Serendipity in Inborn Errors of Metabolism: Combining Two Genetic Mutations in a Single Patient

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Inborn Errors of Metabolism (IEM) are rare disorders, with the majority having an incidence of less than 10 per 100,000 births, and a total incidence of all metabolic diseases reaching 40 per 100,000 births. The incidences of Amino Acid disorders excluding Phenylketonuria and Glycogen Storage disease are 7.6 and 2.3 per 100,000 live births respectively1. The incidence and the frequency of inborn error of metabolisms vary worldwide and are based on racial and ethnic structure of population. Specific genetic mutations can be more frequently seen in specific ethnicities. For instance, sickle cell anemia, caused by a genetic mutation in the HBB gene, is more common in people of African, African-American, and Mediterranean origins. Ashkenazi Jewish are also well known to host the mutation responsible for Tay-Sachs disease2. In other words, diseases tend to “cluster” in families because genes are passed down from ancestors to offspring. Therefore, variations in the incidence and prevalence of genetic mutations are quite different from one geographic area to another and can be partially attributed to the evolutionary factors that influence genetic variation in general.

Alkaptonuria is a rare autosomal recessive disease with an incidence of 1:250,000 - 1,000,000 (Table 1). The homogentisate 1,2-dioxygenase (HGD) gene codes for the enzyme necessary to break down the homogentisic acid into maleylacetoacetic acid. This HGD gene is expressed in many organs mainly the liver, kidney, prostate, osteoarticular joints, and the gastrointestinal tract. Alkaptonuria is caused by the accumulation of homogentisic acid (HGA) in the body, with some secreted in the urine. HGA undergoes oxidation and then polymerization in connective tissues, cardiovascular, and ocular tissues, causing a bluish discoloration, an entity named as Ochronosis. Moreover, individuals with Alkaptonuria might have dark urine that turns black when exposed to air, a pathognomonic sign of this disease. Features of Alkaptonuria include increased risk of developing aortic valve and coronary artery diseases, and osteoarthritis-like damage in joints and spine. Moreover, Alkaptonuria is also classified as a secondary amyloidosis because of the deposition of serum amyloid A related to HGA ochronotic pigments; the latter was proven to be co-localized with Alkaptonuria type II Amyloids in the tissues3,4.

Pompe is a lysosomal storage disease type II. It is a rare autosomal recessive genetic disease caused by a mutation in the GAA gene responsible for the production of acid α-glucosidase (also known as acid maltase) enzyme. This enzyme breaks down glycogen into
PKU: Phenylketonuria; MDS: Mitochondrial DNA Depletion Syndrome


No statistical data are available regarding other inborn Acids Oxidation defects (26.9% and 0.0% respectively).

of metabolisms namely Organic Acids disorders and Fatty acids disorders (70.4%) is the highest among other inborn errors estimated to be 1:40,000 (Table 1)\(^5,6\).

Cardiomegaly, and failure to thrive. Pompe incidence is seen in adulthood period to a severe infantile-onset muscle disease. The disease appears. These are known as classic infantile-onset, non-classic infantile-onset, and late-onset. Severity can range from a mild slowly progressive muscle myopathy seen in adulthood period to a severe infantile-onset muscle hypertrophy, weakness, and hypertrophic myopathy. So, patients experience myopathy, hypotonia, hepatomegaly, cardiomegaly, and failure to thrive. Pompe incidence is estimated to be 1:40,000 (Table 1)\(^7\).

The prevalence of IEMs in Lebanon is considered one of the highest in the region and even worldwide. Out of 49,750 newborns screened, 44 cases were detected to have genetic mutations. The total incidence in Lebanon is 1:1130 compared to 1:1381 in Saudi Arabia, 1:9300 in Japan, 1:2105 in Italy, 1:3315 in Germany, 1:3367 - 4310 in the United States\(^7\). The prevalence of Amino Acids disorders (70.4%) is the highest among other inborn errors of metabolisms namely Organic Acids disorders and Fatty Acid Oxidation defects (26.9% and 0.0% respectively).

No statistical data are available regarding other inborn errors mainly Glycogen Storage disease\(^7\). Alkaptonuria and Pompe diseases are likely to occur in our geographic area (Middle East). Hence, having a single mutation in a patient is common as previously described hundreds of times in the literature. However, having more than one mutation in the same patient is an extremely rare occurrence. Human data presented in Table 1 were selected after an extensive search through PubMed and Medline. Different words were used to search for cases having combined genetic mutations in the literature, specifically in the inborn error of metabolism. “OR” and/or “AND” operators were used during the search. Only seven cases were found, including our case. Full data regarding case \#4 is not available. Three out of six cases (50%) have parents with consanguineous type of marriage. Based on the incidence of Pompe disease and Alkaptonuria as depicted in Table 1, the incidence to have both Pompe disease and Alkaptonuria at the same time is approximately 1:10,000,000,000 - 40,000,000,000. Our case is the only one reported in the literature up to our knowledge. This is a very unusual finding to have such combination of two recessive mutations.

One of the main causes of the high incidence of IEM in the Middle East is the high rates of consanguinity. In the Arab world, consanguinity includes first cousins, double first cousins, and second cousins. Consanguinity also includes individuals with at least one common ancestor as seen in tribes, endogenous marriages, and intra-communities. The main cause of high consanguinity in the Arab region is due to socio-cultural factors such as preservation of family structure and properties, ease of marriage, and better relations with in-laws. The overall consanguinity ranges from 12.8% to 80.4%. Consanguinity rate in the Lebanese population is considered low with respect to other Arab countries (12.8% - 42%). This rate can be as high as 80.4% as noted in Nubia in Egypt\(^8\). IEMs are relatively common in our region. This is mainly due to the high rates of consanguinity as stated above. The total consanguinity can reach up to 67%, reaching 49% with first cousin consanguinity\(^8\). Our case has no family history of parental consanguinity. Both his parents and his sister are clinically well. One of his cousins reported to have episodes of dark urine but was not reachable. Another one died at age of 14 years after having pneumonia and hepatosplenomegaly.

<table>
<thead>
<tr>
<th>Case #</th>
<th>Reference</th>
<th>Disease 1</th>
<th>Disease 2</th>
<th>Incidence for Disease 1</th>
<th>Incidence for Disease 2</th>
<th>Mutation type for Disease 1</th>
<th>Mutation type for Disease 2</th>
<th>Marriage Type</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>(^11)</td>
<td>PKU</td>
<td>Fabry Disease</td>
<td>1:15,000</td>
<td>1:80,000-117,000</td>
<td>Autosomal Recessive</td>
<td>X-Linked</td>
<td>Non-Consanguineous</td>
</tr>
<tr>
<td>2</td>
<td>(^12)</td>
<td>McArdle Disease</td>
<td>MDS</td>
<td>1:100,000</td>
<td>Unknown</td>
<td>Autosomal Recessive</td>
<td>Autosomal Recessive</td>
<td>Consanguineous</td>
</tr>
<tr>
<td>3</td>
<td>(^13-15)</td>
<td>Tyrosinemia Type 1</td>
<td>Angelman Syndrome</td>
<td>1:100,000</td>
<td>1:12,000-20,000</td>
<td>Autosomal Recessive</td>
<td>Autosomal Recessive</td>
<td>Paternal Uniparental Isodisomy</td>
</tr>
<tr>
<td>4</td>
<td>(^3, 4, 16-18)</td>
<td>Alkaptonuria</td>
<td>Huntington Disease</td>
<td>1:250,000-1,000,000</td>
<td>1:30,000</td>
<td>Autosomal Recessive</td>
<td>Autosomal Dominant</td>
<td>Data not available</td>
</tr>
<tr>
<td>5</td>
<td>(^19-21)</td>
<td>Glutaric Aciduria Type I</td>
<td>Isobutyrylglycinuria</td>
<td>1:300-100,000</td>
<td>Rare, with only 5 cases reported</td>
<td>Autosomal Recessive</td>
<td>Autosomal Recessive</td>
<td>Consanguineous</td>
</tr>
<tr>
<td>6</td>
<td>(^3, 4, 16)</td>
<td>Alkaptonuria</td>
<td>Sucrase-Isomaltase Deficiency</td>
<td>1:250,000-1,000,000</td>
<td>Unknown with 1:500-2,000 in Europe</td>
<td>Autosomal Recessive</td>
<td>Autosomal Recessive</td>
<td>Consanguineous</td>
</tr>
<tr>
<td>7</td>
<td>(^3, 4, 6, 22, 23)</td>
<td>Alkaptonuria</td>
<td>Pompe Disease</td>
<td>1:250,000-1,000,000</td>
<td>1:40,000</td>
<td>Autosomal Recessive</td>
<td>Autosomal Recessive</td>
<td>Non-Consanguineous</td>
</tr>
</tbody>
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PKU: Phenylketonuria; MDS: Mitochondrial DNA Depletion Syndrome

Table 1: Cases of double Inborn Error of Metabolisms, their incidence, mutation and marriage types.
Having parents not consanguineous makes the probability of finding a similar case even rarer.

Metabolic disorders can have early or late onset of signs and symptoms due to the deficiency or accumulation of biological products. The best clinical outcome of IEM patients is to have a prompt evaluation or screening and then referral for evaluation and management. Any delay in treatment can have drastic consequences and might cause death. Although neonatal screening programs have some false positivity and negativity mainly due to early screening, interference with drugs, etc.; however, it is a simple and reliable test available for screening newborns. Moreover, some diseases do not have a screening method during the neonatal period. Recently, the Pompe Disease Diagnostic Working Group reported the use of dried blood spot (DBS) as a minimally invasive, easy to collect and

<table>
<thead>
<tr>
<th>Case #</th>
<th>Reference</th>
<th>Age, Gender</th>
<th>Signs &amp; Symptoms</th>
<th>PMH</th>
<th>Laboratory Results</th>
<th>Additional Testing</th>
<th>Molecular</th>
<th>Treatment</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>11</td>
<td>3 years, Male</td>
<td>Slow growth, lack of appetite, severe abdominal pain, episodes of gastroenteritis</td>
<td>-</td>
<td>BH4: normal level</td>
<td>Plasma Galactosidase A: 0.32 nmol/h/ml, Leukocyte Galactosidase A: 0.65 nmol/h/mg.prot&lt;sup&gt;1&lt;/sup&gt;</td>
<td>GLA sequencing: hemizygous missense mutation g.5234G&gt;A in the exon 2</td>
<td>Diet therapy for PKU</td>
</tr>
<tr>
<td>2</td>
<td>12</td>
<td>At birth, Female</td>
<td>Hepatomegaly, jaundice, ascites, liver failure</td>
<td>Severe hypotonia, and thin muscles, at birth</td>
<td>Hypoglycemia, hyperlactacidemia (89 mg/dL, N &lt;20 mg/dL), increased creatine kinase</td>
<td>Southern Blot: 90% depletion of mtDNA, phosphorylase activity undetectable with MRCE activity normal in muscle, MRCE activity decreased in Liver</td>
<td>FAH gene: homozygous with IVS12+5G&gt;A</td>
<td>Died at 5 months of age</td>
</tr>
<tr>
<td>3</td>
<td>13-15</td>
<td>4 months, Male</td>
<td>Bronchiolitis, liver failure, decreased level of consciousness, psychomotor retardation, bouts of laughter, absence of speech, upper limb tremor</td>
<td>-</td>
<td>Tyrosine: 499 umol/L (N: 19-199), urine succinylacetone: 658 mOs/mol of creatinine (N: undetectable), plasma succinylacetone: 29 umOsm/L, αfetoprotein: 175,000 ng/mL</td>
<td>Karyotype and Fragile X studies normal, 15q12 methylation test: presence of unmethylated paternal &amp; loss of methylated maternal alleles</td>
<td>GAICDH sequencing: homozygous c.482G&gt;A in exon 7</td>
<td>Tyrosine and phenylalanine restricted diet, NTBC</td>
</tr>
<tr>
<td>4</td>
<td>NA</td>
<td>NA</td>
<td>Detected by neonatal screening at birth, pyelonephritis &amp; respiratory tract infection at 8 &amp; 11 months, enlarged subarachnoid space, cerebral ventricles, &amp; retarded myelination</td>
<td>-</td>
<td>-</td>
<td>Dried blood newborn screening: glutarylcarnitine (acylcarnitine C5DC):1.03 umol/L (N: 0.0-0.017), α-acetylcarnitine: 1.23 umol/L (N: 0-1.11)</td>
<td>GCDH sequencing: homozygous missense mutation c.482G&gt;A in exon 5, ACAD8 sequencing: splice-site mutation c.481+3G&gt;C in exon 6</td>
<td>Dietary treatment, supplementation of L-carnitine, early interventions for intercurrent illnesses</td>
</tr>
<tr>
<td>5</td>
<td>19-21</td>
<td>3 days, Female</td>
<td>Progressive motor weakness &amp; failure to thrive, hepatomegaly, black spot in the right eye</td>
<td>Recurrent respiratory infection, greenish mucoid stool, dark urine, GE reflux</td>
<td>CPK: 1679 IU/L, α-Glucosidase: 4.0 nm/mg&lt;sup&gt;*&lt;/sup&gt;, Urine homogentisic acid: 15027 mmol/mol Creatine</td>
<td>Urine 1,4 α-Glucosidase deficiency by GC-MS</td>
<td>GAA sequencing: 2 heterozygous (C.670&gt;T in exon 3, C.1064&gt;T&gt;C in exon 6), HGD gene: homozygous deletion of 649 bp in exon 2 &amp; surrounding DNA sequences in flanking introns</td>
<td>Orfadin (Nitisinone)</td>
</tr>
</tbody>
</table>

NA: not available; mtDNA: Mitochondrial DNA; MRCE activity: Mitochondrial respiratory chain enzyme activity; NTBC: 2-(2-(2-nitro-4-trifluoromethylbenzoyl)-1,3-cyclohexanedione); BH4: Tetrahydrobiopterin; PMH: Past Medical History; CPK: Creatinine Phosphokinase; GE reflux: Gastroesophageal reflux; GC-MS: Gas Chromatographic-Mass Spectrometric. <sup>1</sup> Leukocyte Galactosidase A normal range: 26-80 nmol/h/mg.prot; N: Normal White Blood Cells α-Glucosidase control level: 127 nm/mg.

**Table 2:** Cases of double Inborn Errors of Metabolism, their demographic, clinical, and laboratory data.
transport sample for measuring α-glucosidase activity. Moreover, Jacomelli et al validated a rapid color intensity method for semi-quantification of homogentisic acid (HGA) using a dried urine spot (DUS) paper by comparing HGA concentrations in urine and plasma to those obtained from High Performance Liquid Chromatography (HPLC). The brown color intensity obtained on the paper is proportional to HGA concentration from 1.4 mmol/L to 46 mmol/L. Such method fits very well for Alkaptonuria screening even before the detection of dark urine because of its safety, reliability, and ease of use. After screening, additional confirmatory testing is necessary that include measuring enzymes activities and/or molecular analysis for gene mutations.

In our case, the patient was referred to the American University of Beirut Medical Center because of progressive motor weakness and failure to thrive at age of 6 years. His past medical history includes recurrent respiratory tract infections, greenish mucoid stools, frequent episodes of dark urine, and gastroesophageal reflux. Initial laboratory testing include a comprehensive panel for hematologic and biochemical tests in addition to urine organic acid measurements. All tests were within normal range except for creatinine phosphokinase (CPK) enzyme, plasma α-glucosidase, homogentisic acid, and urine 1,4 α-glucosidase. Demographic, clinical, and laboratory data are summarized in Table 2. Further molecular tests for GAA and HGD genes mutations were performed. GAA sequencing revealed a 2 heterozygous mutations, the C.670>T in exon 3, and the C.1064T>C in exon 6. HGD sequencing also revealed a homozygous deletion of 649 base pairs in exon 2 and the surrounding DNA sequences in flanking introns. The patient received orfadin ( nitrogen) that improved his Alkaptonuria condition by decreasing HGA. The patient was lost to follow up since 2013 where he was waiting to receive enzyme replacement therapy (ERT).

As a conclusion, occurrence of a double genetic mutations in one person is extremely rare, specifically in our case where neither the disease prevalence, nor consanguinity are in favor to have such a rare outcome. However, it should be kept in mind there the possibility of having more than one mutation in a patient is still there, especially when the clinical presentation of the patient is complex and signs and symptoms do not fit in one disease. Presence of parental consanguinity can be an additional key point to suspect such cases. Moreover, it is of utmost importance to screen neonates and pediatric patients if clinical signs and symptoms indicate the presence of IEM as soon as possible using the dried blood neonatal cards in order to prevent the outcome of such debilitating and fatal diseases.

Conflict of Interest statement

The authors have no competing conflicts of interest in the field to disclose. This manuscript is neither submitted for publication, nor published elsewhere in the same form, in English or in any other language, without written consent of the copyright holder. The authors have contributed significantly to this paper and understand it and they all agreed to this submission. In addition, the authors have no relationship, financial or otherwise, with any manufacturers or distributors of products evaluated in this paper.

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