Brief Communication

Extensive Hematoma in a Patient with Hereditary Hypersegmentation of Neutrophils

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Abstract

Hypercoagulable states are a group of conditions associated with an enhanced tendency toward blood clotting. Although usual clinical manifestations of hypercoagulable states are thrombotic events such as deep venous thrombosis, hematoma can also occur as a result of hypercoagulability in some patients. Several inherited or acquired conditions may lead to hypercoagulable states. Some of them include myeloproliferative syndromes, over activity of coagulation factors and methyl tetrahydrofolate reductase (MTHFR) polymorphisms. MTHFR is required for converting the amino acid homocysteine to methionine. Another significant role of an aptly functioning MTHFR enzyme is nucleic acid biosynthesis. Therefore MTHFR polymorphisms are expected to be associated with hypersegmentation of neutrophils because of a defect in DNA metabolism. Neutrophil hypersegmentation is one of the most sensitive hematological features of cobalamin or folate deficiency with normal serum vitamin B12-folic acid and iron levels. Hypersegmentation of neutrophils and hematoma or both of them suspected to be due to gene variations of MTHFR.

Here we report a 37-years old female who simultaneously affected by hereditary hypersegmentation and extensive hematoma. Laboratory analysis revealed normal serum vitamin B12, folic acid and iron levels. Routine and specific coagulation tests were normal in except of factor VIIIc that was high. Results of complete blood cell count (CBC) test were normal. Although this is just an idea, but simultaneous presentations of these two conditions can have a common origin.

Keywords: Hematoma, Neutrophil hypersegmentation, Hypercoagulability


Introduction

Hypercoagulable states can be defined as a group of conditions associated with an enhanced tendency toward blood clotting in the veins or arteries. Hypercoagulability, also known as thrombophilia, can result from several inherited and acquired conditions such as deficiencies in protein C, protein S, or antithrombin III, overactivity of
coagulation factors, antiphospholipid syndrome, inflammation and obesity (1-3). Polymorphisms of methylene tetrahydrofolate reductase (MTHFR) gene are also associated with an increased risk of heritable thrombophilia (4, 5). MTHFR gene produces the enzyme that plays an important role in amino acids processing, particularly the conversion of homocysteine to methionine. Two common polymorphisms in MTHFR gene including C677T and A1298C, lead to decreased enzyme activity and therefore elevation of homocysteine level. C677T exists in the MTHFR gene that encodes the catalytic domain of the MTHFR enzyme, in which a C>T substitution at position 677 results in a substitution of alanine to valine. The single amino acid substitution results in impaired folate binding and reduced activity of the MTHFR enzyme (6-8). Heterozygotes carrying this thermolabile variant have a reduced enzyme activity to 65% of normal, while homozygotes have only 30% of normal activity (9). It is found in homozygous state in 10% of the population and both forms of homozygotes and heterozygotes, cause elevated levels of plasma homocysteine (10). However usual clinical manifestations of hypercoagulable states are thrombotic events such as deep venous thrombosis (DVT) which are reported in associated with MTHFR polymorphisms (11), hematoma is also mentioned as a result of hypercoagulability in some patients. It is suggested that the presence of a hematoma may be the first indicator of an underlying thrombophilia (12). MTHFR plays a central role in folate metabolism via converting 5,10-methylenetetrahydrofolate to 5-methylenetetrahydrofolate that is the primary circulating form of folate and is vital for DNA synthesis. Polymorphism in MTHFR and other genes which play a role in folate metabolism may lead to deficient DNA synthesis and probably to neutrophil hypersegmentation (13). Coexistence of MTHFR polymorphism and hypersegmented neutrophils was reported in patients with hemolysis and cobalamin deficiency (14).

**Brief Report**

A 37-years old female was refer with extensive hematoma of both forearms. Initially routine and a number of specific coagulation tests including prothrombin time (PT) and activated partial thromboplastin time (aPTT), protein C and protein S as well as antithrombin III (ATIII) activity were performed. Factor VIII activity (FVIII:C) also was measured. All of these tests were performed by ACL top 300-autoanalyzer (Instrumentation Laboratory, Milan, Italy) (Table 1).

**Discussion**

NPC is caused by mutations in one of the two genes called NPC1 or NPC2. 95% of cases have NPC1 mutation which encodes a large glycoprotein in

![Fig. 1](image_url)

**Fig. 1.** Hypersegmented neutrophils in peripheral smear. (a) Patient blood smear. (b) Patient’s mother blood film Examination of peripheral blood films of patient and her mother reveals 4% neutrophils with more than five nuclear lobes and 12% neutrophils, having five nuclear lobes in patient and 6% neutrophil with five nuclear lobes in patient’s mother.
This gene mapped to chromosome 18q11-q12, spans 56 kbp and contains 25 exons. NPC2 encodes a small soluble lysosomal protein which binds cholesterol with high affinity (8). NPC2 mapped to chromosome 14q24.3, spans 13.5 Kbp and contains 5 exons. Deficiency in both types causes impairment in processing and utilization of endocytosed cholesterol (1, 8). The cellular hallmark of this disease is inability of cholesterol to transport from late endosomes to plasma membrane or reticulum endoplasmic therefore accumulation of cholesterol and products will occur (Figure 1) (1).

Therefore cholesterol storage is impeded and resulted in a different pattern of accumulation in neuronal and extra-neuronal tissues. These changes cause sphingomyelin metabolism alteration in extra neuronal tissues. U nesterified cholesterol, sphingomyelin, bisphosphate, glycolipids, and free sphingosine and sphinganine will accumulate in liver and spleen (1). In neurons, accumulation of Glycosphingolipids including GM2 and GM3 gangliosides occurs and cause meganeurite formation, growth of ectopic dendrites, neurofibrillar tangles

**Table 1:** Results of Routine and specific coagulation tests.

<table>
<thead>
<tr>
<th>Test</th>
<th>Result</th>
<th>Normal range</th>
</tr>
</thead>
<tbody>
<tr>
<td>PT</td>
<td>10.5</td>
<td>9.5-12.5 sec</td>
</tr>
<tr>
<td>aPTT</td>
<td>27.2</td>
<td>24.9-29.1 sec</td>
</tr>
<tr>
<td>FVIII:C</td>
<td>162</td>
<td>60-150 (%)</td>
</tr>
<tr>
<td>ATIII</td>
<td>100</td>
<td>80-120 (%)</td>
</tr>
<tr>
<td>PC</td>
<td>84%</td>
<td>70-140 (%)</td>
</tr>
<tr>
<td>PS</td>
<td>103</td>
<td>60-135.5 (%)</td>
</tr>
<tr>
<td>Fibrinogen</td>
<td>345</td>
<td>220-496</td>
</tr>
</tbody>
</table>

PT: Prothrombin time, aPTT: activated partial thromboplastin time, PS: Protein S, PC: Protein C, ATIII: Antithrombin III activity, FVIII:C: Factor VIII activity, Sec: seconds

Late endosomal location. This gene mapped to chromosome 18q11-q12, spans 56 kbp and contains 25 exons. NPC2 encodes a small soluble lysosomal protein which binds cholesterol with high affinity (8). NPC2 mapped to chromosome 14q24.3, spans 13.5 Kbp and contains 5 exons. Deficiency in both types causes impairment in processing and utilization of endocytosed cholesterol (1, 8). The cellular hallmark of this disease is inability of cholesterol to transport from late endosomes to plasma membrane or reticulum endoplasmic therefore accumulation of cholesterol and products will occur (Figure 1) (1).

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**Table 2:** Complete blood cell count (CBC) results of patients with extensive hematoma and heredity neutrophil hypersegmentation.

<table>
<thead>
<tr>
<th>Index</th>
<th>Result</th>
<th>Unite</th>
<th>Interpretation</th>
</tr>
</thead>
<tbody>
<tr>
<td>WBC</td>
<td>7000</td>
<td>/ul</td>
<td>Normal</td>
</tr>
<tr>
<td>RBC</td>
<td>376000</td>
<td>/ul</td>
<td>Low</td>
</tr>
<tr>
<td>Hemoglobin</td>
<td>11.8</td>
<td>g/dl</td>
<td>Low</td>
</tr>
<tr>
<td>Hematocrit</td>
<td>33.6</td>
<td>%</td>
<td>Normal</td>
</tr>
<tr>
<td>MCV</td>
<td>89.4</td>
<td>Fl</td>
<td>Normal</td>
</tr>
<tr>
<td>MCH</td>
<td>31.4</td>
<td>Pg</td>
<td>Normal</td>
</tr>
<tr>
<td>MCHC</td>
<td>35.1</td>
<td>g/dl</td>
<td>Normal</td>
</tr>
<tr>
<td>Platelet</td>
<td>192000</td>
<td>/ul</td>
<td>Normal</td>
</tr>
<tr>
<td>Neutrophil%</td>
<td>55%</td>
<td>%</td>
<td>Normal</td>
</tr>
<tr>
<td>Lymphocyte%</td>
<td>37%</td>
<td>%</td>
<td>Normal</td>
</tr>
<tr>
<td>Monocyte%</td>
<td>2%</td>
<td>%</td>
<td>Normal</td>
</tr>
<tr>
<td>Eosinophil%</td>
<td>3%</td>
<td>%</td>
<td>Normal</td>
</tr>
</tbody>
</table>
formation, neuroinflammation, and neuroaxonal dystrophy (8). On the other hand, some proteins like Rab9 or mannose-6-phosphate receptors transfer to cell membrane by late lysosomal system. Cholesterol accumulation could also impair this trafficking (1). Unexplainably neural death occurs mostly in Purkinje cells of the cerebellum. The function of these proteins are not defined yet so the exact mechanism and pathophysiology remains a mystery (1).

The only approved therapy for NPC so far is Miglustat (N-butyl-deoxyojirimycin). It is an iminosugar inhibitor of glucosylceramide synthase (1). This drug can stabilize the neurological manifestations including dysphagia (6). Miglustat by inhibiting glucosylceramide synthase reduce the synthesis of glucosylceramide-based glycosphingolipid in CNS (6). But long term clinical outcomes are still unclear (6). Other drugs used for symptom therapy including antiepileptic drugs (treatment of seizure), clomipramine, protriptyline, or modafinil (treatment of Cataplexy), anticholinergic agents (treatment of dystonia and tremor), Melotonin (treatment of insomnia). Physiotherapy for management of muscle spasticity and contracture is useful. As the major cause of mortality in these patients is aspiration pneumonia, the most important part of managing this disease is handling feeding abnormalities (1, 6). Current researches in the field of treating this disease is based on animal models mostly transgenic mice and cats and also testing various compounds like imatinib, curcumin, NSAIDs, neurosteroids (allopregnolone) and 2-HP-ß-cyclodextrin. The last compound showed significant improvement in disease natural history in animal models but needs further investigations (1).

The exact function of NPC genes are unknown therefore the pathophysiology of this disease is still a mystery (4). So the target metabolite in brain causing the neuroinflammatory responses remains unknown. These results are in lack of a biochemical blood test for evaluating prognosis or diagnosis (1, 4). Currently the gold standard for diagnosis is skin biopsy however recent advances suggest oxysterol profile as a biomarker for NPC. Although more researches are needed to define it as an indicator, these findings could change the future treatment and research developments (1, 4).

As mentioned above, NPC is a disease which involves multiple organs including central nervous system, respiratory system and gastrointestinal system (hepatosplenomegaly). Therefore this disease could interfere with routine anesthetic plans (3, 5). Restrictive lung disease arise from recurrent aspirations demand specific consideration in ventilator setup, so lower tidal volumes and increase in respiratory rate could be helpful for these patients. Another problem is the liver damage caused by storage of lipids, which needs careful selection of anesthetic drugs (5). Also hepatomegaly could accompany ascites which leads to decreased Functional Residual Capacity (3). there are some reports indicating the association between NPC and difficult airway therefore considering options for intubation would be wise (1). on the other hand, thrombocytopenia accompanied by this disease, could potentiate the risk of bleeding (1). Due to vigorous secretions anticholinergic drugs prior to anesthesia could improve the outcome of the procedure (3). It is demonstrated that hyperventilation plus high concentrations of Sevoflurane could induce seizures. Therefore to avoid this phenomenon continuing antiepileptic agents in perioperative period and use of TIVA instead of volatiles is recommended (3). The technique of choice is the one in which satisfactory condition for procedure is established rapidly and safely and also the recovery should be safe and predicted with minimal sequel for the patient (5).

Then a complete blood cell count (CBC) was performed using an automated counting device (Sysmex KX-21, SYMSMEX, Japan). All blood cell indices were normal in except of a mild anemia with decreased red blood cell (RBC) count (Table 2).

All blood cell indices were normal except red blood cell (RBC) count and hemoglobin concentration. Both are less than the normal range. Patient was treated with folic acid and iron supplement for more than 3 months before laboratory analyses. The patient's serum vitamin B12, folic acid and homocysteine concentrations were 24.8pg/ml, 12.02ng/ml and 13μmol/l, respectively. All of them were in normal range according to age and sex. Serum iron was also measured in order to determine possible cause of neutrophil hypersegmentation. The serum iron was 80μg/dL, it was normal.
Conclusion

Neutrophil hypersegmentation is an expected finding in folate and vitamin B12 deficiency (15, 16). There are several reports about present of neutrophil hypersegmentation in patients with iron deficiency (17, 18). This female had neutrophil hypersegmentation with normal serum iron and folate as well as vitamin B12.

Normal gene variations in MTHFR gene can alter folate metabolism and subsequently defective DNA synthesis (13). In the other hand this common gene variations have a strong relationship with hypercoagulability and several studies showed various signs of hypercoagulable states in different patients (19, 20). Although hematoma is a common presentation in patients with bleeding but in hypercoagulable states hematoma was also reported (12, 21). A hematoma is a collection of blood, usually clotted, outside of a blood veins that may occur because of an injury to the wall of a blood veins allowing blood to leak out into tissues where it does not belong. The damaged blood vessel may be an artery, vein, or capillary (22-24).

In this patient only FVIII: C is moderately increased and other signs of hypercoagulability are absent but gene variations of MTHFR also can be accompanied by normal level of homocysteine in vascular events (25). In the current case, presence of heredity hypersegmentation can be suggestive of a defect in DNA metabolism. This defect in DNA synthesis can also provokes coagulation activity and therefore hypercoagulability associated presentations (26-28). Neutrophil hypersegmentation may also be accompanied by other conditions such as thrombocytosis that was reported with normal folate and vitamin B12 serum levels (17). Neutrophil hypersegmentation with folic acid deficiency was reported in pregnancy but this situation is transient and is solved several weeks after delivery (29). This case was not pregnant and her neutrophil hypersegmentation was not transient. Since neutrophil hypersegmentation is an autosomal dominant disorder (30) we assessed patient’s mother for presence of hypersegmented neutrophils and we found 6% neutrophil with five nuclear lobes in her blood film (Figure 1). Neutrophil hypersegmentation is defined as any neutrophil having six or more nuclear lobes or more than 3% of neutrophils having five nuclear lobes, when 100 neutrophils are examined (ICSH recommendations for the standardization of nomenclature and grading). This case emphasized that neutrophil hypersegmentation can be accompanied by hypercoagulability. This situation can be a genetically disorder or acquired state, but according to this case, this situation seems to be due to a gene defect in folate metabolism but further studies are required.

Acknowledgment

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Conflicts of Interest

The authors declare that there are no conflicts of interest.

References