

# **Genetic and clinical characteristics of Wilson disease in patients from Hungary**

Ph.D. thesis

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## INTRODUCTION

Wilson disease (WD) is an autosomal recessive genetic disorder of copper metabolism caused by the mutation of the ATP7B gene, resulting in a defect of biliary copper excretion. Excessive copper in the liver, brain and in other organs leads to hepatologic and neuropsychiatric symptoms.

WD is a relatively rare disease with huge importance because of its serious symptoms and, without treatment, potentially fatal outcome.

The worldwide prevalence of the disease is estimated to be 30 per 1 million, with a gene frequency of 0.56% and a carrier frequency of 1 in 90. The number of WD patients in Hungary is estimated to approximately three hundred.

The WD gene (ATP7B), identified on chromosome 13q in 14-q21 positions, encodes a copper transporting P-type ATPase, which is predominantly expressed in the liver with reduced expression in the kidney, brain and placenta. This gene contains 22 exons spanning a DNA region of ~100 kb encoding a protein of 1465 amino acids.

Since the gene was identified in 1993, the number of known mutations within this region exceeds 350 variations, according to recent data on WD mutation. The most common one is the H1069Q point mutation (50-80%) in Hungary, similarly to other Central, Eastern or Northern European countries.

Genetic analysis has a huge clinical importance. Detection of mutations on both alleles makes the diagnosis and treatment possible before the appearance of the first clinical symptoms.

## OBJECTIVES

The aim of the present study was to analyze the demographic characteristics and the distribution of the patients and to assess the impact of genetic testing for diagnosis of WD in patients (n=142) from Hungary. In order to discover the rare, new, not yet described mutations, we performed the sequencing on most WD gene exons in collaboration with Prof. Peter Ferenci from the Vienna University.

In my research the following questions were studied:

### 1. Genetic mapping of Wilson disease in Hungary:

- What is the prevalence of H1069Q point mutation in patients with Wilson disease from Hungary?
- **Is there any difference between** data of WD patients from **Hungary** compared to those **from neighbouring or distant countries**?

### 2. Which mutations of ATP7B gene causing disease can be detected beyond the H1069Q in Hungarian Wilson disease patients?

### 3. Genotype-phenotype correlation analysis:

- Is there any **correlation between genotypes and several phenotypic manifestation** of the illness?
- What is the practical **clinical impact of genetic testing** for the diagnosis of WD?
- What are the **demographic characteristics of WD** patients in Hungary?

### 4. Assessment of the occurrence of acute liver failure in Hungarian Wilson patients:

- **What is the incidence of acute liver failure** among the Hungarian Wilson patients? How many **liver transplantation** was done **due to Wilson disease**? What was the prognosis of these patients?
- What were the genotypes of the affected patients? Is there any possible correlation between the mutations of *ATP7B* and the clinical manifestation?

## PATIENTS AND METHODS

We examined 142 patients from 128 families. As a hepatological centre, we are indebted to the colleagues for sending samples from throughout the country, which made it possible to reach this, even on an international scale, large number. There were 81 male and 61 female patients. Their mean age at the time of the onset of symptoms was 19 years, while at the time of diagnosis it was 22 years. So, on average, three years passed until the diagnosis; however, in some exceptional cases, regrettably, this period exceeds more than ten years. We analyzed the data of eight patients post mortem, who had died from fulminant liver failure.

Our diagnosis was established by the international scoring system based on the typical clinical symptoms and laboratory findings. Only patients with a score of  $\geq 4$  were included.

Seventy patients had neuropsychiatric symptoms, 66 had hepatic manifestation of the disease, and 6 patients had no clinical symptoms at all. Among the siblings of index patients, 47 were healthy and 14 had Wilson disease.

We also examined 60 healthy blood donors as control subjects to exclude that the new mutations reflect polymorphisms present in the Hungarian population.

Genomic DNA was isolated according to standard protocols. The most common mutation was assessed by seminested PCR-based RFLP assay. Heterozygotes and negative samples were then screened for mutations by denaturing HPLC and then sequenced.

## RESULTS

We found 35 mutations altogether, 10 of which have not been described and listed even in the international database yet. Forty-eight patients were homozygous for one of them (including forty patients homozygous for H1069Q), and 44 patients were compound heterozygous. In 6 asymptomatic siblings we proved WD by detection of two disease-causing mutations. In 123 patients we found at least one disease-causing mutation.

The most common point mutation was present in 70% of patients ( $n=100$ ), indicating that the detection of this mutation is very important in the diagnosis of WD in Hungary. Forty patients were homozygous (28%), 60 patients were heterozygous (42%), and 42 patients (30%) were negative for this mutation. This distribution within the WD population according to the H1069Q genotype is close to the equilibrium calculated from the Hardy-Weinberg estimation using the measured 49.3% allele frequency.

We analyzed the distribution of patients according to the age and manifestation of the disease at the detection of the first symptom. Patients with hepatic manifestation were younger than patients with neurological symptoms. The youngest patient was 3 years old and presented with liver disease; the oldest one was 49 years old presented with neurological symptoms.

After classifying patients according to clinical manifestation and presence of H1069Q mutation, we found that the Kayser-Fleischer ring was more frequent in H1069Q homozygous patients, and their mean age at the time of diagnosis was higher than in patients either heterozygous or negative for H1069Q.

In altogether 60 patients, mutations other than H1069Q were found, among which there were 26 missense, 1 nonsense, 1 splice substitution and 6 deletion (5/6 frameshift and the V1217-1218L with deletion of 6 bases). All of these mutations except Q1351X and K844K-fs appeared only in a few patients or in the index patients' family members.

By further analysis, we identified mutations on both alleles in 92 patients (65%) and one mutation in 31 patients (22%). In 19 patients (13%) both mutations are as of yet unknown, although they are clinically proved WD patients.

Most of the further mutations had already been observed in European patients—except R778G, which was described in Turkish patients, whereas the R778L is frequent in Far-Eastern (28-44%) but not in Central-European countries.

Since this is a hereditary disease, we performed the family screening tests in each index patients' case. As a consequence of recessive inheritance, WD in two consecutive generations is expected to be particularly rare. However, among Hungarian patients with Wilson disease, we found three such families. Therefore, this occurrence is not so rare among Hungarian Wilson patients.

We studied the occurrence of acute liver failure in Wilson disease.

Among the 142 WD patients tested genetically and the 8 young patients who had died in fulminant liver failure – most likely due to WD –, 28 had severe liver failure, among whom 19 patients (15 female and 4 male, mean age:  $16 \pm 4$ ) had acute or subacute liver failure. The juvenile deceased patients' data were analyzed post mortem, and in their several siblings Wilson disease was confirmed.

Fourteen orthotopic liver transplantations were done due to Wilson disease at the Department of Transplantation and Surgery of the Semmelweis University. More than 10 years have passed since the first transplantation. This patient is well, is asymptomatic, his ceruloplasmin level is in the normal range, and the Kayser-Fleischer ring has disappeared.

There was no difference in the survival rate calculated by the Kaplan-Meier method between the 23 patients who had received transplants due to metabolic liver disease (among them 14 WD patients) and those who had undergone liver transplantation because of other etiology (p=0,6).

#### Survival data of OLT patients from Hungary

Survival	Metabolic (n=23)	Non-metabolic (n=298)
1 yr	78%	77%
3 yrs	73%	71%
5 yrs	48%	66%

Twelve out of nineteen patients died in hepatic coma, several of whom had been on the waiting list for transplantation, and some of whom had not even been on the list, because the diagnosis had been established so late.

Fifteen out of nineteen patients with ALF had Coombs negative hemolysis, while in all of the Hungarian Wilson population it was detectable only in 25/142 patients.

Serum ceruloplasmin level was almost normal in fulminant form of WD, since it is an acute phase protein; AST was higher, ALT lower, serum bilirubin was much higher, and alkaline phosphatase was much lower than in the other subgroups. (AST/ALT= 2.25 vs 0.5 and 1).

The fulminant hepatic manifestation of Wilson disease is more frequent in women. It can be assumed that female sex hormones may aggravate hepatic copper accumulation.

#### CONCLUSIONS

Our previous observations were extended with the data of 142 Hungarian Wilson disease patients screened for most of the exons for mutations. The 35 different mutations found, eight of which had been described only in Turkish, Italian and Albanian patients before, reflect the genetic heterogeneity of WD in Hungary, which could be the genetic consequence of the 150-year Turkish occupation of Hungary in the 16th and 17th centuries.

The detection of H1069Q point mutation is very important in the diagnosis of WD in Hungary. By standard means (on clinical and chemical findings alone), 44 patients presented with liver disease (66/142) and 7 patients with neurological manifestation (70/142) would have failed to be diagnosed, while,

by adding genetic testing to the diagnostic algorithm, diagnosis could be established, which made further diagnostic tests unnecessary. On the other hand, by using molecular testing, the timeframe from symptoms to diagnosis was shortened in some cases.

Mutation analysis may be helpful in patients in whom typical clinical symptoms or findings of WD are absent: in patients presenting hepatic disease and in asymptomatic siblings. However, mutation analysis is not useable for first line screening, because of the huge number of mutations of the WD gene.

Examination of the index patients' siblings is very important. The presence of disease-causing mutations on both alleles is sufficient to diagnose Wilson disease, even in the absence of any actual clinical symptoms. Treatment started in preclinical stage could prevent the development of clinical symptoms.

#### OUR NEW RESULTS AND THEIR PRACTICAL SIGNIFICANCE

- In a nationwide collaboration, we extended our previous observations with the analysis of the data **from an internationally significant number of Hungarian Wilson disease patients (n=142)**. We assessed that, on average, a long time (several **years**) **had passed**, regrettably, **from the first symptom until the diagnosis**. The activity of our workgroup has also contributed to abridging this timeframe.
- We confirmed that **in Hungary**, similarly to other Central-Northern-Eastern European countries, **H1069Q was the most frequent** mutation, detected in 100 patients (**70%**), which means a **49% allele frequency**.
- **Thirty-four further mutations** were found by sequencing exon 6-20, including **ten new ones not described before** and not even listed in the international WD mutation database: **L517-fs (c1549-1559del11), N676I, S693Y, Y715H, M769L, W939C, V1001G, V1039F, P1273S and G1281D**. Eight of them had been described, apart from Hungary, only in Turkish, Italian and Albanian patients before, which could be the genetic consequence of the 150-year Turkish occupation of Hungary in the 16th and 17th centuries.
- Our results confirmed the general view of the impact of genetic testing in patients presented with hepatic disease or the asymptomatic siblings of an index case, as we documented in several case reports. Examination of index patients' siblings is especially important, since, in case of two disease-causing mutations, the beginning of the therapy

can be prophylactic in the development of clinical symptoms in preclinical stage, as we confirmed with several case reports.

- Wilson patients' distribution according to gender: In the whole Hungarian WD population, male patients were dominant, while in the subgroup of patients suffering from acute liver failure there were more women. Female sex hormones may aggravate hepatic copper accumulation.
- The distribution of patients according to the age and manifestation of the disease at the detection of the first symptom showed that patients with hepatic manifestation were younger than patients with neurological symptoms.
- The occurrence of acute hepatic failure was relatively frequent among the Hungarian Wilson patients and seemed to be associated with H1069Q heterozygous genotype.
- Wilson disease is a relatively rare cause of acute liver failure. However, it must be kept in mind in case of every liver disease with unclarified origin/etiology, considering that in almost 15% of the fulminant WD serum ceruloplasmin is in the normal range, without the presence of the Kayser-Fleischer ring and neurologic symptoms.
- We described three Hungarian families, where Wilson disease appeared in two consecutive generations.

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