

ANALYSIS OF PHENYLKETONURIA (PKU) BY USING INTERPROSCAN

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Abstract:

Phenylalanine hydroxylase (PAH) is the enzyme that converts phenylalanine to tyrosine as a rate-limiting step in phenylalanine catabolism and protein and neurotransmitter biosynthesis. Deficient enzyme activity leads to the disorders like hyperphenylalaninaemia and phenylketonuria. The determination of the crystal structure of PAH now allows the determination of the structural basis of mutations resulting in PAH deficiency.

We present an analysis of the structural basis of 120 mutations with a 'classified' biochemical phenotype and/or available *in vitro* expression data. We have find that the mutations can be grouped into five structural categories, based on the distinct expected structural and functional effects of the mutations in each category.

Structural information helps to formulate some rules that will help to predict the likely effects of unclassified and newly discovered mutations: proteins with truncations and large deletions, fusion proteins and active site mutations generally cause severe phenotypes; domain structure mutations and dimer interface mutations spread over a range of phenotypes.

Keywords: phenylalanine hydroxylase, biochemical phenotype, mutations hyperphenylalaninaemia, phenylketonuria.

INTRODUCTION:

Phenylketonuria (PKU) is an autosomal recessive genetic disorder characterized by a deficiency in the hepatic enzyme phenylalanine hydroxylase (PAH).^[1] This enzyme is necessary to metabolize the amino acid phenylalanine ('Phe') to the amino acid tyrosine. When PAH is deficient, phenylalanine accumulates and is converted into phenylpyruvate (also known as phenylketone), which is detected in the urine.^[2]

Phenylketonuria was discovered by the Norwegian physician Ivar Asbjørn Folling in 1934^[3] when he noticed that hyperphenylalaninemia (HPA) was associated with mental retardation. In Norway, this disorder is known as

Falling's disease, named after its discoverer.^[4]

PKU is normally detected using the HPLC test, but some clinics still use the Guthrie test, part of national biochemical screening programs. Most babies in developed countries are screened for PKU soon after birth.^[5]

Structures

Phenylalanine hydroxylase is a tetramer composed of four monomers, that is, composed of 4 identical subunits. Each subunit is in turn composed of three domains, a regulatory domain, a catalytic domain, and a tetramerization domain.

The *regulatory domain* is composed of the approximately 115 amino acids

nearest the amino terminal of the subunit.

The *catalytic domain* is composed of the next approximately 300 amino acids, and is responsible for all of the catalytic activity of the enzyme.

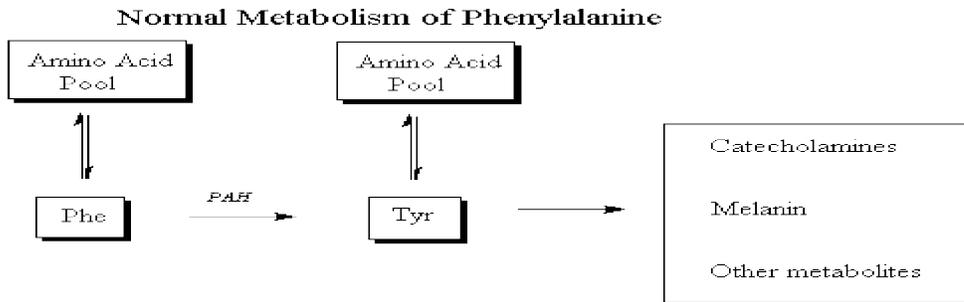
The *tetramerization domain* consists of the remaining amino acids and through the formation of a coiled-coil arrangement of amino acids, holds the tetrameric structure of the holoenzyme together with a leucine zipper. Phenylalanine hydroxylase contains one bound iron atom per subunit which is necessary for catalytic activity.

Normal catabolism of phenylalanine

In humans, phenylalanine is normally oxidized by the enzyme phenylalanine hydroxylase (*PAH*) to form the amino acid tyrosine; this is in fact the normal biosynthetic route to tyrosine for humans. From tyrosine, there are further connections to the biosynthesis of catecholamines, melanin, hormones, etc. Usually, dietary intake of Phe and Tyr, and the body's demand for Phe and Tyr, are fairly closely balanced. However, when there is too much phenylalanine in the body's pool of amino acids, it must be eliminated, either by excretion or by biochemical reaction.

Fig 1: Normal Metabolism of phenylalanine

There are two routes by which the (Tyr), and transamination to



excess Phe can be metabolized: oxidation to tyrosine (the normal and main route for degradation of Phe, and the normal route for biosynthesis of

phenylpyruvate and subsequent further metabolism (a minor route, which comes to the fore when the main route is blocked).

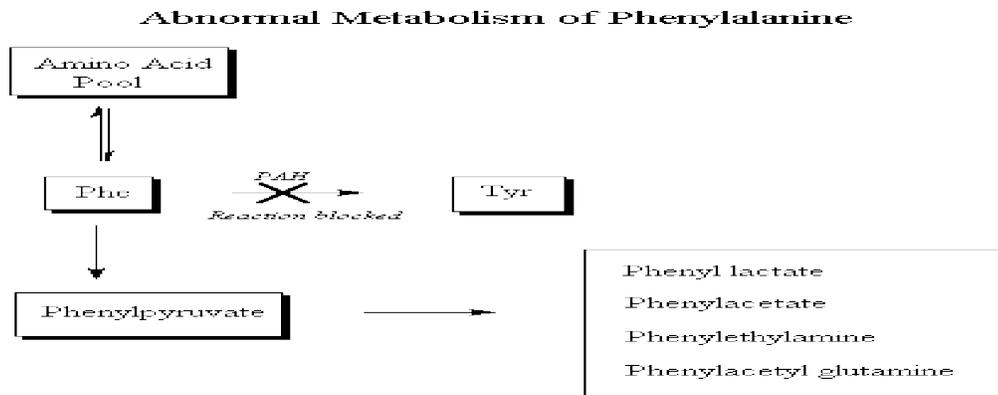


Fig 2: Abnormal Metabolism of Phenylalanine

Consequences of a Block in the

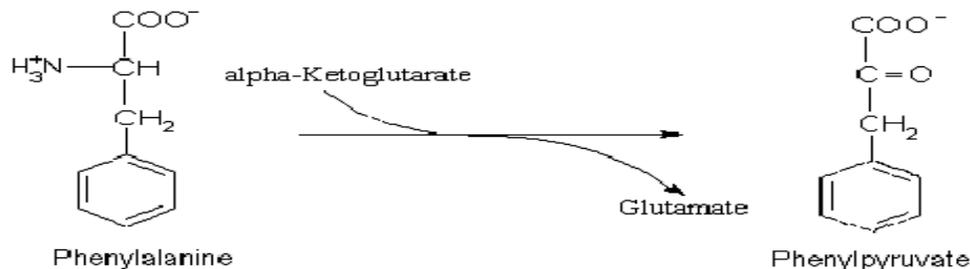
Oxidation of Phenylalanine

When the main route of degradation of Phe is blocked, then serum concentrations of Phe and its metabolites will rise. Three forms of

Table 1:

Compound Concentration	Normal	<i>Benign</i> HPA	<i>Variant</i> HPA	<i>Classic</i> HPA / <i>Classic</i> PKU
Phenylalanine	approx. 1 mg/dL (0.061 mM)	4-10 mg/dL (0.242-0.605 mM)	10-20 mg/dL (0.605-1.21 mM)	above 20 mg/dL (above 1.21 mM)

hyperphenylalaninemia (HPA) can be distinguished, based on serum concentrations of phenylalanine: *benign*, *variant*, and *classic*.



Under conditions of hyperphenylalaninemia a minor route of Phe metabolism becomes important. The minor route for degradation of Phe starts with the transamination of Phe to phenylpyruvate:

Fig3: Transamination of Phenylalanine to Phenylpyruvate

This transamination reaction is a standard one in the metabolism of amino acids. Note the role of α -ketoglutarate as the acceptor of the amino group from Phe, with consequent formation of glutamate.

The phenylpyruvate is further metabolized. Decarboxylation of phenylpyruvate gives phenylacetate, while a reduction reaction gives phenyl lactate. The phenylacetate can be further conjugated with glutamine to give phenylacetyl glutamine. All of these

metabolites can be detected in serum and urine by suitable clinical tests.

Materials and Methods:

1) NCBI-ENTREZ

CROSSDATABASE SEARCH

When it comes to getting information on sequences and databases, many paths lead to the National Center for Biotechnology Information (NCBI). The ENTREZ databases and querying

system developed at NCBI offers a powerful and relatively simple way to search most of NCBI's numerous databases. The main drawback with ENTREZ was that only one database could be accessed at a time.

mRna Sequence of PAH [NM_000277]

Fig: 4 Results: Interproscan

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InterProScan Results

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SEQUENCE: NP_000268.1 CRC64: 018F00EBBDDCE2F LENGTH: 452 aa

InterPro IPR001273 Family	Aromatic amino acid hydroxylase PR00372 Q3D5A1-10-800-10 PTHR11473 PF00351 SSF56534	FYWHYDRXLASE Aaa_hydroxylase Aaa_hydroxylase Bioterin_H Aaa_hydroxylase
InterPro IPR002912 Domain	Amino acid-binding ACT PF01842	ACT
InterPro IPR005961 Family	Phenylalanine-4-hydroxylase, tetrameric form TIGR01288	Phe4hydrox_tetr
noIPF unintegrated	unintegrated PD002559 PIRSF000336 SSF55021	Aaa_hydroxylase TH SSF55021

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ACKNOWLEDGEMENT:-

We would like to express deep sense of gratitude to Mr. Sanjaykumar Choubey for his guidance, persuasion, sincere work, and we also thankful to Mr. Pramod Deshmukh for his valuable, guidance, suggestion and enthusiasm, which helped us a lot.

CONCLUSION:

A Deficiency in PAH activity cause PKU. This PKU can be detected through Patients urine and serum.: A person suffering from this disease they have to face many problems like mental retardation and seizures.

However, PKU is one of the few genetic diseases that can be controlled by diet. A diet low in phenylalanine and high in tyrosine can be a very effective treatment. There is no cure. Damage done is irreversible so early detection is crucial.

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