# Molecular characterization of beta-ketothiolase deficiency in 10 Indians: Discovery of 4 novel mutations in *ACAT1* gene

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- 4) MILS International, India.





#### **Disclosure Information**

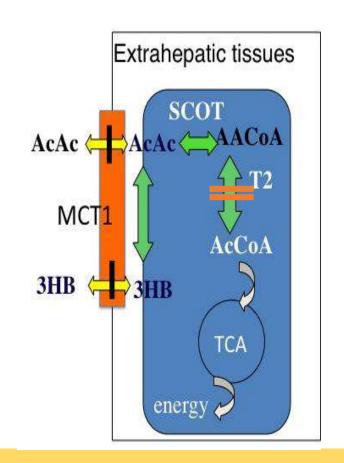
Elsayed Abdelkreem

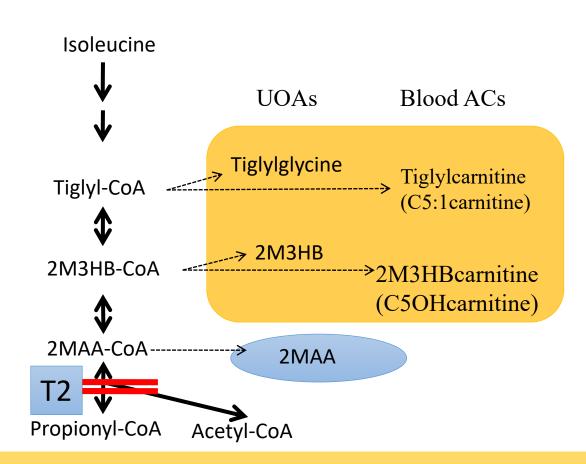
I have no financial relationships to disclose

#### **Beta-ketothiolase deficiency**

- Also known as mitochondrial acetoacetyl-CoA thiolase or T2 deficiency.
- Affects ketone body utilization and isoleucine catabolism.

## Mitochondrial acetoacetyl-CoA thiolase (T2) deficiency

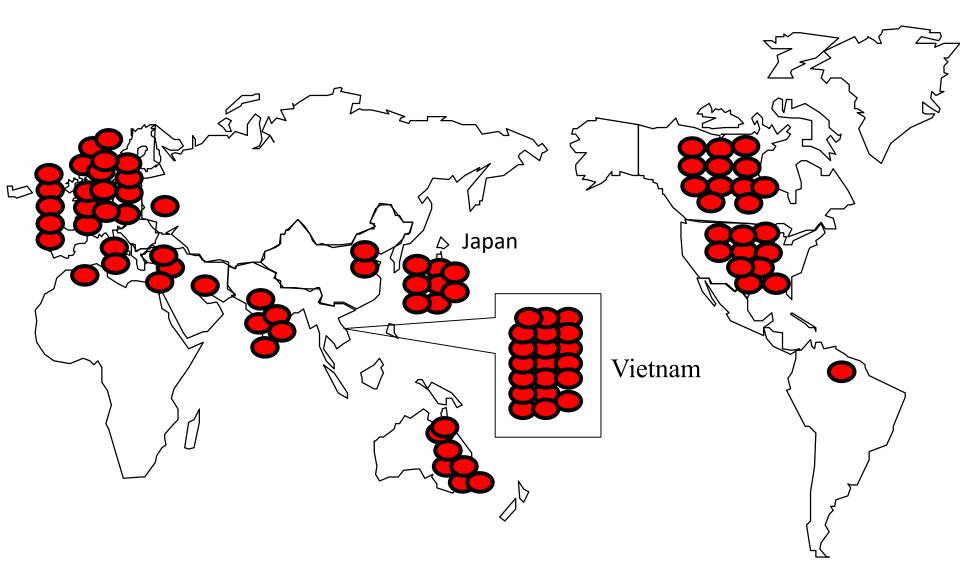




### Typical T2 deficiency:

- > Presents with ketoacidotic episodes.
- Characteristic biochemical abnormalities.

## T2 deficient patients whose mutations were confirmed in Gifu Univ



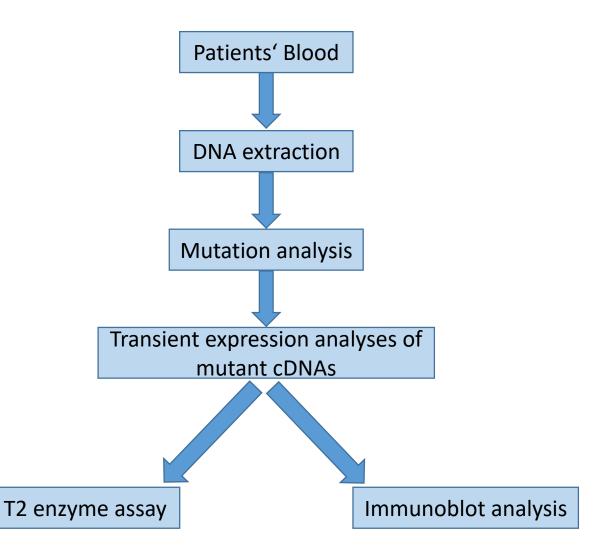
- T2 deficiency is an AR disease encoded by ACAT1 gene.
- ACAT1 gene is located at chromosome 11q22.3-q23.1.
- Spans about 27kb long, contains 12 exons and 11 introns.
- cDNA is about 1.5kb long and encodes a precursor protein of 427 amino acids.
- To date, more than 70 different mutations have been identified.
- No clear genotype-phenotype correlation.

 Herein, we report 10 Indians with beta-ketothiolase deficiency, describe their molecular characterization, and expression analysis of the identified mutations in *ACAT1* gene.

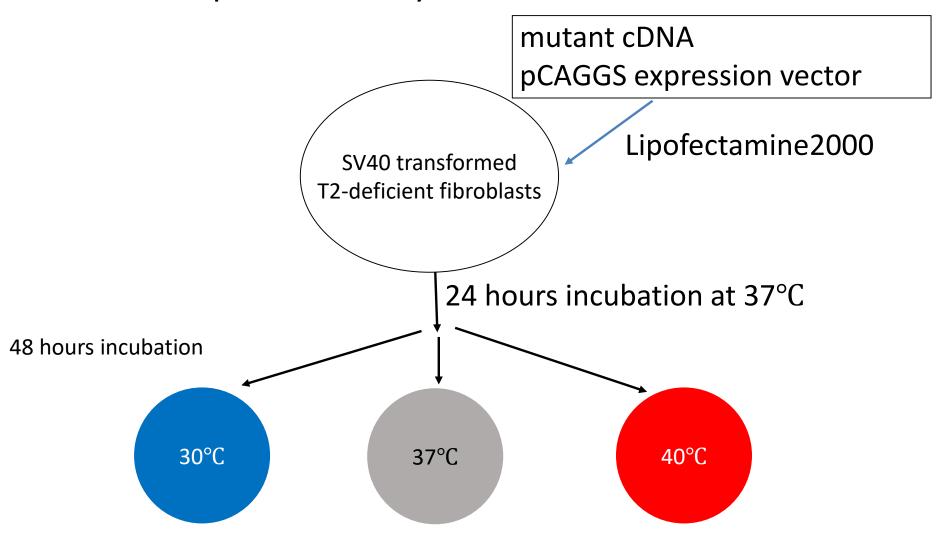
## **Patients**

Identifier	Gender	Predisposing condition	Age at onset (months)	Frequency	рН	Outcome
GK94	M	Fever, cough	7	1	6.5	Normal
GK98	M	Diarrhea	4	1	6.8	GDD, epilepsy
GK108	F	Poor feeding	11	1	6.9	Death
GK109	M	Vomiting	9	1	6.8	Normal
GK110	F	Fever, cough	11	1	7.0	Normal
GK111	M	Poor feeding	6	1	7.1	Normal
GK112	F	Fever, cough	7	3	7.15	Some hypotonia
GK113	M	Fever, cough	12	2	7.1	Normal
GK114	M	Fever, cough	19	1	6.9	Normal
GK99	[	Data are currently ur	navailable			

## **Methods** (summary)



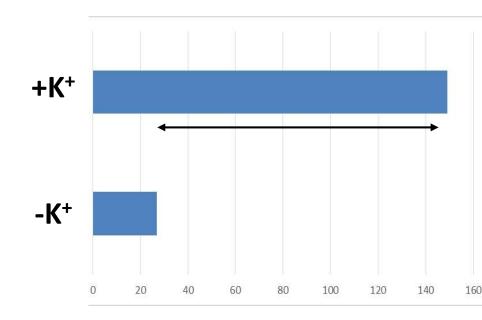
#### Transient expression analysis of T2 cDNA



enzyme assay and immunoblot analysis

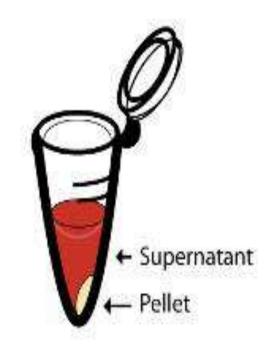
## Enzyme assay of Mitochondrial acetoacetyl CoAthiolase (T2)

- Based on the fact that T2
  is the only thiolase
  activated in presence of K<sup>+</sup>.
- Hence, the difference of thiolase activity in presence and absence of K+ represents the T2 activity.



#### Immunoblot analysis

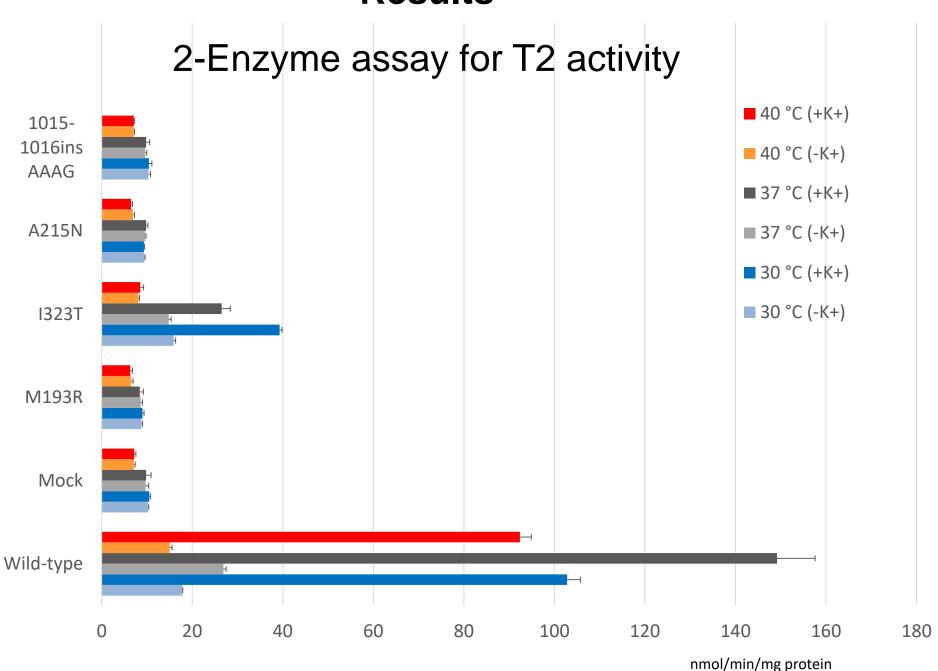
- The 1<sup>st</sup> antibody was a mixture of an anti-T2 antibody and anti-succinyl-CoA: 3oxoacid CoA transferase (SCOT) antibody.
- To assess the effects of mutations on protein solubility, we analysed not only the supernatants that were used for the enzyme assay but also the pellets of the cell extracts.



## 1- Mutation Analysis

Patients	Mutations in <i>A</i>	CAT1 gene
GK94	253_255delGAA (E85del)	253_255delGAA (E85del)
GK98	578T>G (M193R)	578T>G (M193R)
GK99	578T>G (M193R)	578T>G (M193R)
GK108	578T>G (M193R)	578T>G (M193R)
GK109	1015-1016insAAAG	1015-1016insAAAG
GK110	IVS7+1g>a	IVS7+1g>a
GK111	578T>G (M193R)	968T>C (I323T)
GK112	c.1124A>G	c.1124A>G
GK113	c.643_644delGCinsAA (A215N)	c.643_644delGCinsAA (A215N)
GK114	578T>G (M193R)	578T>G (M193R)

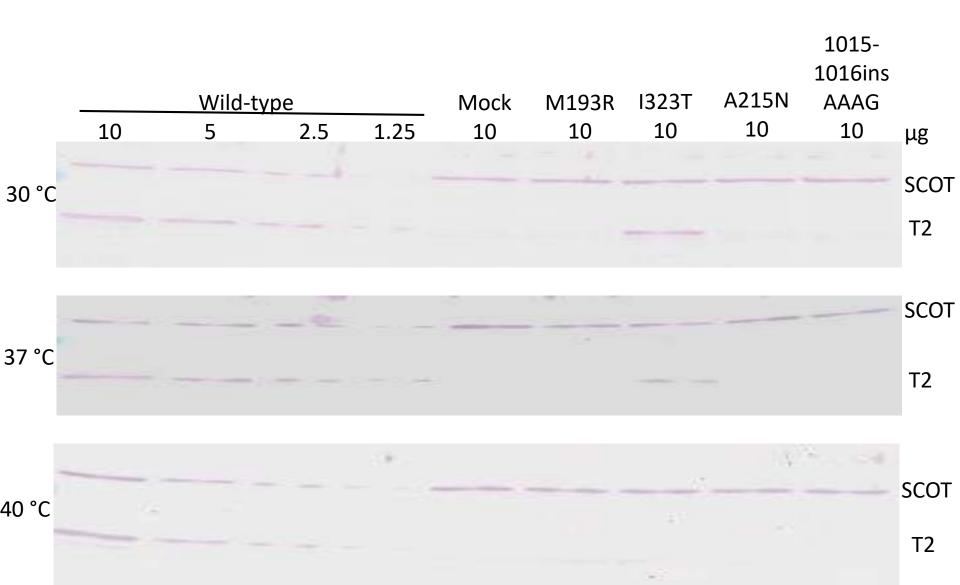




## 3-Immunoblot analysis (*supernatant*)

		Wild-ty	pe		Mock	M193R	1323T	A215N	1015- 1016ins AAAG	
	10	5	2.5	1.25	10	10	10	10	10	μg
					-	-				SCOT
37 °C										T2

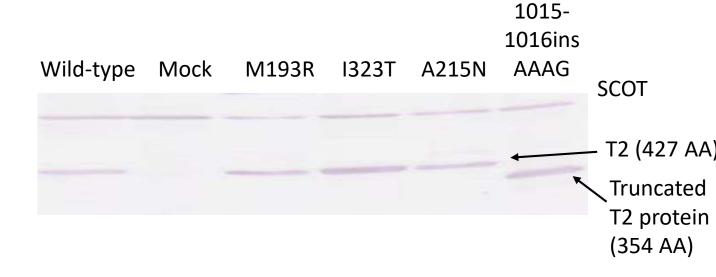
## 3-Immunoblot analysis (*supernatant*)



### 3-Immunoblot analysis (*Supernatant vs Pellet*)

Super	natant	S						1015-	
		_						1016ins	
	Wild-t	ype		Mock	M193R	1323T	A215N	AAAG	
10	5	2.5	1.25	10	10	10	10	10	μg
		_							SCOT
	_								T2





Mutations in ACAT1
 gene are highly heterogeneous;
 Only a few common mutations were identified.

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Case	Mu	utations	Case	Mu	tations
GK70	R208X	c.163_167del5ins2	GK89	IVS10−1g>c	IVS10-1g>c
GK72	R208X	R208X	GK90	R208X	A410V
GK73	R208X	R208X	GK91	R208X	R208X
GK74	IVS10−1g>c	R208X	GK93	R208X	R208X
GK75	R208X	R208X	GK100	IVS10-1g>c	Ex6-11del
GK76	R208X	R208X	GK101	R208X	R208X
GK79	R208X	R208X	GK102	R208X	S284N
GK80	R208X	R208X	GK103	R208X	IVS10−1g>c
GK86	IVS10-1g>c	IVS10-1g>c	GK104	R208X	R208X
GK87	R208X	R208X	GK105	R208X	c.1032_1033insA

Mutant allele frequency

R208X 70%
IVS10-1g>c 17.50%

#### **Indians**

#### **Vietnames**

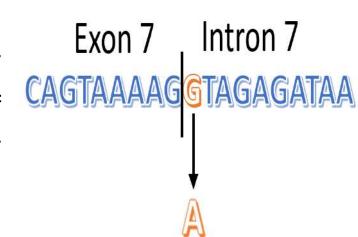
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GK73	R208X	R208X	GK91	R208X	R208X
GK74	IVS10−1g>c	R208X	GK93	R208X	R208X
GK75	R208X	R208X	GK100	IVS10−1g>c	Ex6-11del
GK76	R208X	R208X	GK101	R208X	R208X
GK79	R208X	R208X	GK102	R208X	S284N
GK80	R208X	R208X	GK103	R208X	IVS10−1g>c
GK86	IVS10-1g>c	IVS10-1g>c	GK104	R208X	R208X
GK87	R208X	R208X	GK105	R208X	c.1032_1033insA

Mutant allele frequency

R208X 70% IVS10-1g>c 17.50%

- Genetic analysis of the 10 Indians revealed
   4 novel mutations in ACAT1 gene:
- IVS7+1g>a mutation: affects a highly conserved point at the splice donor site of intron 7; hence, aberrant splicing (exon 7 skipping) is the usual result.
- Transient expression analysis of mutant T2 cDNA showed that M193R, A215N, and 1015-1016insAAAG are null mutations (no T2 enzymatic activity) and the corresponding mutant T2 proteins are insoluble.



contrast, • In **I323T** mutation resulted in an T2 unstable mutant protein that retained some activity (20%) at 37°C; Such activity increased to 40% at 30°C but ablated at 40 °C.

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#### **Summary and conclusions**

- We confirmed the diagnosis of beta-ketothiolase deficiency in 10 new patients from India.
- We identified 4 novel mutations in *ACAT1* gene.
- M193R appears to be a common mutation in T2-deficient Indians.

## Thank you