

		LABORATO	RY REPORT			
Name	:Mr ANUP SHAF	RMA	Sex/Age : N	lale/28 Years	Case ID	:40521602317
Ref By	:DR VINAY CHO	PRA	Dis.Loc. :		Pt ID	:
Bill. Loc.	:KOS DIAGNOST	TIC LAB			Pt. Loc.	:
Registratio	n Date & Time	: 13-May-2024	Sample Type	: Heparin Whole Blood - Na	Ph#	:
		08:52				
Sample Da	ite & Time	: 13-May-2024	Sample Coll.By	<i>'</i> :	Ref Id	•
		08:52				
Report Da	Report Date & Time : 20-Ma		Acc. Remarks	:	Ref Id 2	:
		01:46				

Chromosome Analysis Report

Clinical History	No clinical history available.
Karyotype (ISCN Nomenclature 2020)	46,XY

<u>Interpretation</u>	Normal Karyotype

Banding Method : GTG : 72hrs PHA stimulated **Culture Type Banding Resolution** : Approx 550 **Metaphases Counted** :20 **Metaphases Analyzed** :20 Metaphase Karyotyped :05 **Proliferative Index** : Good **Quality of Metaphases** : Good

For specimens received from non NCGM locations, it is presumed that it belongsto the patient as identified on the labels of the container/Test Requisition Formand it has been verified as per GCLP (Good Clinical Lab Practices) by the referrer atthetime of collection of the specimen. NCGM's responsibility is limited to the analytical part of the assay performed.

Dr. Sandip Shah M.D. (Path. & Bact.)

Consultant Pathologist

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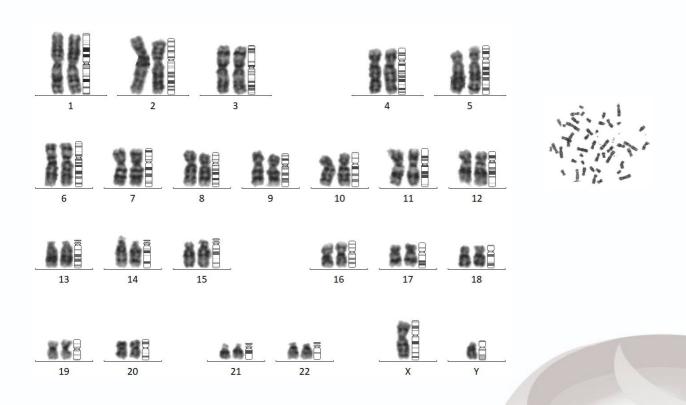
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Karyogram and Metaphase



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Bodakdev, Ahmedabad -380059



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		01:46				

Limitation

The error rate of the test is 0.5%. The normal report does not rule out very Low grade mosaicism, minor chromosomal anomalies, and deletion, Duplication or Inversion at very subtle level. The report should be interpreted in accordance with the counselling provided before the test and with the report. A standard G-banded Karyotype usually has a resolution of around 5 Mb.

<u>Disclaimer</u>

Polymorphic variants have not been reported as these variants are not associated with specific disease or phenotype. Cytogenetically visible polymorphic variants include variants involving heterochromatin (variant size), satellite size, pericentric inversions (heterochromatic or euchromatic regions) [e.g., 1qh+/qh-, 9qh+/qh-, 16qh+/qh-, acrocentric p+ or p-, Yqh+/qh-, inv(9)(p11q13), inv(2)(p11.2q13)] and also euchromatic variants (e.g., located on 4p16, 8p23.1, 9p12, 9q13-q21.12, 15q11.2, 16p11.2).

Reference: Silva, M., de Leeuw, N., Mann, K., Schuring-Blom, H., Morgan, S., Giardino, D., Rack, K. and Hastings, R., 2019. European guidelines for constitutional cytogenomic analysis. *European Journal of Human Genetics*, 27(1), pp.1-16.

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Registration	on Date & Time	: 13-May-2024 08:52	Sample Type	: Heparin Whole Blood - N	la Ph#	:
•		: 13-May-2024 08:52	Sample Coll.By	:	Ref ld	:
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----- End Of Report -----

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