

Name	CRM NO	Lab ID	Father Name
Package TMS 52	Gender Female	DOB/Age	Referred By
Hospital name	City NOIDA	Pre Term / Full Term Full Term	Birth Weight (Kg)
Specimen Source Heel Prick	Specimen Notes Acceptable	Collection Date & Time	Received Date & Time
Date of Report & time	Blood Transfusion No	Transfusion Date No	Special Feeds / IVF / TPN/ Supplements No

INBORN ERRORS OF METABOLISM (IEM) SUMMARY REPORT

Sr. No.	Test Methodology	Result	Test Type
1	TANDEM MASS SPECTROMETRY SCREENING REPORT	NEGATIVE	Screening

DISCLAIMER

All investigations have their limitations which are imposed by the limits of sensitivity and specificity of individual assay procedures as well as the quality of specimen received by the laboratory.

It is presumed that the received specimen belongs to the patient named or identified in the test request form.

Partial reproduction of this report is not permitted.

It is important to monitor baby's health and take a proactive approach should there be any unhealthy condition(s) within or beyond scope of this test.

A negative screening assay result does not rule out the possibility of an underlying metabolic / genetic disease.

The metabolites and health conditions mapping is not mutually unique and there can be other forms in which an abnormality may be apparent. Similarly, the health condition may not exclusively be an outcome of the disorders in consideration

If the newborn is premature, retest is suggested when adjusted gestational age is 40 weeks.

HB-Variant hemoglobins may require further testing by the laboratory to diagnose.

HB-Presence of Hb Bart's with any newborn screen result indicates the presence of alpha thalassemia of unknown severity.

HB-If an increased amount of adult hemoglobin is noted, the infant is presumed to have been transfused (Including Intrauterine transfusion).

HB-Carriers of beta thalassaemia and HbLepore cannot be excluded at this age.

Investigations performed is not intended to replace the medical advice and or treatment obtained from a qualified healthcare practitioner as it doesnot cover all the metabolic disorders

HB (Covered under Hemoglobinopathies package)



Dr. Ankur Jindal (Ph.D)
Consultant Cytogenomics

Name	CRM NO	Lab ID	Father Name

TANDEM MASS SPECTROMETRY SCREENING REPORT

REPORT- SUMMARY

Observations:

All the analytes are within the normal range.

Interpretation:

The tested sample is screened **NEGATIVE** for the list of disorders in Annexure 1.

Recommendation:

Please correlate the results with other clinical and diagnostic findings.

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 Initial Report

 Duplicate Report

 Revision Report

 Version No

Analyte Name	Analyte Result(μmol/L)	Analyte Reference	MoM*	MoM Value	MoM Reference
Alanine	280.000	71.00-1361.24		1.14	NMT 5.56
Arginine	13.200	0.78-60.00		5.43	NMT 24.69
Citrulline	13.800	3.86-62.11		1.21	NMT 5.45
Glutamine	936.000	48.24-1309.25		1.83	NMT 2.56
Glycine	358.000	117.35-1270.00		0.93	NMT 3.29
leucine\isoleucine\hydroxyproline	134.000	23.93-383.93		1.06	NMT 3.02
Methionine	14.200	4.63-48.00		0.87	NMT 2.93
Ornithine	90.100	22.75 - 330.00		1.03	NMT 3.77
Phenylalanine	56.300	17.10 - 193.00		1.05	NMT 3.61
Proline	156.000	30.00 - 448.00		1.14	NMT 3.27
Tyrosine	72.700	15.34-455.88		0.97	NMT 6.10
Valine	117.000	31.28 - 450.00		1.03	NMT 3.95
Free Carnitine	25.900	7.00-121.04		1.03	NMT 4.82
Acetylcarnitine	15.600	2.49-65.00		0.95	NMT 3.81
Propionylcarnitine	2.780	0.12-7.02		1.71	NMT 4.31
Butyrylcarnitine	0.230	0.06-1.30		1.05	NMT 6.27
Isovalerylcarnitine	0.090	0.00-0.60		0.75	NMT 5.00
Hexanoylcarnitine	0.020	0.00-0.30		0.5	NMT 7.50
Octanoylcarnitine	0.040	0.00-0.37		0.67	NMT 6.17
Octenoylcarnitine	0.030	0.00-0.32		1	NMT 10.67
Decenoylcarnitine	0.060	0.01-0.24		1.5	NMT 6.00
Decadienoylcarnitine	0.010	0.00-0.07		1	NMT 7.00
Tetradecenoylcarnitine	0.120	0.01-0.51		1.71	NMT 7.30
Hexadecenoylcarnitine	4.090	0.00-9.04		1.14	NMT 2.53
3-Hydroxy-hexadecenoylcarnitine	0.030	0.00-0.09		1.5	NMT 4.50
Octadecenoylcarnitine	0.990	0.00-2.41		1.06	NMT 2.59
Octadecenoylcarnitine	1.710	0.00-3.25		1.37	NMT 2.60
Succinylacetone	0.200	0.00-1.00		0.8	NMT 4.00
Adenosine	0.450	0.00-3.52		0.76	NMT 5.97
2-deoxyadenosine	0.020	0.00-0.10		1	NMT 5.00
C26:0-lysophosphatidylcholine	0.650	0.0 - 1.06		2.17	NMT 3.53
Argininosuccinic acid	0.150	0.00-2.02		0.65	NMT 8.70
Glutamic acid	300.000	47.00-915.00		1.2	NMT 3.65
Propionylcarnitine/Acetylcarnitine	0.180	0.00-0.55		1.64	NMT 5.00
Methylmalonyl	0.170	0.00-1.50		1	NMT 8.82
Glutaryl carnitine	0.050	0.00-0.35		0.71	NMT 5.00
Malonylcarnitine	0.280	0.00-1.00		2.8	NMT 10.00

* This Graph represents the value corresponding to Multiple of Median(MoM). Each box represents the interval between 1%ile and 99%ile. The MoM Shown as dot.

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ANNEXURE - 1 LIST OF DISORDERS

Amino Acid Disorders	Organic Acid Disorders
Citrullinemia Type I	Malonic acidemia
Citrullinemia Type II	Methylmalonic acidemia (Cobalmin Disorders)
Hyperornithinemia with Gyral Atrophy	Methylmalonic acidemia (Mutase)
5-Oxoprolinuria	3-Methylglutaconic aciduria
Transient Neonatal Tyrosinemia	2-Methyl-Butyryl-Glycinuria
Argininosuccinic Acidemia	2-Methyl-3-HydroxyButyric Aciduria
Argininemia	3-Methylcrotonyl-CoA carboxylase deficiency
Defects of bipterin cofactor biosynthesis	Multiple Carboxylase Deficiency (also known as Holocarboxylase synthase deficiency)
Disorders of bipterin cofactor regeneration	3-Hydroxy-3-Methyl-Glutaric aciduria
Benign Hyperphenylalaninemia	Propionic acidemia
Homocystinuria	
Hypermethioninemia	Other Conditions
Non-ketotic hyperglycinemia	Adenosine deaminase (ADA)–severe combined
Phenylketonuria	Carbamoyl phosphate synthetase I
Tyrosinemia type I	Hyperornithinemia-Hyperammonemia-Homocitrullinuria
Tyrosinemia type II	X-linked adrenoleukodystrophy
Tyrosinemia type III	Treatment with antibiotics/Pyvalic Acid
Maple syrup (urine) disease	Treatment with Benzoate or Valproic Acid
Fatty Acid Oxidation Disorders	Hyperalimantation (TPN)
Medium/short-chain L-3-hydroxyacyl-CoA dehydrogenase deficiency	Liver Complications
Very long-chain acyl-CoA dehydrogenase deficiency	Medium Chain Triglyceride (MCT) Oil Administration
Short-Chain Acyl-CoA Dehydrogenase Deficiency	N-acetylglutamate synthase deficiency
Trifunctional protein deficiency	Ornithine transcarbamylase deficiency
Glutaric acidemia type II	
Medium-Chain Acyl-CoA Dehydrogenase Deficiency	
Medium-chain ketoacyl-CoA thiolase deficiency	
Carnitine Acyl-Carnitine Translocase deficiency	
Carnitine Palmitoyl-Transferase type II deficiency	
2,4 Dienoyl-CoA reductase deficiency	
Carnitine Palmitoyl Transferase Type I Deficiency	
Carnitine Uptake Defect	
Organic Acid Disorders	
Isobutyryl-CoA-Dehydrogenase Deficiency	
Glutaric Acidemia type I	
Iso-Valeric Acidemia	
Beta-KetoThiolase deficiency	
Long-chain L-3 hydroxyacyl-CoA dehydrogenase deficiency	

REFERENCES

1. Ombrone D, Giocaliere E, Forni G, Malvagia S, la Marca G. Expanded newborn screening by mass spectrometry: New tests, future perspectives. Mass Spectrom Rev. 2016 Feb;35(1):71?84.

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2. It is to be presumed that the tests performed pertain to the specimen/sample attributed to the Customer's name or identification. It is presumed that the verification particulars have been cleared out by the customer or his/her representation at the point of generation of said specimen / sample. It is hereby clarified that the reports furnished are restricted solely to the given specimen only.
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