

Subcartical encefalopathy plus suicidal ideation: clinical presentation of uncommon genotype of CADASIL



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BACKGROUND AND OBJECTIVES

Background: CADASIL (cerebral autosomal dominant arteriopathy with subcortical infarcts and leucoencephalopathy) is a systemic artery disease responsible for more or less diffuse white-matter lesions associated with small deep infarcts. The disease affects middle-aged adults and it is caused by mutations of the Notch3 gene located on chromosome 19. CADASIL is clinically characterized by the recurrence of ischaemic stroke leading to a pseudobulbar palsy and subcortical dementia, attacks of migraine with aura and mood disorders.

Objectives: To describe a case of CADASIL with uncommon genotype

CASE REPORT: A 48-year-old-right-handed man, with familiar history of seizures, absence of brain damage in childhood and history of hypertension and behavioral disorders and suicidal ideation, was admitted to our department with a sudden left severe hemiparesis.

ON ADMISSION TO OUR DEPARTMENT:

- On neurological examination, sudden left severe hemiparesis, dysarthria (NIHSS=11). Depression mood was highlighted.
- First level laboratory tests were normal.
- First Brain CT was normal
- Brain CT after 48 hours showed: ipodensity in the right semioval center right and leukoencephalopathy
- ■Both heart and neck vassel ultrasonography examinations were normal
- •4-hour Holter ambulatory ECG monitoring was normal
- Serum research for tumoral markers was negative

MRI showed an extensive and diffuse DWI <u>hyperintense signal in in the right semioval</u> <u>center</u> and extensive small deep infarcts and leukoencephalopathy. (C-D-E)

CSF analysis including total proteins, count cell, glucose and isoelettrofocusing was normal

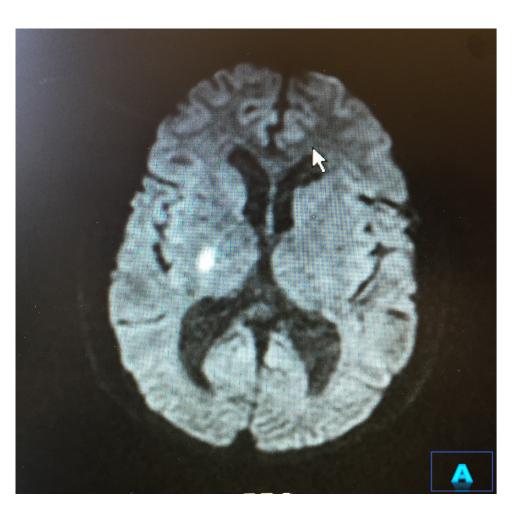
Genetic tests for <u>Fabry disease</u> and <u>CADASIL</u> showed: <u>S497L in exon 9 for NOTCH3</u>

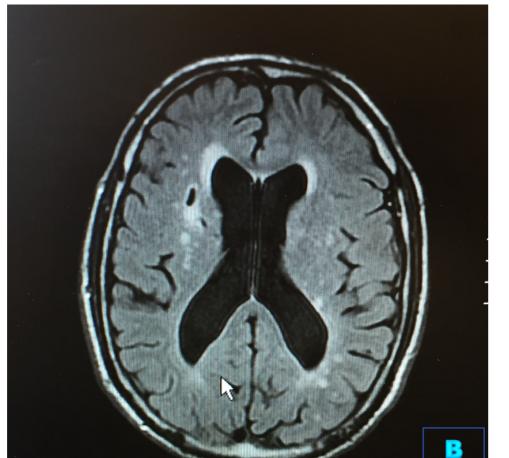
Over the next two weeks: Improve general condictions.

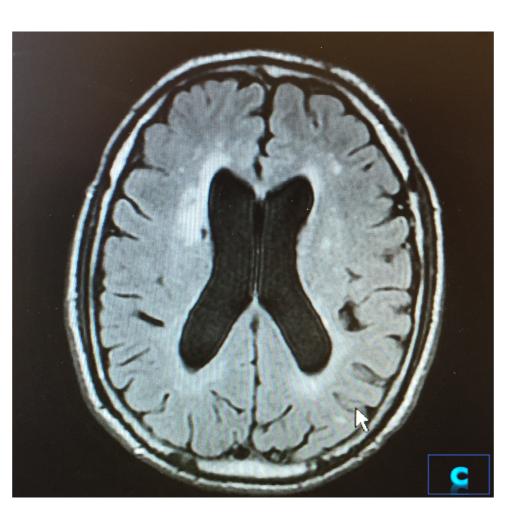
Motor and logopedic <u>rehabilitation therapy</u> associated with <u>farmacological therapy</u> for depressive symptoms and behavioral disorders was started as soon as possible with improvement of motor impairmant

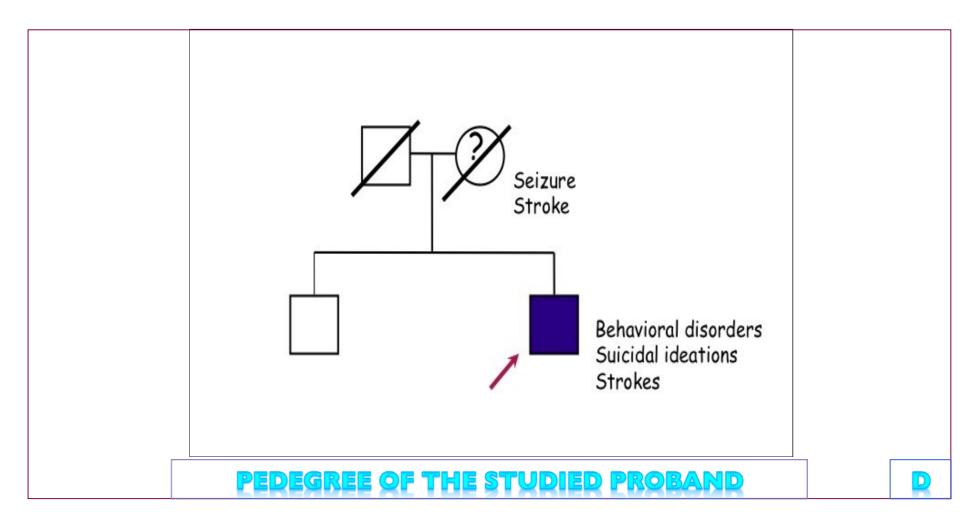
According to publish literature data

FINAL DIAGNOSIS WAS: CADASIL







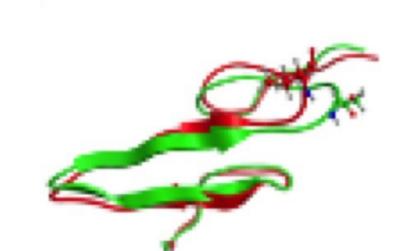


REVIEW OF LITERATURE

• NOTCH3 has 33 exons but all CADASIL mutations occur in exons 2–24, which encode the 34 EGF like region.

All mutations lead to an odd number of cysteine

- ✓ 95% missense mutations
- ✓ 5% small in-frame deletions or splice-site mutations
- ✓ polymorfisms
- •In one series of 229 CADASIL Italian patients, an heterogeneous mutational spectrum has been observed.
- A new type of mutation has been showed recently: <u>no-cys mutations</u>
- A silico-model study has revelead a no-cys mutation for the variation S497L



Structural superposition of the wild type (green) and Ser497Leu (red) Notch3 protein.

An "in silico study" has revealed a rather complex molecular mechanism of Notch3 on the structural level; based on the nature and position of S497L mutation, a consensus significant loss of beta-sheet structure is observed.

A series of Notch3 mutations in CADASIL; insights from 3D molecular modelling and evolutionary analyses

Journal of Molecular Biochemistry (2014) 3, 97-105

DISCUSSION AND CONCLUSION

- We described a case of CADASIL with psychiatric onset and stroke due to no-cys mutation S497L in exon 9 for NOTCH3.
- Strokes and behavioral disorders are the most common symthoms in CADASIL in a large series from Italy, but cognitive impairment, migraine with aura and seizures can occur.
- All classic mutations for NOTCH3 lead to an odd number of cysteine EGFR.
- No-cys mutations could be a key to undestand polymorfisms and new mutations for NOTCH3.
- Full sequencing of exons 2-24 is mandatory for CADASIL screening.

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- •Vlachakis, Spyridon Champeris Tsaniras, Katerina Ioannidou, Louis Papageorgiou, Marc Baumann and Sophia Kossida A series of Notch3 mutations in CADASIL; insights from 3D molecular modelling and evolutionary analyses. Journal of Molecular Biochemistry (2014) 3, 97-105.