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Expanding the clinical and EEG spectrum of CNKSR2-related encephalopathy with status epilepticus during slow sleep (ESES)



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HIGHLIGHTS

• Worsening of epilepsy associated with increment of awake spike-wave-index.

- A frontal topography of sleep EEG epileptic activity in the active phase of ESES.
- · Language disorder due to speech/oro-motor dyspraxia.

ABSTRACT

Objective: To investigate the clinical and EEG features of Encephalopathy with Status Epilepticus during slow Sleep (ESES) related to CNKSR2 pathogenic variants.

Methods: Detailed clinical history, repeated wakefulness/overnight sleep EEGs, brain MRI were collected in five patients, including one female, with CNKSR2-related ESES.

Results: Neurodevelopment in infancy was normal in two patients, delayed in three. Epilepsy onset (age range: 2-6 years) was associated with appearance or aggravation of cognitive impairment, language regression and/or behavioral disorders. Worsening of epilepsy and of cognitive/behavioral disturbances paralleled by enhancement of non-rapid eye movement (NREM) sleep-related, frontally predominant, EEG epileptic discharges [spike-wave-index (SWI): range 60–96%] was consistent with ESES. In three patients, episodes of absence status epilepticus or aggravation of atypical absences occurred, in this latter

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Abbreviations: AED, anti-epileptic drugs; array-CGH, array-comparative genomic hybridization; EAS, epilepsia aphasia spectrum disorders; EEG, electroencephalogram; ESES, encephalopathy with status epilepticus during slow sleep; MRI, magnetic resonance imaging; NGS, next generation sequencing; NREM, non-rapid eye movement; REM, rapid eye movement; SWI, spike wave index; WES, whole exome sequencing; WISC-R, Wechsler intelligence scale for children revised; WPPSI, Wechsler preschool and primary scale of intelligence.

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case associated with striking increment of awake SWI. Speech/oro-motor dyspraxia was diagnosed in four patients. In two patients, long-term follow-up showed epilepsy remission and persistence of mild/-moderate cognitive disorders and behavioral disturbances into adulthood.

Conclusions: Novel findings of our study are occurrence also in females, normal neurodevelopment before epilepsy onset, epilepsy aggravation associated with enhanced awake SWI, mild/moderate evolution in adulthood and language disorder due to speech/oro-motor dyspraxia.

Significance: Our findings expand the phenotypic spectrum of CNKSR2-related ESES.

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1. Introduction

Encephalopathy with status epilepticus during slow sleep (ESES) (Tassinari et al., 1977) is a self-limiting, age-related, epilepsy syndrome characterized by different seizure types, neuropsychological regression, behavioral disorders and a typical electroencephalogram (EEG) pattern of extreme activation of epileptic discharges during non-rapid eye movement (NREM) sleep (Patry et al., 1971). Recently, pathogenic variants in several genes, including GRIN2A and CNKSR2, have been reported to be associated with epilepsia-aphasia-spectrum disorders (EAS), a spectrum of conditions that encompasses ESES (Kessi et al., 2018; Lesca et al., 2019). So far pathogenic variants in the CNKSR2 gene have been reported only in few patients with ESES (Lesca et al., 2012; Vaags et al., 2014; Damiano et al., 2017; Sun et al, 2018). Few additional adult patients in whom ESES was speculated to have occurred in childhood, have also been described (Vaags et al., 2014; Aypar et al., 2015; Hu et al., 2016; Damiano et al., 2017).

CNKSR2 gene (OMIM *300724), located on chromosome Xp22.12 encodes a multidomain synaptic scaffold and adaptor protein, connector enhancer of KSR2. The protein is exclusively expressed in the brain, mostly in the hippocampus, amygdala, caudate nucleus and cerebellum (Nagase et al. 1998). It connects with densin-180, PSD95, and S-SCAM in the neuronal postsynaptic density (PSD) (Yao et al. 2000, Ohtakara et al. 2002) and participates in the downstream pathway of RAS-MAPK signal transduction (Therrien et al., 1998, Lanigan et al. 2003, Wellbrock et al., 2004) regulating neuronal complexity and synaptogenesis (Lanigan et al. 2003, Hu et al. 2016) as well as neuronal proliferation, migration, differentiation and death (Bumeister et al. 2004, Liu et al. 2009). Study on CNKSR2-depleted animals, in addition, demonstrated a reduction of the complexity and length of hippocampal neurons (Hu et al. 2016) confirming the possible role in synaptogenesis.

Here we report the clinical and EEG findings of five novel patients with *CNKSR2*-related ESES, aiming to expand the phenotypic spectrum of this rare disorder.

2. Material and methods

Five patients with ESES associated with *CNKSR2* pathogenic variants were collected through data sharing with European Epilepsy and Genetic Centers. All patients underwent brain magnetic resonance imaging (MRI) scanning and repeated wakefulness/ overnight sleep EEG recordings. The duration of the wakefulness EEGs could vary from 20 to 60 minutes, whereas the duration of the wakefulness/overnight EEG could vary from 12 to 18 hours in the different centers. During the course of the disease, sleep EEGs during daytime including a full sleep cycle were also performed, usually to evaluate response to treatment.

Array-Comparative Genomic Hybridization (array-CGH), targeted gene panel sequencing performed by Next Generation Sequencing (NGS) or whole exome sequencing (WES) were performed as part of the formal diagnostic work-up in all patients. Written informed consent from parents/legal guardians and approval from the local ethical committees were obtained.

2.1. EEG recording and analysis

EEG signals were recorded through 19 silver-silver chloride electrodes placed over the scalp according to the International 10-20/10-10 system. In patient #1, data were exported to .edf (European data format) and further processed with MATLAB (The Math Works Inc., Natick, MA). An anti-aliasing low-pass FIR filter was applied and downsampling to 128 Hz was performed. Sleep stage scoring was based on 20 second epochs. Due to the extremely altered sleep-EEG pattern, sleep stage scoring was simplified in "wakefulness", "NREM-sleep" and "REM-like sleep", as in previous studies (Bölsterli et al., 2017, Pavlidis et al, 2019). Epochs disturbed by artifacts were eliminated. Spike amplitude and density, and electrode site showing spike activity leading to secondary bilateral synchrony were the criteria used to identify the EEG epileptic focus. A semi-automated spike search method implemented in the BESA software (BESA[®] Research 6.0) was used to calculate the spike-wave-index (SWI). Spikes were detected by template matching, in which an average of the visually identified, representative spikes for each EEG served as template. A 4-40 Hz zero phase band pass filter was applied to minimize the possible influence of sleep slow waves on the accuracy of template matching (Larsson et al., 2009). A virtual average montage over all channels was used for spike search with a correlation percentage of 80%. Calculation and graphical representation of the SWI was performed in a MATLAB environment (MATLAB, version 7.3.0 R2006b). In patients #3 and #4, SWI was assessed visually by expert neurophysiologists on paper-EEG; in these patients, SWI was assessed by calculating the amount of NREM sleep (in seconds) occupied by epileptic discharges divided by the total time of NREM sleep (in seconds) and then the ratio was transformed in percentage. In patients #2 and #5, a similar method was used on digital-EEG.

3. Results – Electro-clinical phenotyping (Table 1)

3.1. Patient 1

This Danish 5-years 8-months-old boy was born after an uneventful pregnancy, second child of healthy unrelated parents. He presented with breastfeeding problems and difficulty in swallowing since the first months of age. Developmental delay was first noticed at 5 months of age. When he was 20 months old, neurological examination showed poor language, gross motor function delay, poor balance due to diffuse hypotonia, joint hypermobility, and impaired walking, possible only with support. He had adequate fine motor function and he could manipulate toys with good coordination, using both hands with pincher grasp. His language consisted of several different words with meaning and he could pronounce two-word sentences; language understanding was overall good. At the age of 2 years, he started to suffer from nocturnal epileptic seizures, characterized by staring, complex motor manifestations and sometimes vomiting, often related to fever. Epilepsy was diagnosed at the age of 3 years 6 months, after the appearance of diurnal seizures, characterized by slurred speech without impairment of consciousness. Awake EEG disclosed multifocal 3-4 Hz spike-waves discharges. At this age, genetic tests (array-CGH, X-fragile) and metabolic screening were negative. Brain MRI was normal. At 3 years 7 months of age, fine motor functions worsened; in addition, he was diagnosed with speech and oro-motor dyspraxia. In the following years, physiotherapy, speech education and occupational therapy improved motor skills and language. At the age of 5 years, the detection of diffuse, continuous spikes/sharp waves at 2-2.5 Hz, with bilateral fronto-centroparietal predominance, during NREM sleep coupled with seizure aggravation, worsening of motor coordination, cognitive regression, lack of eye contact, and complete loss of speech and language comprehension led to the diagnosis of ESES. SWI was 28% during wakefulness, 93% and 56% during NREM and REM sleep respectively (Fig. 1A). Several anti-epileptic drugs (AEDs) failed to achieve seizure control; a course of oral prednisolone worsened impulsivity and hyperactivity. Four months later, a wakefulness/overnight EEG (Fig. 1B) confirmed the extreme activation of the frontocentral spike/polyspikes-and-wave discharges during sleep and it showed an increment of the epileptic abnormalities during wakefulness: SWI was 44% during wakefulness, 95% and 47% during NREM and REM sleep respectively. At the last follow-up at the age of 5 years 6 months, he had regained eye contact; fine motor skills and cognitive status were mildly improved with partial recovery of expressive language (mainly echolalia). Hypotonia, hyperkinetic behavior, apraxic gait and speech/oro-motor dyspraxia were unchanged. In addition, he presented with daily multiple atypical absences, with staring and arrest of ongoing activities up to 40-45 seconds. A wakefulness/overnight EEG showed a striking enhancement of the epileptic discharges during wakefulness with awake SWI up to 90%: NREM-SWI was 96% (Fig. 1C).

At 5 years of age, genetic analysis, performed by NGS with a targeted panel for 582 genes associated with epilepsy, intellectual disability or autism spectrum disorder, disclosed a novel *de novo* hemizygous mutation of *CNKSR2*, c.2024_2027delAGAG leading to p.(Glu675Glyfs*41) on exon 18.

3.2. Patient 2

This Spanish 21-year-old man, second child of healthy unrelated parents, was born after an uneventful pregnancy. He had a normal psychomotor development until the age of 3 years. At this age, he presented his first seizure characterized by a prolonged episode (1 hour) of unresponsiveness and upward eyes deviation. One month later, he started to suffer from seizures characterized by stiffening of the body followed by tonic-clonic manifestations, occurring during sleep or in the sleep/wakefulness transition. In the post-ictal recovery phase, an expressive aphasia was evident. EEGs showed abundant 2.5 Hz spike-wave discharges in the frontal regions, with right side predominance, both during wakefulness and sleep. After seizure onset, he started to present with hyperactive behavior. At 6 years of age, worsening of hyperactivity associated with a remarkable enhancement of epileptiform discharges with frontal predominance during NREM sleep (NREM-SWI > 90%) (Fig. 2), led to the diagnosis of ESES. Brain MRI was normal. At 8 years of age a neuropsychological evaluation showed an intelligent quotient (IQ) in the low average range with attention deficit possibly related to medications, poor visuomotor coordination with normal verbal/auditory memory and an estimated age of 5 years. Sleep-related seizures continued uncontrolled despite different combinations of eight different AEDs, corticosteroids and adrenocorticotropic hormone (ACTH) treatment until the age of 12 years. At this age, seizures disappeared; EEG normalized by the age of 13 years. At present, he is seizure free and out of medications; he still has hyperactive behaviors and a mild cognitive impairment, with reading and writing difficulties, despite logopedic treatment.

At 16 years of age, genetic testing by WES disclosed a novel variant of *CNKSR2* gene, c.246-247delAG, leading to a frameshift p.T83Kfs*30. Parents were asymptomatic and tested negative for the mutation.

3.3. Patient 3

This French 12-year-old boy, second child (out of three) of healthy unrelated parents, was born after an uneventful pregnancy. Mild developmental delay was diagnosed soon after birth. He was able to sit at 10 months of age and to walk without support at 28 months of age. He also presented with stereotypies and a language disorder that was diagnosed as speech dyspraxia. Myoclonic seizures during sleep/wakefulness transition appeared at the age of 2 years 6 months. Epilepsy was later diagnosed at 4 years of age after the appearance of clonic seizures. At the age of 7 years, he started to present also atypical absences with head drops. EEG at epilepsy onset showed poorly organized background activity with abundant bilateral fronto-temporal epileptiform discharges. At 5 years of age, his language was severely impaired (he could pronounce only single words). Brain MRI at this age was normal. At 6 years 6 months of age, worsening of behavior associated with the appearance of continuous, bilateral fronto-temporal spikewaves during sleep (NREM-SWI > 80%) (Fig. 3), was consistent with the diagnosis of ESES. Language further regressed at the age of 8 years. Two prolonged episodes of absence status epilepticus occurred at the age of 7 years 6 months and 11 years, respectively. At present, he still suffers from uncontrolled seizures. At the last follow-up at the age of 12 years, he had moderate/severe neurodevelopmental delay (neuropsychological testing was not possible due to lack of cooperation) and his language was severely impaired (he could pronounce about 20 single words), with minimal improvement after speech therapy. Sleep EEG showed an overt decrease of epileptic discharges during NREM sleep, compatible with the remission of ESES. A trio-based WES, at 12 years of age, disclosed a novel deletion on chromosome X (c.457_461del) of CNKSR2 gene leading to a pathogenic frameshift variant (p.Tyr153-Serfs*5). Segregation analysis of the parents disclosed a maternal mosaicism (5/156 reads) in the asymptomatic mother.

3.4. Patient 4

This French 41-years-old woman is the only child of healthy unrelated parents. She normally developed until 6 years 3 months of age, when she started suffering from tonic-clonic seizures. A treatment with valproic acid was initiated. At the age of 7 years, atypical absences with falls and cognitive disturbances such as difficulties in reading, writing and understanding complex instructions appeared. On Wechsler Preschool and Primary Scale of Intelligence (WPPSI) testing, she had a full-scale IQ of 62, verbal IQ of 66 and performance IQ of 57. Awake EEG showed generalized spike-wave discharges and right temporal spikes. At 8 years of age, complete loss of language, apraxia and unresponsiveness to external noises and episodes of absence status epilepticus were reported. Moreover, she could alternate periods of apathy with periods of hyperactivity and compulsive behavior. At this age, a striking activation of bilateral spike-and-wave discharges during sleep (NREM-SWI > 60%) led to the diagnosis of ESES. Brain MRI was normal. At



Fig. 1. Wakefulness and NREM sleep EEG recording in patient #1. On the left EEG tracing in wakefulness and NREM sleep are shown. On the right, graphs illustrating the SWI in wakefulness and sleep are reported during 18-hour recording (start of recording: 2 p.m.; end of recording 8 a.m. the day after). (A) At ESES onset (age 5 years) wakefulness EEG showed multifocal bilateral spike/spike-waves discharges (wakefulness SWI: 28%). During NREM sleep, diffuse 2.5 Hz spike-wave activity, with predominance over the bilateral fronto-centro-parietal regions (NREM-SWI: 93%) appeared. (B) First follow-up (age 5 years, 4 months): during wakefulness, striking activation of 2.5 Hz spike-wave activity, predominant in fronto-central regions and on the right side (awake SWI: 44%). During sleep, the NREM-SWI is 95%. (C) At the last follow-up (age 5 years 6 months): epileptic activity during wakefulness was further enhanced (awake SWI: 90%) whereas during NREM sleep SWI was unchanged as compared to the previous recordings (NREM-SWI: 96%). The SWI was calculated by using a semi-automated spike search method implemented in the BESA software (BESA[®] Research 6.0) (Larsson et al., 2009). Leggent: EEG: Electroencephalogram; ESES: encephalopathy related to status epilepticus during slow sleep; m: months; NREM: non-rapid eye movement sleep; Pt.: patient; SWI: spike-wave index; y: years; Wkf: wakefulness.

 Table 1

 Clinical, EEG and genetic features of our cohort and already published patients affected by CNKSR2-related ESES.

Patient	Patient #1	Patient #2	Patient #3	Patient #4	Patient #5	Sun et al., 2018	Damiano et al.,	Damiano	Vaags et al.,	Vaags et al.,	Vaags et al.,
							2017*	et al., 2017*	2014**	2014**	2014
Gender/Age Origin	M / 5y8m Danish	M / 21y Spanish	M / 12y French	F / 41y French	M / 9y8m Spanish	M / 8y8m Chinese	M / 18y Ashkenazi Jews	M / 12y Ashkenazi Jews	M / 6y Canadian	M / 8y Canadian	M / 8y Norwegian
Mutation	c.2024_2027delAGAG, p.Glu675Glyfs*41	c.246-247delAG, p.T83Kfs*30	c.457_461del, p. Tyr153Serfs*5	Deletion Xp22.12 (21523673– 21558329) ^{#§}	Deletion Xp22.12 (21609392– 21619786) [#]	c.2185C > T, p. Arg729*	c.2314C > T, p. Arg712*	c.2314C > T, p.Arg712₊	Deletion Xp22.12 (20,297,696– 21,471,387) [#]	Deletion Xp22.12 (20,297,696– 21,471,387) [#]	Deletion Xp22.12 (21,375,312– 21,609,484) [#]
Occurrence	De novo	De novo	Maternal mosaicism (5/156 reads)	Parents unavailable	De novo	De novo	Maternal (unaffected)	Maternal (unaffected)	Maternal (unaffected)	Maternal (unaffected)	Maternal (unaffected)
Sz onset - age	2y Sleep-related Sz with	3y Prolonged episode	4y TCS	бу тсs	3y2m Staring with	~ 2y Episodes of	3y6m	3y6m	2y Sleep-related	2y Sleep-related	2y6m Staring
52 type at onset	staring and complex motor behavior	of unresponsiveness and upward eyes deviation			bilateral "tremors" of upper limbs	loss of consciousness, staring, limbs jerks		NA NA	TCS	TCS	spells
Other Sz types	Episodes of slurred speech without impairment of consciousness	Sleep-related Sz with body stiffening evolving to TCS	Myoclonic Sz, atypical absences with head drops, abs-SE	Atypical absences with falls, abs-SE	Loss of tone followed by confusion/ agitation. TCS, hemiclonic seizures +/- loss of awareness	No	NA	NA	No	TCS	TCS
Sz outcome	Multiple daily atypical absences	12y: seizure free	Atypical absences with head drops	12y: seizure free	9y8m: 1–2 Sz/y	Improvement	NA	NA	Seizure free	Seizure free	Seizure free
WKF-EEG	3y6m: multifocal SW	3y: spikes, SW in L-O and biFr regions	4y6m: normal. 3y: disorganized BGA, multifocal spikes, biFr-T SW	6y3m: R-T spikes and GSW	2y10m: normal BGA. triphasic high-voltage spikes (L Fr- C-T and R F- T region)	NA	NA	NA	NA	NA	NA
ESES onset – age	5y	6y	6y6m	~ 8y	2y10m	NA	~ 4y	3y6m	2y	~ 3y	7y
ESES topography	5y: biFr	>90% biFr (R > L)	>80% biFr-T (L > R)	>60% Diffuse (R > L)	biFr-T	biT and Fr	biC-T or Fr	biC-T or Fr	80–100% Fr-T	∽80% Fr-T	INA Fr-T
ESES evolution	Sy6m: diffuse Ongoing	13y: remission	12y: remission	12y: remission	(R > L) 9y8m: ongoing	9y8m: speech improvement. No EEG.	NA	NA	4y10m: NDD. EEG: SWI 90%	NA	NA
AEDs [∞]	OXC, VPA, OCS, <u>CLB,</u> STM, <u>LEV</u>	ETS, LTG, CBZ, OXC, LEV, PHT, VPA, CLB, OCS, ACTH	CLB, <u>VPA</u> , <u>LEV</u> , ZNS, CNZ	VPA, ETS, <u>ICS.</u> <u>OCS</u>	VPA, LEV, CLB, ETS, STM, ZNS, RUF, LTG, PER, LCS, OCS, KD	IgIV, OCS, LTG, VPA, LEV	NA	NA	VPA, DZP, OCS	LTG, CBZ, <u>VPA</u>	<u>LTG</u>
ND before Sz onset	Mild NDD. 20 m: hypotonia, gross motor delay, speech/ oro-motor dyspraxia, instability	Normal	Mild NDD. 10 m: sitting; 20 m: walk with support; 28 m: walk without support	Normal	Moderate NDD. 20m: hypotonia, language delay	NDD, ADHD	NDD and language delay	NDD and language delay	Mild NDD	Mild NDD	Mild NDD

Table 1 (continued)

Patient	Patient #1	Patient #2	Patient #3	Patient #4	Patient #5	Sun et al., 2018	Damiano et al., 2017*	Damiano et al., 2017*	Vaags et al., 2014 ^{**}	Vaags et al., 2014 ^{**}	Vaags et al., 2014
Cognition, language and behavior before ESES onset	Mild NDD. Hypotonia, gross motor delay, speech/oro-motor dyspraxia, instability. 3y6m: fine motor delay	Hyperactivity	Stereotypies, speech dyspraxia. 5y: 50 single words	Mild NDD. Difficulties in reading, writing, comprehension	Moderate NDD. 20m: hypotonia, language delay	ND regression	NDD and language delay. Attention deficit, hyperactivity	NDD and language delay. Attention deficit, hyperactivity	Mild NDD	Language regression	6y: single words ADHD
Cognition, language and behavior after ESES onset	Worsening of motor coordination, cognitive regression, complete loss of speech and language comprehension, lack of eye contact	ND regression	Moderate/severe NDD. 8y: language regression, 20 single words	ND regression. 10y: dysarthria, aphasia, bucco- lingual-facial dyspraxia, hyperactivity 12y: WISC-R: FSIQ 40.	3y: language delay (3–5 words), dyslalia, mild orolingual dyspraxia	Further ND regression. No speech, ASD	4y: ND regression. No speech	12y: useful language, attendance at regular school	ND Regression. 5y: no speech. Hyperactivity	ND regression. 4y: no speech. Hyperactivity	Severe NDD, inattention, impulsivity
Long-term evolution after ESES remission	NA	Mild NDD. 21y: specific disorder of reading and writing. Hyperactivity	NA	29y: WISC-R: FSIQ 54. 41y: basic reading and mathematics. Family dependent. Limited social life	NA	NA	18y: institutionalized	NA	NA	NA	NA
MRI scanning	Normal (3y)	Normal (6y)	Normal (5y)	Normal (8y)	Normal (3y)	Normal	NA	NA	Normal	Normal	Normal

Legend: abs-SE: absence status epilepticus; ACTH: Adrenocorticotropic hormone; ADHD: attention deficit hyperactivity disorder; AEDs: anti-epileptic drugs; ASD: autism spectrum disorder; BGA: background activity; bi-: bilateral; C: central; CBZ: carbamazepine; CLB: clobazam; CNZ: clonazepam; DZP: diazepam; EEG: Electroencephalogram; ESES: encephalopathy related to status epilepticus during slow sleep; ETS: ethosuximide; F: female; Fr: frontal; FSIQ: full scale intelligence quotient; F-UP: follow-up; GSW: generalized spike-waves; ICS: intravenous corticosteroids; IgIV: intravenous immunoglobulin; KD: ketogenic diet; L: left; LCS: lacosamide; LEV: levetiracetam; LTG: lamotrigine; M: male; m: months; MRI: magnetic resonance imaging; NA: not available; ND: neurodevelopment; NDD: neurodevelopmental delay; NREM: non-rapid eye movement sleep; O: occipital; OCS: oral corticosteroids; OXC: oxcarbazepine; PER: perampanel; PHT: phenytoin; R: right; RUF: rufinamide; STM: sulthiame; SW: spike-waves; SWI: spike-wave index; Sz: seizures; T: temporal; TCS: tonic-clonic seizures; VPA: valproic acid; y: years; WISC-R: Wechsler Intelligence Scale for Children Revised; WKF: wakefulness; ZNS: zonisamide.

Patients belong to the same family.

** Patients belong to the same family.

[#] Human Genome version 19 by UCSC genome browser.

[§] Genetic data already published by Lesca et al. (2012).

[®] AEDs with reported best efficacy are underlined. Current AEDs at last available evaluation are highlighted in bold.



Fig. 2. Wakefulness and NREM sleep EEG recording in patient #2. At the age of 6 years, wakefulness EEG (left panel) showed sporadic bifrontal high amplitude spike-waves. During NREM sleep (right panel), exaggeration of bifrontal spike/spike-wave discharges with right side predominance (NREM-SWI: >90%). Legend: EEG: electroencephalogram; ESES: encephalopathy related to status epilepticus during slow sleep; NREM: non-rapid eye movement sleep; Pt.: patient; y: years; Wkf: wakefulness.

9 years of age, EEG showed frequent short bursts (1-2 sec) of generalized spike-and-waves during wakefulness and unchanged NREM-SWI. Short courses of intravenous hydrocortisone until the age of 10 years improved transitorily her cognitive status and behavior. EEG recording showed a decrement of epileptic abnormalities during wakefulness, whereas NREM-SWI was still around 60%. At 10 years of age, she further deteriorated presenting with dysarthria, oro-lingual-facial dyspraxia and severe aphasia. After a further cycle of prednisolone, her clinical status and EEG progressively improved. By the age of 11 years 11 months, the sleep EEG was normal: neuropsychological tests showed persistence of cognitive deficits (at Wechsler Intelligence Scale for Children Revised (WISC-R) testing: full-scale IQ 40, verbal IQ 49, performance IQ 45). No data on physiotherapy or rehabilitation are available. At age of 12 years 11 months, both awake and sleep EEG were normal. She is seizure free without medications since the age of 16 years. A neuropsychological assessment at the age of 29 years demonstrated moderate intellectual disability (at WISC-R testing: full-scale IQ 54, verbal IQ 60, performance IQ 52). At the last follow-up at the age of 41 years, she had very basic reading and mathematics abilities, and she was dependent on her family with limited social life. At the age of 34 years, an array-CGH disclosed a copy number change of 35 kb size on Xp22.12, ChrX:21523673-21558329 causing the loss of part of CNKSR2 gene (genetic findings were previously published by Lesca et al., 2012). Parents were not available for testing.

3.5. Patient 5

This Spanish 9-years 8-months old boy was born at 36 gestational weeks from C-section after a bicorial biamniotic pregnancy. His mother suffered from epilepsy since the age of 18 years, controlled by antiepileptic treatment, and a borderline behavioural disorder. Soon after birth, mild generalized hypotonia became evident associated with unspecific developmental delay. He walked at 20 months of age; at this age, he presented with attention deficit, and a severe language delay, characterized by only few single words and mild oro-lingual dyspraxia. First overnight-EEG, performed at 2 years 10 months of age, as part of formal work-up for the language delay showed during wakefulness a normal background with frequent runs of triphasic high-voltage spikes, occurring asynchronously in left fronto-centro-temporal and right fronto-temporal regions, with slightly right predominance. During sleep this activity became almost continuous, with a NREM-sleep SWI > 80% (Fig. 4). Valproic acid was started with some improvement of the attention disorder and no effect on language. At 3 years of age, he used about 3-5 referential two-syllable words with abundant dyslalia and mild oro-lingual dyspraxia. At the age of 3 years 2 months, he started to present sporadic and brief (up to 10 seconds) staring episodes without other associated signs. At this age a brain MRI was normal. Epilepsy was diagnosed at the age of 3 years 6 months after a seizure during sleep/wakefulness transition characterized by staring and bilateral "tremors" of the upper extremities. A neuropsychological evaluation at that time showed a global IQ of 46 (language 39, coordination 50, socialization 43). At 4 years of age a neuropsychological evaluation estimated a developmental age corresponding to 1 year. Since this age, despite the association of different treatments (clobazam, ethosuximide, sulthiame, zonisamide, rufinamide, lamotrigine, perampanel, lacosamide, corticosteroids and ketogenic diet) he continued to have weekly generalized tonic-clonic seizures, and hemiclonic seizures (on both sides) with or without loss of awareness, predominantly during sleep or awakening. Over the years, seizure frequency progressively decreased, and since the age of 7 years he is seizure free on valproic acid. Overnight EEGs continue to show a NREM SWI > 80%. At last follow-up at the age of 9 years, he presented motor aphasia, mild oro-lingual dyspraxia, and abnormal behaviour characterized mainly by impulsivity and irritability. He needed support to attend to a special school and for daily activities, and he continued speech therapy.

At 5 years of age, an array-CGH showed a deletion of 10 kb on Xp22.12, ChrX:21609392-21619786 involving the region of *CNKSR2* gene. Parents were tested and found negative.

4. Discussion

In this study we report the electro-clinical phenotypes of five novel patients with *CNKSR2*-related ESES, comparing them with



Fig. 3. Wakefulness and NREM sleep EEG recording in patient #3. At the age of 6 years 6 months, wakefulness (left panel) and NREM sleep EEG (right panel) in patient #3. During wakefulness (left panel), EEG disclosed multifocal spike-wave discharges. During NREM sleep (right panel), striking increment of bilateral fronto-temporal spike-wave discharges, with left predominance (NREM-SWI: >80%). Legend: EEG: electroencephalogram; ESES: encephalopathy related to status epilepticus during slow sleep; m: months; NREM: non-rapid eye movement sleep; Pt.: patient; y: years; Wkf: wakefulness.

previously published cases (Vaags et al., 2014; Damiano et al., 2017; Sun et al., 2018) (Table 1). Three of our patients (#1, #3, #5) presented with neurodevelopmental delay with severe language impairment since early infancy, as already described (Vaags et al., 2014; Damiano et al., 2017; Sun et al., 2018), whereas, the other two patients (#2, #4) developed normally until the onset of epilepsy at the age of 3 and 6 years respectively. Seizure onset was associated with the appearance (#2, #4) or aggravation (#1, #3, #5) of cognitive and/or behavioral disturbances. During the course of the disease, a further worsening of the cognitive and behavioral disorders was observed in coincidence with the detection of strikingly enhanced sleep-related EEG epileptiform discharges, leading to the diagnosis of ESES. In patients #2 and #4, long-term follow-up showed the remission of epilepsy and mild/moderate cognitive impairment, indicating that CNKSR2-related ESES can present with a phenotype milder than previously reported in adult patients harboring CNKSR2 pathogenic variants and presumed ESES in childhood (Vaags et al., 2014).

Bilateral diffuse, or unilateral, more or less focal (frontal, centrotemporal, parietal, occipital), subcontinuous epileptic discharges during sleep are the EEG features defining ESES (Tassinari et al., 2019). In 4/5 of our patients, topography of sleep-related EEG epileptic discharges during ESES was primarily frontal with spreading to central or temporal regions, in agreement with previous observations in patients with CNKSR2-related ESES (Table 1). This finding may suggest that the involvement of frontal areas by the epileptic activity, in particular during sleep, might be a distinctive EEG feature of CNKSR2-related ESES, at variance with ESES associated with GRIN2A pathogenic variants, in which the EEG focus is most often located more posteriorly, in centro-parietal regions (Gabrielle Rudolf, personal communication). Indeed, a recent paper has shown in two patients with GRIN2A-related ESES, a multifocal (parietal and temporo-parietal) topography of the EEG focus (Pavlidis et al., 2019). However, at present the data are very scanty to establish whether there are distinctive EEG patterns in the different genetically-determined ESES. The lack of a systematic neuropsychological assessment in our patients renders difficult to establish whether the type of cognitive/behavioral disorders may be related to the predominant frontal topography of sleep-related epileptic discharges or alternatively may depend on a more diffuse brain dysfunction, possibly related to impaired sleep homeostasis, as recently suggested (Pavlidis et al., 2019, Rubboli et al., 2019) as well as mediated by remote inhibition mechanisms (De Tiege et al., 2008).

All previously published patients harboring CNKSR2 pathogenic variants are reported to suffer from language impairment, not further specified. Four out of five of our subjects presented with speech and/or oro-motor dyspraxia, unreported yet in CNKSR2-related disorders. Speech dyspraxia refers to impaired motor planning and programming whereas oro-motor dyspraxia impairs speech execution (Turner et al., 2015). As observed in GRIN2Arelated EAS conditions (Turner et al., 2015), the combination of these speech dysfunctions could account for the severe language disturbance also in CNKSR2-related disorders. The observation of these speech disturbances in three patients (#1, #3, #5) before the diagnosis of ESES may suggest that speech/oro-motor apraxia may be a feature of CNKSR2-related encephalopathy rather than directly linked to ESES. These speech deficits, reported in all published patient with CNKSR2 pathogenic variants, suggest that CNKSR2 plays a role in speech production. In fact, CNKSR2 has been shown to be expressed, since the prenatal age, in structures such as cerebellum and nucleus caudatus which are involved in speech production (Liegeois and Morgan, 2012).

Episodes of prolonged absence status epilepticus or aggravation of atypical absences have been observed in 3/5 patients (#1, #3, #4), in concomitance with the diagnosis of ESES. Based on previous data suggesting that worsening of epileptic seizures might herald the onset of ESES (Saltik et al., 2005), aggravation of epilepsy in patients harboring CNKSR2 pathogenic variants should prompt clinicians to perform a sleep EEG to verify the occurrence of an ESES EEG pattern. Interestingly, in patient #1 aggravation of atypical absences was associated with a striking increment of EEG epileptic discharges during wakefulness (awake SWI: 90%), with persistence of continuous spike-wave activity in between the absence episodes. Further studies may clarify whether modifications/increase of awake SWI may correlate with epilepsy worsening during the course of ESES. It is noteworthy that the increment of epileptic EEG activity during wakefulness was not accompanied by further worsening of the cognitive and behavioral disturbances. This is in keeping with data suggesting that cognitive derangement in ESES might be linked to the impairment, caused by exaggerated EEG epileptic activity during NREM sleep, of physiological sleep-related cortical plasticity processes that underlies learning and memory consolidation, particularly in the developmental age (Tononi and Cirelli, 2014; Rubboli et al., 2019).

Herein we describe the first female (#4) with *CNKSR2*-related ESES. Previously three female patients with *CNKSR2* pathogenic variants were described (Damiano et al. 2017, Polla et al. 2019)



Fig. 4. Wakefulness and NREM sleep EEG recording in patient #5. At the age of 3 years, wakefulness EEG (left panel) showed high amplitude triphasic high-voltage spikes, in the right fronto-temporal regions and sporadic runs of delta activities in bifrontal regions were detectable. NREM sleep (right panel) was characterized by a remarkable enhancement of spike-wave activity (SWI > 80%), with right fronto-central/fronto-temporal predominance. Legend: EEG: electroencephalogram; ESES: encephalopathy related to status epilepticus during slow sleep; NREM: non-rapid eye movement sleep; Pt.: patient; y: years; Wkf: wakefulness.

associated with a mild phenotype of various different types of seizures with or without intellectual disability. One of them, sister of two brothers with *CNKSR2*-related ESES suffered from benign epilepsy with centro-temporal spikes, part of the spectrum of EAS (Damiano et al., 2017). Recently, an increasing number of Xlinked conditions initially identified in affected males have also been reported in severely affected females (such us for instance *KIAA2022, IQSEC2*-related disorders, recently published by de Lange et al., 2016; Mignot et al., 2019), suggesting that additional mechanisms may contribute to the pathogenesis of these disorders other than those related to the classical heterozygous inheritance.

In conclusion, with this study, we expand the phenotypic spectrum of *CNKSR2*-related ESES including, as novel findings, occurrence also in females, normal neurodevelopment before epilepsy onset, mild/moderate intellectual disability in adulthood, epilepsy aggravation associated with enhanced awake SWI and speech/oromotor dyspraxia as the cause of language impairment. Additional studies may corroborate our findings helping clinicians in providing a correct and timely diagnosis.

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Declaration of Competing Interest

None of the authors have potential conflicts of interest to be disclosed.

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Author's contributions

CMB and GR reviewed the literature, performed clinical data collection and drafted the manuscript. CM, JMS, GL, BGG, RM, GR, KMJ, PMG contributed in collecting clinical data from patient's chart and from the literature. EG and CR revised and analyzed electrophysiological data. All authors revised the manuscript and approved the final article. GR, EG, RSM, CAT conceived the idea of the present paper, critically reviewed and finally approved the manuscript.

References

- Aypar U, Wirrell EC, Hoppman NL. CNKSR2 deletions: a novel cause of X-linked intellectual disability and seizures. Am J Med Genet A 2015;167:1668–70. <u>https://doi.org/10.1002/ajmg.a.36902</u>.
- Bölsterli BK, Gardella E, Pavlidis E, Wehrle FM, Tassinari CA, Huber R, et al. Remission of encephalopathy with status epilepticus (ESES) during sleep renormalizes regulation of slow wave sleep. Epilepsia 2017;58:1892–901. https://doi.org/10.1111/epi.13910.
- Bumeister R, Rosse C, Anselmo A, Camonis J, White MA. CNK2 couples NGF signal propagation to multiple regulatory cascades driving cell differentiation. Curr Biol 2004;14:439–45. <u>https://doi.org/10.1016/j.cub.2004.02.037</u>.
- Damiano JA, Burgess R, Kivity S, Lerman-Sagie T, Afawi Z, Scheffer IE, et al. Frequency of CNKSR2 mutation in the X-linked epilepsy-aphasia spectrum. Epilepsia 2017;58:e40–3. <u>https://doi.org/10.1111/epi.13666</u>.
- de Lange IM, Helbig KL, Weckhuysen S, Møller RS, Velinov M, Dolzhanskaya N, et al. De novo mutations of KIAA2022 in females cause intellectual disability and intractable epilepsy. J Med Genet 2016;53:850–8. <u>https://doi.org/10.1136/ imedgenet-2016-103909</u>.
- De Tiege X, Ligot N, Goldman S, Poznanski N, de Saint Martin A, Van Bogaert P. Metabolic evidence for remote inhibition in epilepsies with continuous spikewaves during sleep. NeuroImage 2008;40:802–10. <u>https://doi.org/10.1016/j.</u> neuroimage.2007.11.043.
- Hu H, Haas SA, Chelly J, Van Esch H, Raynaud M, de Brouwer AP, et al. X-exome sequencing of 405 unresolved families identifies seven novel intellectual disability genes. Mol Psychiatry 2016;21:133–48. <u>https://doi.org/10.1038/</u> mp.2014.193.
- Kessi M, Peng J, Yang L, Xiong J, Duan H, Pang N, et al. Genetic etiologies of the electrical status epilepticus during slow wave sleep: systematic review. BMC Genet 2018;19:40. <u>https://doi.org/10.1186/s12863-018-0628-5</u>.
- Lanigan TM, Liu A, Huang YZ, Mei L, Margolis B, Guan K-L. Human homologue of drosophila CNK interacts with Ras effector proteins Raf and Rlf. FASEB J 2003;17:2048-60. <u>https://doi.org/10.1096/fj.02-1096com</u>.

- Larsson PG, Wilson J, Eeg-Olofsson O. A new method for quantification and assessment of epileptiform activity in EEG with special reference to focal nocturnal epileptiform activity. Brain Topogr 2009;22:52–9. <u>https://doi.org/ 10.1007/s10548-008-0072-3</u>.
- Lesca G, Rudolf G, Labalme A, Hirsch E, Arzimanoglou A, Genton P, et al. Epileptic encephalopathies of the Landau-Kleffner and continuous spike and waves during slow-wave sleep types: genomic dissection makes the link with autism. Epilepsia 2012;53:1526–38. <u>https://doi.org/10.1111/j.1528-1167.2012.03559</u>.
- Lesca G, Møller RS, Rudolf G, Hirsch E, Hjalgrim H, Szepetowski P. Update on the genetics of the epilepsy-aphasia spectrum and role of GRIN2A mutations. Epileptic Disord 2019;31. <u>https://doi.org/10.1684/epd.2019.1056</u>.
- Liegeois F, Morgan A. Neural basis of childhood speech disorders: lateralization and plasticity for speech function during development. Neurosci Biobehav Rev 2012;36:439–58. <u>https://doi.org/10.1016/j.neubiorev.2011.07.011</u>.
- Liu L, Channavajhala PL, Rao VR, Moutsatsos I, Wu L, Zhang Y, et al. Proteomic characterization of the dynamic KSR-2 interactome, a signaling scaffold complex in MAPK pathway. Biochim Biophys Acta 2009;1794:1485–95. <u>https://doi.org/10.1016/i.bbapap.2009.06.016</u>.
- Mignot C, McMahon AC, Bar C, Campeau PM, Davidson C, Buratti J, et al. IQSEC2related encephalopathy in males and females: a comparative study including 37 novel patients. Genet Med 2019;21:837–49. <u>https://doi.org/10.1038/s41436-018-0268-1</u>.
- Nagase T, Ishikawa K, Suyama M, Kikuno R, Hirosawa M, Miyajima N, et al Prediction of the coding sequences of unidentified human genes. XII. The complete sequences of 100 new cDNA clones from brain which code for large proteins in vitro. DNA Res. 1998;5:355–64.
- Ohtakara K, Nishizawa M, Izawa I, Hata Y, Matsushima S, Taki W, et al. Densin-180, a synaptic protein, links to PSD-95 through its direct interaction with MAGUIN-1. Genes Cells 2002;7:1149–60. <u>https://doi.org/10.1046/j.1365-2443.2002.00589.x</u>.
- Patry G, Lyagoubi S, Tassinari CA. Subclinical electrical status epilepticus induced by sleep in children. A clinical and electroencephalographic study of six cases. Arch Neurol 1971;24:242–52.
- Pavlidis E, Møller RS, Nikanorova M, Sophie Kölmel MS, Stendevad P, Beniczky S, et al. Idiopathic encephalopathy related to status epilepticus during slow sleep (ESES) as a "pure" model of epileptic encephalopathy. An electro-clinical, genetic and follow-up study. Epilepsy Behav 2019;97:244–52. <u>https://doi.org/</u> 10.1016/j.vebeh.2019.05.030.

- Polla DL, Saunders HR, de Vries BBA, van Bokhoven H, de Brouwer APM. A de novo variant in the X-linked gene CNKSR2 is associated with seizures and mild intellectual disability in a female patient. Mol Genet Genomic Med 2019;7: e00861. <u>https://doi.org/10.1002/mgg3.861</u>.
- Rubboli G, Huber R, Tononi G, Tassinari CA. Encephalopathy related to Status Epilepticus during slow Sleep: a link with sleep homeostasis?. Epileptic Disord 2019;21:62–70. <u>https://doi.org/10.1684/epd.2019.1059</u>.
- Saltik S, Uluduz D, Cokar O, Demirbilek V, Dervent A. A clinical and EEG study on idiopathic partial epilepsies with evolution into ESES spectrum disorders. Epilepsia 2005;46:524–33.
- Sun Y, Liu YD, Xu ZF, Kong QX, Wang YL. CNKSR2 mutation causes the X-linked epilepsy-aphasia syndrome: a case report and review of literature. World J Clin Cases 2018;6:570–6. <u>https://doi.org/10.12998/wjcc.v6.i12.570</u>.
- Tassinari CA, Dravet C, Roger J. Encephalopathy related to electrical status epilepticus during slow sleep. Electroenceph Clin Neurophysiol 1977;43:529–30.
- Tassinari CA, Cantalupo G, Dalla Bernardina B, Darra F, Bureau M, Cirelli C, et al. Encephalopathy related to status epilepticus during slow sleep (ESES) including Landau-Kleffner Syndrome. In: Bureau M, Genton P, Dravet C, Delgado-Escueta A, Tassinari CA, Thomas P & Wolf P, editors. Epileptic Syndromes in Infancy, Childhood and Adolescence, 6th ed. John Libbey Eurotext Ltd; 2019, p. 261–83.
- Therrien M, Wong AM, Rubin GM. CNK, a RAF-binding multidomain protein required for RAS signaling. Cell 1998;95:343–53.
- Tononi G, Cirelli C. Sleep and the price of plasticity: from synaptic and cellular homeostasis to memory consolidation and integration. Neuron 2014;81:12–34. <u>https://doi.org/10.1016/i.neuron.2013.12.025</u>.
- Turner SJ, Mayes AK, Verhoeven A, Mandelstam SA, Morgan AT, Scheffer IE. GRIN2A: An aptly named gene for speech dysfunction. Neurology 2015;84:586–93. https://doi.org/10.1212/WNL.000000000001228.
- Vaags AK, Bowdin S, Smith ML, Gilbert-Dussardier B, Brocke-Holmefjord KS, Sinopoli K, et al. Absent CNKSR2 causes seizures and intellectual, attention, and language deficits. Ann Neurol 2014;76:758–64. <u>https://doi.org/10.1002/ ana.24274</u>.
- Wellbrock C, Karasarides M, Marais R. The RAF proteins take centre stage. Nat Rev Mol Cell Biol 2004;5:875–85.
- Yao I, Ohtsuka T, Kawabe H, Matsuura Y, Takai Y, Hata Y. Association of membraneassociated guanylate kinase-interacting protein-1 with Raf-1. Biochem Biophys Res Commun 2000;270:538–42. <u>https://doi.org/10.1006/bbrc.2000.2475</u>.