A Novel CIAS1 Mutation and Plasma/Cerebrospinal Fluid Cytokine Profile in a German Patient With Neonatal-Onset Multisystem Inflammatory Disease Responsive to Methotrexate Therapy

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ABSTRACT. The clinical features, the underlying CIAS1 mutation, and the results of cytokine analyses are described for a 10-year-old German boy with neonatal-onset multisystem inflammatory disease, whose condition improved with age. Disease onset occurred at 26 months of age with predominantly cutaneous (urticarial rash) and neurologic (headache, chronic meningitis) symptoms including early bilateral optic nerve atrophy, whereas articular manifestations were mild. Sequence analysis of exon 3 of the CIAS1 gene revealed heterozygosity for a novel missense mutation. A T515C transition led to the replacement of isoleucine by threonine at amino acid position 172 (I172T) in a region of cryopyrin flanking the PYRIN and NACHT domains. This mutation was not present in the parents or in 11 controls and therefore was considered to be de novo. Enzyme-linked immunosorbent assays were performed to determine interleukin-6 and soluble tumor necrosis factor receptor superfamily 1B levels in the patient’s serum and cerebrospinal fluid (CSF). Concentrations were highly elevated in the CSF, whereas corresponding serum levels remained low. The strong cytokine activation in the CSF corresponded with the neurologic symptoms. Local activation of intrathecal macrophages may therefore be an important pathogenetic mechanism. CSF cytokine levels decreased to normal under corticosteroid and intrathecal methotrexate therapy. When the boy reached the age of 5.5 years, treatment was stopped, and he has remained relapse-free. Pediatrics 2004;114:e124–e127.

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NOMID, CINCA, chronic infantile neurological cutaneous and articular syndrome, CIAS1 mutation, cytokines, aseptic meningitis, treatment.

ABBREVIATIONS. NOMID, neonatal-onset multisystem inflammatory disease; CINCA, chronic infantile neurological cutaneous and articular syndrome; CNS, central nervous system; MWS, Muckle-Wells syndrome; FCAS, familial cold autoinflammatory syndrome; IL, interleukin; MTX, methotrexate; CRP, C-reactive protein; ESR, erythrocyte sedimentation rate; Ig, immunoglobulin; CSF, cerebrospinal fluid; PCR, polymerase chain reaction; sTNFRSF1B, soluble tumor necrosis factor receptor superfamily 1B; ELISA, enzyme-linked immunosorbent assay; TNF, tumor necrosis factor.

NOMID/CINCA is a rare congenital autoinflammatory disorder (with ~100 reported cases worldwide) that results from missense mutations within exon 3 of the CIAS1 gene on chromosome 1q44. Alterations of this particular exon have been reported previously also to cause Muckle-Wells syndrome (MWS) and familial cold autoinflammatory syndrome (FCAS). Other autoinflammatory diseases associated with fever, arthralgia, and urticaria. Actually, >20 different CIAS1 mutations have been described to cause the 3 overlapping syndromes, with FCAS representing the mildest and NOMID/CINCA representing the most severe disease expression. Similar to FCAS and MWS, an autosomal dominant inheritance pattern is also assumed for NOMID/CINCA.

The CIAS1 gene encodes the protein cryopyrin (also called PYPAF1 or NALP3), which is highly expressed in polymorphonuclear cells and chondrocytes. It contains a PYRIN domain, involved in the regulation of inflammation and apoptosis. Because cryopyrin activates nuclear factor κB and also may regulate interleukin-1 (IL-1) processing through PYRIN domain-mediated caspase 1 signaling, CIAS1 mutations have been proposed to have a profound proinflammatory effect. Accordingly, during (and often between) febrile episodes, laboratory investigations reveal an inflammatory state with elevated acute-phase reactants.
Here we report a de novo CIAS1 mutation and the resulting cerebral cytokine activation in a patient of German origin, whose clinical picture resembled NOMID/CINCA and whose condition surprisingly improved with age after a long-term methotrexate (MTX) treatment over 2.5 years.

METHODS

The proband and his healthy, nonconsanguineous parents of German origin were investigated. Written informed consent was obtained from the patient’s parents. The study was conducted according to Helsinki Committee standards.

Patient

After an uneventful pregnancy, the 10-year-old boy was born at 40 weeks’ gestation with a birth weight of 3220 g. Since the age of 26 months, he exhibited a nonpruritic urticarial rash located mainly on his trunk. Three months later, the child showed recurrent fever, arthralgia (knees, ankles, feet), and a sudden, progressive visual impairment. At 32 months of age, he developed the clinical picture of acute meningitis, which was accompanied by high, spiking fever, exanthema, and arthralgia. Physical examination revealed a disproportionate increase of the cranial circumference and a striking delay in the closure of the fontanel. Ventricular enlargement was confirmed by cranial magnetic resonance imaging. Electroencephalography repeatedly showed diffuse slow waves. Neurologic examination demonstrated a normally developed boy without signs of psychomotor or psychomental retardation. Ophthalmologic evaluation revealed a bilaterally reduced vision, an optic atrophy, which was more pronounced on the left side, and pathologic visually evoked potentials. Laboratory findings showed neutrophilic leukocytosis (up to 85%; maximum: 32,500 per µL) with eosinophilia (9%), thrombocytosis (maximum: 575,000 per µL), C-reactive protein (CRP) levels up to 23 mg/dL, and an increased erythrocyte sedimentation rate (ESR) (maximum: 124/150 mm/hour, Westergren). Additional pathologic findings were subfebrile temperatures with exercise-induced fever, and slightly increased acute-phase reactant levels (CRP: 2.78 mg/dL; ESR: 57/93 mm/hour), but he had no neurologic symptoms, a normal electroencephalogram, and no signs of arthropathy or growth retardation. His visual impairment is slowly progressive, whereas his mild hearing deficit remains stable.

DNA Extraction

EDTA blood samples were collected from the patient and his parents, and the genomic DNA was isolated from white blood cells by the help of the QiAamp blood minikit (Qiagen, Hilden, Germany).

Polymerase Chain Reaction Amplification and DNA Sequence Analysis of the CIAS1 Gene

Exon 3 of the CIAS1 gene was amplified from high-molecular weight DNA with 2 oligonucleotides located in the adjacent intronic regions ~70 to 80 bases up- and downstream from the respective exon/intron border. The polymerase chain reaction (PCR) products were purified with the QIAquick PCR purification kit (Qiagen) and sequenced with the ABI PRISM Big Dye Terminator v3.1 ready reaction cycle sequencing kit (Applied Biosystems, Foster City, CA) by using 5 different internal primers. Sequences were analyzed on an ABI PRISM 377 DNA sequencer (Applied Biosystems) using the Sequence Analysis 3.4.5 program (Applied Biosystems).

Measurement of IL-6 and Soluble Tumor Necrosis Factor Receptor Superfamily 1B Levels

During disease exacerbation as well as under steroid and MTX treatment, cytokine measurements were performed in serum and CSF samples from the patient with commercially available enzyme-linked immunosorbent assays (ELISAs). The proinflammatory cytokine IL-6 was measured with a sandwich ELISA kit manufactured by DiaNova-Immunotech (Hamburg, Germany; sample volumes: 100 µL; normal value: <10 pg/mL). To determine serum and CSF concentrations of the soluble tumor necrosis factor receptor superfamily 1B (sTNFRSF1B, p75), an ELISA kit distributed by DPC-Biermann (Bad Nauheim, Germany) was applied. sTNFRSF1B levels increase during inflammation and in autoimmune diseases; however, even in healthy control persons, constitutive concentrations are present in the range of ≥1700 pg/mL. Absorbance was measured in an appropriate photometer (Dy- natech Laboratories, Inc, Chantilly, VA), and cytokine concentrations were calculated with an ELISA software program based on standard curves and positive controls.

RESULTS

We detected a novel T→C transition at cDNA nucleotide position 515 (with respect to the ATG start codon) located in exon 3 of the CIAS1 gene, which results in the exchange of isoleucine (ATC) by threonine (ACC) at amino acid position 172 (I172T). The boy is a heterozygous carrier of this missense mutation, which was not found in the patient’s healthy parents and on 22 control chromosomes and which seems to represent therefore a de novo mutation. It is also the first NOMID/CINCA-associated mutation that is located in the cryopyrin region that flanks the PYRIN and NACHT domain.

Because the boy mainly presented with signs of chronic meningitis including slowly progressive optic nerve atrophy and visual impairment, we also studied his cytokine profile in serum and CSF samples before and after therapy with prednisolone as well as before and during oral and intrathecal MTX treatment over a period of 10 months. Serum sTNFRSF1B levels measured before and after oral steroid therapy were always within the normal range (between 1600 and 2100 pg/mL). Similarly, serum...
IL-6 measurements revealed normal or only slightly increased concentrations during steroid as well as MTX therapy, ranging from 5 to 27 pg/mL. Thus, proinflammatory IL-6 and antiinflammatory sTNFRSF1B serum levels were not increased, although the patient showed elevated acute-phase reactants, with an ESR of up to 124/150 mm/hour and with maximum CRP levels of 23 mg/dL. Simultaneous measurement of the boy’s CSF cytokine concentrations during chronic meningitis, however, revealed markedly elevated levels of IL-6 (1190 pg/mL) and sTNFRSF1B (1130 pg/mL; Fig 1). After initiation of high-dose glucocorticoid therapy, cytokine levels as well as CSF cellularity decreased rapidly (IL-6: 120 pg/mL; sTNFRSF1B: 415 pg/mL; CSF: 7 cells per μL). After reduction of the steroid dosage, IL-6 concentrations increased again to values of ~800 pg/mL, which was accompanied by a CSF pleocytosis of 100 cells per μL. Onset of oral MTX treatment led to clinical improvement and normalization of CSF cellularity (3 cells per μL), whereas IL-6 levels initially remained elevated. During the following 2 MTX treatment months, a CSF pleocytosis (100 cells per μL) again was present, whereas IL-6 concentrations declined to ~400 pg/mL but did not return to normal. Because of the deterioration of his vision with the danger of imminent blindness, intrathecal MTX and prednisolone applications were started at the age of 41 months. Although CSF cellularity nearly normalized to 15 cells per μL, IL-6 levels increased to ~740 pg/mL, which was followed by a slow but continuous decline to 115 pg/mL over the next 3 months in the presence of mild pleocytosis (30–45 cells per μL).

**DISCUSSION**

NOMID/CINCA is the most severe of 3 overlapping syndromes also including FCAS and MWS, which were recognized recently to be the consequence of mutations in exon 3 of the CIAS1 gene.\(^7,8\) We have identified a novel I172T missense mutation also encoded by CIAS1 exon 3 in a German patient, who presented with clinical features compatible with NOMID/CINCA. This mutation is the first amino acid substitution associated with this disorder, which is located next to the PYRIN domain of cryopyrin. Only 1 additional mutation has been reported to be located in the region flanking the PYRIN and NACHT domain (V198M), leading to FCAS instead.\(^6\)

The CIAS1 gene is highly expressed in leukocytes (especially in monocytes and granulocytes) as well as in chondrocytes. Accordingly, our patient had predominantly neutrophils and monocytes in the peripheral blood as well as in the CSF. Cryopyrin was shown recently to be a potent upstream activator of nuclear factor κB signaling.\(^10\) Despite this, inflammatory mediators (especially cytokines) have been studied only sporadically in NOMID/CINCA\(^8,14\) and MWS\(^15\) by using serum or plasma samples but not CSF. Cytokine analyses in our proband revealed an intense activation of both proinflammatory IL-6 and antiinflammatory sTNFRSF1B in the CSF. Despite simultaneously increased acute-phase reactants, corresponding cytokine levels in serum remained normal. This result indicates a predominantly local inflammation of the CNS, which is consistent with the clinical phenotype of chronic meningitis in this patient. Because IL-6 plays a central role in acute-phase protein induction, one could speculate that the inflammatory process in the CNS induces a peripheral inflammation via a destroyed blood-brain barrier, with IL-6 not being measurable because of lower peripheral levels and a short half-life. Alternatively, the blood-brain barrier may be intact and peripheral IL-6 is therefore low.

In contrast to our findings, Aksentijevich et al\(^8\) detected increased serum levels of IL-6 and tumor necrosis factor (TNF) β1 in a patient with the CIAS1 D303N mutation, who presented with arthropathy as well as CNS inflammation, which was accompanied by an accumulation of cytokines in the peripheral blood. In accordance with the latter finding were the elevated IL-1β, IL-3, IL-5, and IL-6 concentrations measured in monocyte lysates of this patient. Hutenlocher et al,\(^14\) on the other hand, described a patient with severe arthropathy (but without CNS...
involvement) who had normal serum levels of IL-1, IL-6, and TNF-α, whereas TNF-β levels were elevated.

With regard to the clinical features, our proband suffered from a milder NOMID/CINCA variant, because he showed prominent CNS and eye involvement as well as urticarial exanthema but only mild arthralgia without swelling or development of any joint deformity or bony prominence. Furthermore, the age of the patient at onset of symptoms was 26 months, thus being outside the typical neonatal onset. Besides the lack of arthopathy and the delayed disease onset, this is also the first case of NOMID/CINCA with clinical improvement at later age and especially with unexpected remission of CNS symptoms. Despite 4.5 years of treatment cessation, there was no systemic relapse, and the patient’s visual defect showed only a slight progression. Whether this course is because of the earlier intrathecal MTX and prednisolone therapy or a direct consequence of the specific mutation, which is located next to the PYRIN domain, remains to be elucidated. Up to now, only 2 patients in whom treatment with MTX was efficacious in controlling the symptoms have been described. However, none of the patients received MTX intrathecally. Fortunately, the boy has also not shown any clinical signs of amyloidosis despite evidence of slightly but continuously elevated acute-phase proteins.

CONCLUSIONS

Our findings illustrate the broad spectrum of organ involvement in CIAS1-associated diseases (especially in NOMID/CINCA) and the phenotypic variability of the underlying mutations. Our results may also provide a basis for new therapeutic options such as intrathecal MTX application, sTNF-R-Fc fusion proteins such as etanercept (Enbrel), or monoclonal anti-IL-6 antibodies to control local and systemic inflammation.

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