion of patients with EOAD present symptoms such as language, visuospatial or dysexecutive phenotypes rather than the usual amnesic disorder seen in LOAD and have a genetic predisposition, being 20% of EOAD diagnosed with fAD. In amnesic EOAD a higher frequency of APOε4 allele was verified, while the APOε4 allele frequency in variants is less observed⁶.

There is no scientific evidence showing an increased probability of the offspring get the disease in sAD cases, contrarily to fAD. The risk is higher in both forms when in the presence of APOε2, APOε3 or APOε4 gene mutation.

Patients with AAO earlier than 35 are classified as very early-onset Alzheimer's Disease (VEOAD), which are associated with more aggressive course of the disease and increased mortality, being the majority of cases, fAD cases³⁶.

Conclusions

AD is the commonest cause of dementia worldwide and it is divided in two forms, sporadic (sAD) and familial (fAD). Although fAD represents a much lesser number of cases (approximately 1%), severe features like earlier AAO, more aggressive course of the disease and earlier age of death shows the importance of correct management and treatment. To do so, a better understanding of the pathophysiology with more studies to characterize the variant phenotypes and correlate genotypes, are needed. The comparison between the two AD forms is relevant, in order to recognize similarities and differences between both forms and gain insights into the cascade of pathological events and how they relate to clinical manifestations. To the date studies have shown main differences in pathophysiology such as higher ratio $A\beta_{42}/A\beta_{40}$ in fAD, different mechanisms in the amyloidogenic pathway that leads to an increased production of $A\beta_{42}$ in fAD and a decreased clearance of $A\beta_{40}$ in sAD. Additionally, fAD show more commonly hippocampal sparing and less mesial temporal lobe disease, and a greater posterior neocortical atrophy (parietal and temporoparietal junction) and hypometabolism. Consequently, fAD has an overall less semantic memory impairment and greater attention, executive, praxis, visuospatial and psychosocial difficulties (more associated with depression and anxiety) compared to sAD.

The main similarities reported to date clinically and pathophysiologically are, respectively, progressive memory and cognitive impairment, and amyloid deposition, *tau* hyperphosphorylation, neurofibrillary tangles, a common amyloidogenic pathway, same relationship between $A\beta$ deposition and APO ϵ genotype (APO ϵ 4 allele possession is associated with decreased age at onset in both sporadic and familial AD), decreased synaptic flow, neuronal hypometabolism and cortical atrophy.

To date, the pathophysiology of AD remains unclear, exploring the genetic and pathological basis of this phenotypic heterogeneity can illuminate aspects of the underlying disease mechanism, and is likely to inform our understanding and treatment of AD in the future.

-
- ¹Shao, W., Peng, D., Wang, X. *et al.* Genetics of Alzheimer's disease: From pathogenesis to clinical usage. *Journal of Clinical Neuroscience*, 2017; 45, 1-8.
- ²Høgh P, *et al.* Alzheimer's Disease. 2017; 179(12), 4-5.
- ³Alireza *et al.* The Alzheimer's Disease Clinical Spectrum: Diagnosis and Management. *Medical Clinics of North America*. 2019; 103(02), 263-293.
- ⁴Wu, L., Rosa-Neto *et al.* Early-Onset Familial Alzheimer's Disease (EOFAD). *The Canadian Journal of Neurological Sciences*. 2012; 39(04), 436-445.
- ⁵Qi Qin, Yunsu Yin, Yan Wang, Yuanyuan Lu, Yi Tang, Jianping Jia *et al.* Gene mutations associated with early onset familial Alzheimer's disease in China: an overview and current status. 2020; 8 (10), 11-18.
- ⁶Hardy, J. A., Higgins, G. A. *Et al.* Alzheimer's disease: the amyloid cascade hypothesis. *Science*. 1992; 256, 184-5.
- ⁷M. F. Mendez *et al.* Early-onset Alzheimer Disease and its Variants. *Continuum*. 2019; 25(1), 34-51.
- ⁸G. Raux, L. Guyant-Maréchal, C. Martin *et al.* Molecular diagnosis of autosomal dominant early onset Alzheimer's disease: an update. *Journal of Medical Genetics*. 2005; 42(10), 793-795.
- ⁹Manzano, J. González, A. Marcos, J. Matías-Guiu *et al.* Genetics and Alzheimer's disease. 2009; 24(2), 83-89.
- ¹⁰Jolanta Dorszewska, M. Predecki, Anna Oczkowska, Mateusz Dezor, Wojciech Kozubski *et al.* Molecular Basis of Familial and Sporadic Alzheimer's Disease. 2016; 13(9), 952-963.
- ¹¹Natalie S. Ryan, Martin N. Rossor *et al.* Correlating familial Alzheimer's disease gene mutations with clinical phenotype. *Biomark Med*. 2010; 4(1), 99-112.
- ¹²Heneka M. T., Carson M. J., El Khoury J. *et al.* Neuroinflammation in Alzheimer's disease. *Lancet Neurology*. 2015; 14(4), 388-405.
- ¹³Gómez-Isla, Growdon, McNamara *et al.* The impact of different presenilin 1 and presenilin 2 mutations on amyloid deposition, neurofibrillary changes and neuronal loss in the familial Alzheimer's disease brain. *Brain*. 1999; 122(9), 1709-1719.
- ¹⁴Sperling *et al.* Towards defining the preclinical stages of Alzheimer's disease: Recommendations from the National Institute on Aging-Alzheimer's Association workgroups on diagnostic guidelines for Alzheimer's disease *Alzheimer's & Dementia*. 2011; 7, 280-292.
- ¹⁵R. Armstrong *et al.* Spatial patterns of beta-amyloid deposits in familial and sporadic AD. 2011; 49(3), 153-61.
- ¹⁶Randall J. Bateman, Paul S. Aisen, Bart de Strooper *et al.* Autosomal-dominant Alzheimer's disease: a review and proposal for the prevention of Alzheimer's disease. *Alzheimer's research and therapy*. 2011; 3(1), 97-121.
- ¹⁷Jiao, Tang, Liu *et al.* Mutational analysis in early-onset familial Alzheimer's disease in Mainland China. *Neurobiology of aging*. 2014; 35 (8), 1951-1956.
- ¹⁸Zhangyu Zou, Changyun Liu, Chunhui Che, Huapin Huang *et al.* Clinical genetics of Alzheimer's disease. *BioMed Research International*. 2014; doi:10.1155/2014/291862.
- ¹⁹Steiner, Fluhrer, Haass *et al.* Intramembrane proteolysis by gamma-secretase. *Journal Biol Chem*. 2008; 283 (44), 29627-31.
- ²⁰Giri M., Zhang M., Lu Y. *et al.* Genes associated with Alzheimer's disease: an overview and current status. *Clinical Interv Aging*. 2016; 11, 665-681.
- ²¹Tolia A., Horré K., De Strooper *et al.* Transmembrane domain 9 of presenilin determines the dynamic conformation of the catalytic site of gamma-secretase. *Journal Biol Chem*. 2008; 283 (28), 19793-803.
- ²²Tomoko Miki, Osamu Yokota, Takashi Haraguchi *et al.* Young adult-onset, very slowly progressive cognitive decline with spastic paraparesis in Alzheimer's disease with cotton wool plaques due to a novel presenilin1 G417S mutation. *Acta Neuropathologica Communications*. 2019; 10(02), 7:19.
- ²³Karlstrom H., Brooks W. S., Kwok J. B. *et al.* Variable phenotype of Alzheimer's disease with spastic paraparesis. *Journal Neurochem*. 2008; 104, 573-583.
- ²⁴M. F. Mendez, McMurtry *et al.* Frontotemporal dementia-like phenotypes associated with presenilin-1 mutations. *Am J Alzheimer's Dis Other Dement*. 2006; 21:281-286.

-
- ²⁵ Larner *et al.* Presenilin-1 Mutations in Alzheimer's Disease: Na Update on Genotype-Phenotype Relationships. *Journal of Alzheimer's Disease*. 2013; 37(4), 653-659.
- ²⁶ Perry G. Ridge, Mark Ebbert, John Kauwe *et al.* Genetics of Alzheimer's disease. *BioMed Research International*. 2013; doi:10.1155/2013/254954.
- ²⁷ Wakabayashi T., De Strooper B. *et al.* Presenilins: members of the gamma-secretase quartets, but part-time soloists too. *Physiology (Bethesda)*. 2008; 194-204.
- ²⁸ Yang Z., Tian J., Shang *et al.* Gene mutations analysis of 3 families with early-onset Alzheimer's disease. *Chinese Journal of Neurology*. 2016; 49(9), 682-686.
- ²⁹ Cao, Qiu, Zheng *et al.* Study of mutations of presenilin 1 gene in early onset familial Alzheimer's disease. *Chinese Journal of Medical Genetics*. 2014; 31(3), 298-301.
- ³⁰ Bagyinszky, Youn, Na *et al.* Mutations, associated with early-onset Alzheimer's disease, discovered in Asian countries. *Clinical Interventions in Aging*. 2016; 11, 1467-1488.
- ³¹ Turner, P. R., O'Connor, Tate, Abraham *et al.* Roles of amyloid precursor protein and its fragments in regulating neural activity, plasticity and memory. *Progress in neurobiology*. 2003; 70 (1), 1-32.
- ³² Goate A., Chartier-Harlin M. C., Mullan M. *et al.* Segregation of a missense mutation in the precursor protein gene with familial Alzheimer's disease. *Nature*. 1991; 349 (6311), 704-706.
- ³³ Jiang, Zhou, Li *et al.* Mutation screening in Chinese patients with familial Alzheimer's disease by whole-exome sequencing. *Neurobiology of Aging*. 2019; 76 (7), 1010-1016.
- ³⁴ Huang, Hsiao, Lin *et al.* Amyloid PET pattern with dementia and amyloid angiopathy in Taiwan familial AD with D678H APP mutation. *Journal of Neurological Sciences*. 2019; 398, 107-116.
- ³⁵ Natalie S. Ryan and Martin N. Rossor *et al.* Correlating familial Alzheimer's disease gene mutations with clinical phenotype. *Biomark Med*. 2010; 4(1), 99-112.
- ³⁶ Rossor M. N., Newman S. Frackowiak R. S., Lantos P., Kennedy A. M. *et al.* Alzheimer's disease families, with amyloid precursor protein mutations. *ANN NY Acad Sci*. 1993; 695, 198-202.