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Research article

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### Prediction of target enzyme for diabetes mellitus by isolated blood samples

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#### ABSTRACT

Diabetes Mellitus is a serious problem universally, by and large in both developing as well as underdeveloped nations. Conversely, a synthetic drug management of diabetes mellitus creates an unexpected adverse effect towards all organs and it is not safeguarding too. However, numerous enzymes, which is a real concern and may be accountable for diabetes mellitus and its associated complications, which would certainly afford a plenty of scope and platform to explore out the main target enzyme. These enzymes are to be targeted in treating diabetes mellitus may have a proven advantage in reducing the diabetes mellitus consequences. There are several evidences indicated Protein Tyrosine Phosphatase and variety of isomers are responsible diabetes mellitus, however Protein Tyrosine Phosphatase and its isomers are hardly targeted, due to inappropriate clinically validation lapse. Researchers are clinically validating the indeterminate Protein Tyrosine Phosphatase to establish an efficient method for variability of Protein Tyrosine Phosphatase enzyme levels of various clinical conditions of diabetes mellitus and its associated diseases.

**Keywords:** Diabetes Mellitus, Protein Tyrosine Phosphatase Synthetic drugs, clinically validation and Researcher.

#### INTRODUCTION

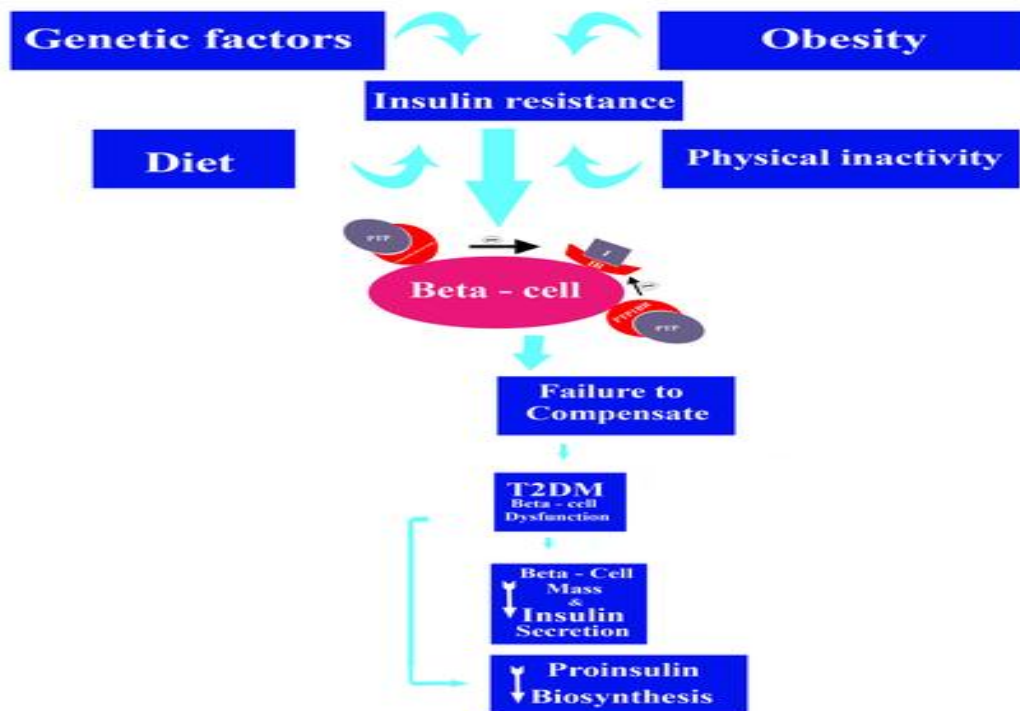
Diabetes mellitus has aggravated prevalence in the human population due to autoimmune obliteration of the  $\beta$ -cells of the pancreas through resulting insulin paucity [1-7]. The World Health Organization (WHO) predictable throughout status to diabetes mellitus and exist 7th foremost motive of casualty in 2030 and also addition it force double flanked by 2005 plus 2030 [8-11]. Present is a immense need for unambiguous enzyme target of precise metabolic complications for diabetes mellitus. Protein Tyrosine Phosphatase receptor is single central opening for triggering the insulin

receptors to liberate the vital capacity of insulin. Protein Tyrosine Phosphatase receptors and its subtypes (PTP1A, PTP1B, PTP1C, PTPN1, PTPN20CP, PTPN3, PTPN4, PTPN5, PTPN6, PTPN7,PTPN9, PTPN11, PTPN12, PTPN13, PTPN14, PTPN18 PTPN20, PTPN21, PTPN22 &PTPN23) offered, which is responsible for diabetes mellitus but its clinical value is mysterious [12-19]. The curative plants and synthetic anti diabetic medicines approaches towards healing of diabetes mellitus but doesn't know the exact intention enzyme for type II diabetes mellitus [12, 42 & 21-23]. Based on the above reasons,

systematic justification is obligatory Protein Tyrosine Phosphatase receptors and its sub types.

### The hypothesis

The hypotheses are proposed to find the target enzyme and its subtypes, by may be collection of blood sample from diabetic patients.



**Fig:1** (Abbreviations: IR-Insulin Receptor, I-Insulin, PTP- Protein Tyrosine Phosphatase) Has indicated Protein Tyrosine Phosphatase 1B (PTP1B) is a molecule that negatively regulates Insulin Receptor. In addition to insulin sensitization, inhibition of PTP1B has potential to promote weight loss, which is a benefit since obesity largely contributes to the type 2 DM pathology.

### Evaluation of the hypothesis

Although Protein Tyrosine Phosphatase and subtypes (Fig-1) for their working mechanism against diabetes mellitus and associated diseases have not been well understood, the two main afferent input to trigger insulin receptor to regulate the insulin release from beta cell and activation of glucose transporters. Other category of causative factors like PPR Gama, DDP-4, glycolysis and gluconeogenesis are clinically validated but more number of drugs promoted, which can't completely cure the diabetes mellitus [24-26]. Research suggestion shall be ready and set onward some Hospital have Institutional Ethical Committee. Consequent to the Institutional Ethical Committee ahead approval for research work shall be initiated. Characteristic guiding principle based, feedback form

might be primed and a consent from diabetologist in the direction of the collection of tolerant data and Pharmaceutical care issues. The variety which control demographic data like age, sex, social history, family history, current treatment regimen, change of prescription drugs and current position of blood glucose level. In my limitation has expressed background for applying different inputs of subsequent two criteria must be measured such as 1. Inclusion criteria: a. Patient age, b. Patient with diabetes mellitus and with extra Co-morbidities, c. Patient diabetes mellitus as In-patient. d. Patient effortlessly to read and write the consent form and 2. Exclusion criteria: a. Patient age, b. Patient visited diabetology department without diabetes mellitus, c. Patient attenders and bystanders and d. Patient who

unable to read and write the consent form [27-45]. Based on the criteria mentioned above, specified number of patients selected for scrutiny, hence segregated different groups such as Type-I, Type-II and associated diseases with Type-I & II for collect the blood samples, which subjected to Protein Tyrosine Phosphatase analysis.

### Protein Tyrosine Phosphatase Estimation

The T-Cell Fluorimetric Assay Kit will be purchased abnova, sigma Aldrich, which will used to estimate the amount of Protein Tyrosine Phosphatase levels in DM and associated diseases conditions and also available two methods such as one step process and two-step process. First 10+10 sample were performed one step + two step methods for the purpose to select the method of PTP analysis, which was economically and also get accurate result.

### RESULTS AND DISCUSSION

Prophylaxis strategies for management of diabetes mellitus are mainly based on patents risk. The Protein Tyrosine Phosphatase and its subtypes most widely used as risk assessment tool for risk stratification in diabetes mellitus [28-29]. However sensitivity and specificity of (Table –I), Shown Protein Tyrosine Phosphatase for an hodgepodge of

clinical conditions of Type-II diabetes mellitus placid augmented in the human body when compared to apiece supplementary allied diseases and additional parameters resembling different types of receptors, which have type of chromosomes and mechanism of action included [52-53]. Above table will help to evaluate the performance of a customized Protein Tyrosine Phosphatase 1/2 and other isomers panel for Diabetes Mellitus -risk assessment with patients [54-55]. The review scrutiny of diabetes mellitus and coupled diseases epidemiological and intention ruling studies in Tamil Nadu, will be established to the occurrence is rising exponentially in our kingdom. Our studies will be confirmed increasing diabetes mellitus and associated level were driving this epidemic. There is a vital necessity to widen appropriate strategies for avoidance of diabetes mellitus and associated disease in India by population based approaches, which may valuable to discover the interaction sandwiched between PTP isomers and its antagonist drugs. This hypothesis will be useful for diabetes mellitus research workers to discover new enzymes, and its subtype for the management of diabetes mellitus by the help of some micro oraganisms [56-60] and to further diminish the occurrence of diabetes mellitus associated risks.

**Table- I, Multiple-enzyme sequencing sheet indicated Protein Tyrosine Phosphatase disturbed various detached situation**

Sl. No	Clinical Condition	Receptors	Chromosomes	Mechanism	PTP Level in Blood	References
1	Normal	PTP and all isomers	12q15-q20	Trigger Insulin Receptor	20-30 µg/100ml	[29]
2	Type-1DM	PTP1A,B,C,N22	20p13,12q15-q21,1q-31-q32 & 2q-35-q36.1	β <sub>2</sub> integrin adhesion site action	Mild increased	[54&30]
3	Type-2 DM	PTP1B,O,N22	12q15-21,2p12p13-p12 & 2q-35-q36.1	RPTPα localize adhesion	Moderately increased	[54&31]
5	Type-2DM Associated Hypertension	PTP1A,D,C	12q15-q20,9p24.1-p23&1q31-q32	Focal adhesion formation	More increased	[54&33]
6	Type-2DM Associated Asthma	PTP1A,B,V	20p13,12q15-q21 & 12q15-q21	Inhibit integrin	Marked level increased	[54&35]
7	Type-2 DM Associated Cardiovascular	PTP1A,B,C,N22,U	20p13,12q15-q21,1q-31-q32, 2q-35-q36.1 & 1p35.3	Activation of mutation	Marked level increased	[54&36]

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