Phelan-McDermid syndrome in two adult brothers: atypical bipolar disorder as its psychopathological phenotype?

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Abstract: The 22q13.3 deletion, or Phelan-McDermid syndrome, is characterized by global intellectual disability, generalized hypotonia, severely delayed or absent speech associated with features of autism spectrum disorder, and minor dysmorphisms. Its behavioral phenotype comprises sleep disturbances, communication deficits, and motor perseverations. Data on psychological dysfunctions are so far not available. Previous studies have suggested that the loss of one copy of the gene SHANK3 is related to the neurobehavioral phenotype. Additional genes proximal to SHANK3 are also likely to play a role in the phenotype of patients with larger deletions. The present paper describes two adult brothers with an identical 2.15 Mb 22qter (22q13.32q13.33) deletion, of whom the youngest was referred for evaluation of recurrent mood changes. In both patients, magnetic resonance imaging of the brain showed hypoplasia of the vermis cerebelli. Extensive clinical examinations led to a final diagnosis of atypical bipolar disorder, of which symptoms fully remitted during treatment with a mood stabilizer. In the older brother, a similar psychopathological picture appeared to be present, although less severe and with a later onset. It is concluded that the behavioral phenotype of the 22q13.3 deletion syndrome comprises absent or delayed speech and perseverations with associated autistic-like features, whereas its psychopathological phenotype comprises an atypical bipolar disorder. The latter may have implications for the treatment regime of the syndrome-related behavioral disturbances.

Keywords: 22q13.3 deletion syndrome, 22qter, Phelan-McDermid, SHANK3, autism spectrum, unstable mood disorder, cerebellar vermis

Introduction

More than a decade ago, a small number of patients were described in whom a telomeric deletion of the chromosomal region 22q13 was present and whose main phenotypical characteristics included marked developmental delay and absence of speech.1–3 Subsequently, the clinical picture of the 22q13 deletion syndrome was systematically elaborated by Phelan et al.4 According to their report, this deletion syndrome is characterized by global intellectual disability (ID), childhood hypotonia, severely delayed or absent speech, features of autism spectrum disorder (ASD) and normal to accelerated growth in the absence of major dysmorphisms and somatic anomalies. Since then, the 22q13 deletion syndrome has also been called Phelan-McDermid syndrome (OMIM: 606232), and more than 100 cases with deletions of variable length have been published.5–7 The exact prevalence is unknown and the syndrome is thought to be underdiagnosed. In a large cohort study among 11,000 patients with ID, it was the
second most frequent subtelomeric rearrangement, suggesting that it is a relatively common chromosomal disorder. The SH3 and multiple ankyrin repeat domains 3 (SHANK3) gene that is localized within the common deleted region were suggested to be responsible for at least the neurobehavioral features, including ID and features of ASD. This was supported by the presence of a balanced translocation disrupting SHANK3 in a patient with the 22q13 deletion syndrome phenotype, and further confirmed by patients with deletions restricted to SHANK3. The latter is a brain-specific gene and has a critical function for the postsynaptic densities in the brain. Several previous studies suggested involvement of this gene in ASD. Recently, Sarasua et al showed that several additional genes proximal to SHANK3 contribute to the phenotype of these patients as well.

Concerning the behavioral phenotype, several reports mention sleep disturbances, communication deficits, and behavioral abnormalities such as hyperactivity, restlessness, attention problems, repetitive motor activities, and increased reactivity to sensory stimuli. Features concerning language and social communication are typically subsumed under the heading of ASD. Apart from the developmental aspects mentioned here, data about psychological dysfunctions or psychiatric symptoms are not available as yet.

In an ongoing cohort study among patients with unexplained ID (Department of Human Genetics, Radboud University Nijmegen Medical Centre, Nijmegen, The Netherlands), genome-wide array analysis disclosed a 2.15 Mb identical 22qter deletion (22q13.32q13.33; Figure 1) in two adult male siblings of whom the youngest was already known (first author) with an unstable mood pattern. In the absence of any relevant information about psychological functioning in this deletion syndrome, both patients were extensively examined in order to identify a possible psychopathological phenotype.

**Cases**

Two male siblings are presented, aged 29 years (Patient 1) and 31 years (Patient 2), respectively, both born after uncomplicated pregnancies from nonconsanguineous parents. Apart from short-lasting episodes of flattened affect in the father, family history shows no load with any major psychiatric disorder, nor with ID or epilepsy.

**Patient 1**

The early childhood of this patient was characterized by sleep disturbances and a general developmental delay, particularly in respect of speech. He frequently displayed hyperactive behaviors with temper tantrums. At the age of 6 years he had a developmental delay of more than 3 years

![Figure 1](image-url) University of California, Santa Cruz genome browser view (GRCh37/hg19) of the deleted region in chromosome 22q13 identified by genome-wide array analysis. **Notes:** The red box indicates the chromosomal location schematically depicted. The relative positions of involved genes in the deleted region are shown. The green color indicates OMIM disease-causing genes, including SHANK3. Other involved genes are involved in diseases with autosomal recessive inheritance.
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Patient 2

Patient 2 is the older brother. His recent history mentioned an unstable pattern of mood and activity with predominant recurrent depressive episodes starting at the age of 27 years for which he was treated with carbamazepine (400 mg daily; 4.5 mg/L) and paroxetine (30 mg daily; 39 µg/L) elsewhere.

Four years later, after establishment of his genetic diagnosis, he was referred for extensive neuropsychiatric evaluation, which did not disclose any psychopathology. In respect of neuropsychology, a developmental age of 3 years 3 months (Vineland screener) was established, with a corresponding cognitive profile and communication problems due to moderately impaired development of speech and language. In particular, deficits in planning and maintaining overview were found. MRI of the brain showed a similar picture as in his younger brother (hypoplasia of the vermis cerebelli with enlarged cisterna magna, and mild enlargement of the lateral ventricles). Also here, in retrospect, a diagnosis of atypical bipolar disorder was made, albeit that relapsing symptomatology was less severe and with a later onset. Because psychiatric symptoms were in full remission, his psychopharmacological treatment regime was kept unchanged.

As shown in Table 1, primary behavioral and psychopathological features of both patients are from the mood and anxiety domains.
Table 1  Behavioral and psychopathological findings in both patients

<table>
<thead>
<tr>
<th></th>
<th>Patient 1</th>
<th>Patient 2</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Intelligence/Adaptation</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Male/29</td>
<td>Male/31</td>
<td></td>
</tr>
<tr>
<td>Vineland screener28</td>
<td>28 (2;4)</td>
<td>39 (3;3)</td>
</tr>
<tr>
<td>(developmental age in years; months)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Communication</td>
<td>13 (1;1)</td>
<td>36 (3;0)</td>
</tr>
<tr>
<td>Daily living skills</td>
<td>35 (2;11)</td>
<td>44 (3;8)</td>
</tr>
<tr>
<td>Socialization</td>
<td>28 (2;4)</td>
<td>37 (3;1)</td>
</tr>
<tr>
<td>Motor skills</td>
<td>37 (3;11)</td>
<td>37 (3;1)</td>
</tr>
<tr>
<td>Total score</td>
<td>28 (2;4)</td>
<td>39 (3;3)</td>
</tr>
<tr>
<td><strong>Behavior and psychopathology</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>PIMRA29 (proxy, maximum per scale = 7)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Schizophrenic disorder</td>
<td>2</td>
<td>1</td>
</tr>
<tr>
<td>Affective disorders</td>
<td>5</td>
<td>4</td>
</tr>
<tr>
<td>Psychosexual disorder</td>
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<td>0</td>
</tr>
<tr>
<td>Adjustment disorder</td>
<td>3</td>
<td>2</td>
</tr>
<tr>
<td>Anxiety disorders</td>
<td>4</td>
<td>2</td>
</tr>
<tr>
<td>Socioform disorders</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Personality disorders</td>
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<td>0</td>
</tr>
<tr>
<td>Inappropriate mental adjustment</td>
<td>3</td>
<td>1</td>
</tr>
<tr>
<td>PIMRA total score (max = 56)</td>
<td>18</td>
<td>10</td>
</tr>
<tr>
<td><strong>Neuropsychiatric rating scale</strong></td>
<td></td>
<td></td>
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<tr>
<td>(NPI30 severity [max = 4] × frequency [max = 3])</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Delusions</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Hallucinations</td>
<td>0</td>
<td>0</td>
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<tr>
<td>Agitation/Aggression</td>
<td>1</td>
<td>2</td>
</tr>
<tr>
<td>Depression/Dysphoria</td>
<td>9</td>
<td>1</td>
</tr>
<tr>
<td>Anxiety</td>
<td>4</td>
<td>0</td>
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<tr>
<td>Euphoria</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Apathy</td>
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<td>0</td>
</tr>
<tr>
<td>Disinhibition</td>
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<td>0</td>
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<tr>
<td>Irritability</td>
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<td>0</td>
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<tr>
<td>Aberrant motor behavior</td>
<td>3</td>
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</tr>
<tr>
<td>Night-time behavior disturbances</td>
<td>6</td>
<td>0</td>
</tr>
<tr>
<td>Appetite and eating abnormalities</td>
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<td>1</td>
</tr>
<tr>
<td>NPI severity × frequency</td>
<td>25</td>
<td>4</td>
</tr>
<tr>
<td>total score (max = 144)</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

**Abbreviations:** NPI, Neuropsychiatric Inventory; PIMRA, Psychopathological Instrument for Mentally Retarded Adults.

Genome-wide submicroscopic chromosomal analysis by 250 k single nucleotide polymorphism array analysis (Affymetrix, Inc, Santa Clara, CA) revealed in both brothers an identical terminal loss of 2.15 Mb in chromosomal region 22q13.32q13.33 (49.03–51.18 Mb according to University of California, Santa Cruz genome browser Hg 19). Chromosomal analysis in the parents gave normal results without any indications for a mosaic pattern (blood lymphocytes and buccal cells).

**Discussion**

In these brothers, with until recently unknown etiology of their severe ID and moderate to marked developmental delay of speech and language, genome-wide array analysis revealed a diagnosis of 22q13 deletion syndrome/Phelan-McDermid syndrome, which appeared to be associated with a long-lasting pattern of unstable mood and behavior. In the absence of a family load with (bipolar) affective disorder, in both brothers a psychiatric diagnosis of atypical bipolar disorder was made, albeit that the youngest patient presented with more pronounced symptoms and an onset at earlier age. Their respective psychiatric histories disclosed no symptoms belonging to the autism spectrum.

Although several authors have suggested that autistic symptoms belong to the behavioral phenotype,4,12,16–18,20,21 the incremental value of such a diagnostic vignette is only marginal, given the co-occurrence with severe ID and, as was also recently stressed by Sarasua et al7, relies heavily on the definition and the validity of the assessment process. In these patients, it is not autism that is at the heart of the phenotypical presentation but the characteristic combination of marked speech and language delay and executive dysfunctions, the latter mainly detectable through a pattern of perseverations and planning problems.

In retrospect, it can be argued that the fundamental lack of expressive language not only intensified the presentation of previously observed symptoms of major depression but also obscured the fluctuations in mood and behavior typical for so-called unstable mood disorder.22 This psychopathological syndrome that refers to an atypical bipolar disorder can frequently be observed in patients with ID irrespective of the etiology.23,24 The dysregulation of affect in these patients is most probably related to hypoplasia of the cerebellar vermis that is typically involved in the modulation of cognition and emotion.25,26

As their symptoms remitted under maintenance therapy with a mood stabilizer, compounds from this class may be the first choice in the treatment of fluctuating challenging behaviors in patients with ID and, more specifically, those with 22q13 deletion syndrome in whom an (atypical) bipolar disorder may be present. These patients also illustrate that categorical diagnoses of autism or ASD should be applied with great caution only. Such vignettes may seriously hinder the in-depth assessment of an individual patient’s functioning following the contextual neuropsychological paradigm.27 The latter is aimed to understand and to explain functional deficits and is relevant for the treatment design.

In conclusion, key elements of the behavioral phenotype of the 22q13.3 deletion syndrome, as previously described in the literature, are delayed or absent speech and perseverations with associated features from the autism spectrum. This is the first report of which the data suggest that the psychopathological phenotype of Phelan-McDermid syndrome comprises...
oscillating abnormalities in mood and behavior that can be attributed to an atypical bipolar disorder.

Acknowledgments
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Disclosure
The authors report no conflicts of interest in this work.

References

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