Joubert Syndrome Presenting With Normal Pyramidal Decussation: A Case Report

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Joubert syndrome (JS) is a rare genetic disorder characterized by a congenital malformation of the hindbrain, and accompanied by axonal decussation abnormalities affecting the corticospinal tract and the superior cerebellar peduncles. To the best of our knowledge, there are no reports of normal pyramidal decussation in JS. Here, we describe the case of an 18-year-old boy presenting midline-crossing corticospinal projections, which were considered normal corticospinal tract trajectories. Diffusion tensor imaging and motor evoked potential study analysis demonstrated the exclusive presence of decussating corticospinal projections in the patient. Based on these results, we suggest that JS might be associated with several, diverse corticospinal motor tract organization patterns.

Keywords Evoked potentials, Developmental disabilities, Joubert syndrome

INTRODUCTION

A congenital hindbrain malformation, Joubert syndrome (JS) is associated with axonal decussation abnormalities affecting the corticospinal tract (CST) and superior cerebellar peduncles [1]. A post-mortem neuropathological study has shown that pyramidal decussation is completely absent in patients with JS [2]. Recently, diffusion tensor imaging (DTI) analyses have also confirmed that the CST fails to decussate in JS [3]. Moreover, a neurophysiological study using transcranial magnetic stimulation (TMS) showed that individuals with JS display strong ipsilateral CST projections [4]. Taken together, these findings suggest that pyramidal tract fibers do not cross the midline in JS.

In this report, however, we present an exceptional case of JS characterized by only normally decussating CST projections.

CASE REPORT

An 18-year-old boy presented with a wide-based, unsteady gait. He had difficulty ambulating on uneven surfaces. Additionally, he presented noticeable oculomotor apraxia and mental retardation. His preferred hand was right, but he displayed severe clumsiness in both hands and had low hand strength. Mirror movements were not observed in either hand. Furthermore, the patient had
displayed hypotonia during infancy, and presented a
global developmental delay. Brain MRI demonstrated the
presence of the molar tooth sign and vermian hypoplas-
ia, typical of JS (Fig. 1A–1C).

We performed whole exome sequencing to look for the
19 genes associated with JS, namely: NPHP1, CEP290,
AHI1, TMEM67 (MKS3), RPRGIP1L, CC2D2A, ARL13B,
INPP5E, OFD1, TMEM216, KIF7, TCTN1, TCTN2,
TMEM237, CEP41, TMEM138, C5orf42, TMEM231, and
TCTN3. Genomic DNA was extracted from peripheral
blood leukocytes using the Wizard Genomic DNA Puri-
fication Kit, as per the manufacturer’s instructions (Pro-
mega, Madison, WI, USA). SureSelect Human All Exon V5
(Agilent Technologies, Santa Clara, CA, USA) was used
for library preparation, and sequencing was done on the
Illumina NextSeq 500 platform (Illumina Inc., San Diego,
CA, USA), generating 2×150 bp paired-end reads.

Results showed that the patient carried two com-
pound-heterozygous mutations, a frameshift mutation
(c.3557delA [p.Lys1186Argfs*22]) and a missense muta-
tion (c.4277T>C [p.Ile1426Thr]) in the C5orf42 gene.
Both these variants in C5orf42 were variants of uncertain
significance, and have not been reported previously.
Sanger sequencing confirmed these two variants in the
C5orf42 gene (Fig. 1D). In the original family described
by Joubert, Srour et al. [5] had reported that mutation of
C5orf42 gene was the cause of JS, and the phenotype of
this gene most closely resembled pure or classic JS.

DTI data were acquired using a Verio 3.0T system (Sie-
mens, Erlangen, Germany) equipped with a 12-chan-

Fig. 1. Axial T2-weighted (A) and
T2-weighted fluid-attenuated
inversion recovery (B) brain mag-
netic resonance imaging scans at
the pontomesencephalic junction
show the molar tooth sign (arrow).
The midsagittal T1-weighted im-
age (C) shows hypoplasia of the
cerebellar vermis (arrowhead).
Sanger sequencing chromatograms verified c.3557delA and
c.4277T>C in the C5orf42 gene
(D). The black indicates a guanine,
red indicates a thymine, blue in-
dicates a cytosine, and green indi-
cates an adenine.

Fig. 2. Axial color-coded map at the level of the pontomesencephalic junction shows the absence of the midline ‘focal
red dot’ which is interpreted as the absent decussation of the fiber tracts within the superior cerebellar peduncles (A).
The transpontine fiber shows no abnormality at the basis pontis (B, C, D). The red indicates a predominant left-right
(x-element), green indicates an anteroposterior (y-element), and blue indicates a superior-inferior (z-element) orien-
tation of the anisotropic component of diffusion within each voxel.
nel sensitivity encoding (SENSE) head coil for single-shot echo planar imaging. The imaging parameters used were: echo time=93 ms, repetition time=7,900 ms, field of view=230 mm×230 mm, sampling matrix size=128×128 reconstructed with homodyne processing to 256×256, SENSE factor=3, EPI=128, and b-value=1,000 s/mm². We acquired 47 contiguous, 3.0 mm thick slices parallel to the anterior commissure-posterior commissure line in 30 different diffusion directions.

DTI data analysis revealed absence of the midline ‘focal red dot’ at the level of the pontomesencephalic junction, representing the absent decussation of the superior cerebellar peduncle (Fig. 2A). At the basis pontis, the dorsal component of the transpontine fibers was found to project normally (Fig. 2B–2D). However, the deep cerebellar nuclei (dentate nuclei) appeared abnormally located more laterally (Fig. 3).

TMS was performed using a Magstim 200 stimulator (Magstim, Whitland, Wales, UK) equipped with a 65-mm figure-eight coil. Motor evoked potentials (MEPs) from the first dorsal interosseous (FDI) muscle of both hands were recorded simultaneously. Stimulation intensity was set at 120% of the resting motor threshold. Each site was stimulated four times; the shortest latency and the average peak-to-peak (PTP) amplitudes were then measured from the recorded MEPs.

TMS of each hemisphere in the patient robustly elicited contralateral MEPs in the respective opposite hand, whereas no ipsilateral motor response could be induced (Fig. 4A, 4B). Optimal stimulation sites for FDI muscles were identified about 5 cm lateral and 1 cm anterior to the vertex on both hemispheres. The excitatory threshold at the optimal stimulation was 77% in the left hemisphere and 87% in the right hemisphere. TMS of the right and left hemisphere elicited contralateral MEPs in the left hand (latency, 22.9±0.4 ms; PTP amplitude, 5.5±0.2 mV) and contralateral MEPs in the right hand (latency, 22.1±0.3 ms; PTP amplitude, 6.1±0.5 mV), respectively.

Fig. 3. Diffuse tensor imaging at the level of the dentate nuclei shows that the dentate nuclei is abnormally located more laterally (arrowheads).

Fig. 4. Recording of the right and left 1st dorsal interossei motor evoked potentials with transcranial magnetic stimulation of left (A) and right (B) cortex shows contralateral responses only, suggesting crossed corticospinal projections from each hemisphere. TMS, transcranial magnetic stimulation.
DISCUSSION

To the best of our knowledge, this study is the first to investigate the organization of the spinal motor system in JS using TMS. Our analysis demonstrated the presence of an exclusively decussated CST in one patient with JS, thus representing the first report of normal CST decussation in JS. Data on a possible presence of decussating fibers in corticospinal and superior cerebellar tracts in JS are scarce. Therefore, this condition is considered an exception to the Valsalva doctrine, which refers to the typical contralateral relationship of the brain to the body [6]. However, rudimentary decussation of the superior cerebellar peduncle fibers in JS has been reported in a few studies [2]. Among these, an fMRI study suggested the occurrence of a partial CST decussation in some JS cases [7]. Moreover, a recent neurophysiological investigation provided evidence of midline-crossing corticospinal projections being present in some patients with JS [4]. However, these individuals concurrently showed ipsilateral corticospinal projections, and similar projections were also observed in the aforementioned fMRI study.

Therefore, until now, no fully normal pyramidal decussation has ever been reported in JS. This patient showed exclusively midline-crossing corticospinal projections, which were considered to be the normal CST trajectories. This indicates that the motor and sensory systems did not show a mismatch between the target of a specific motor output and the source of the corresponding sensory information in each cerebral hemisphere. Therefore, the patient could be functionally advantaged in the performance of voluntary limb movements.

Whole exome sequencing of the patient showed that he carried compound-heterozygous mutations in the C5orf42 gene. There could also be a possibility of normal corticospinal organization according to the genetic abnormality detected. Further studies are needed to elucidate the potential role of genes associated with JS.

We are aware that this study presents several limitations. Although the diagnosis of JS was based on clinical features as well as MRI findings of the patient, the use of molar tooth sign as a pathognomonic marker of JS has been previously criticized [8]. Furthermore, due to the current resolution of DTI, an accurate differentiation between a complete absence versus decreased volume of fibers is still not possible with this technique. Because of the technical limitations, we were unable to perform fiber tractography to show proper pyramidal decussation in the caudal medulla and crossing defects in the superior cerebellar peduncles.

In conclusion, this case report indicates that a variety of corticospinal motor organization patterns might characterize JS, in relation to the severity of hindbrain malformations and the extent of embryonic developmental defects affecting pyramidal decussation processes.

CONFLICT OF INTEREST

No potential conflict of interest relevant to this article was reported.

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