

(A Unit of KOS Healthcare)



Dr. Vinay Chopra MD (Pathology & Microbiology) Chairman & Consultant Pathologist Dr. Yugam Chopra MD (Pathology) CEO & Consultant Pathologist

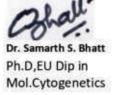
		LABORATO	RY REPORT	0			
Name	:Mr KARAN		Sex/Age	: Male/18 Years	Case ID	:40621600009	
Ref By	:DR VINAY CHOPRA		Dis.Loc. :		Pt ID		
Bill. Loc.	:KOS DIAGNOS	TIC LAB			Pt. Loc.		
Registratio	n Date & Time	: 01-Jun-2024 09:32	Sample Typ	: Heparin Whole Blood	-Na Ph#	3	
Sample Date & Time : 01-Jun-2024 09:32		Sample Co	II.By :	Ref Id			
Report Da	te & Time	: 12-Jun-2024 01:27	Acc. Rema	rks :	Ref ld 2		

## Chromosome Analysis Report

Clinical History	No clinical history available,				
Karyotype (ISCN Nomenclature 2020)		46,XY			
Interpretation		Normal Karyotype			
Banding Method	: GTG	Culture Type	: 72hrs PHA stimulated		
	Annew 550	Matanhagas Counted	: 20		
Banding Resolution	: Approx 550	Metaphases Counted	. 20		

Quality of Metaphases

# For specimens received from non NCGM locations, it is presumed that it belongs to 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 at the time of collection of the specimen. NCGM's responsibility is limited to the analytical part of the assay performed.



Proliferative Index

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: Good



: Good



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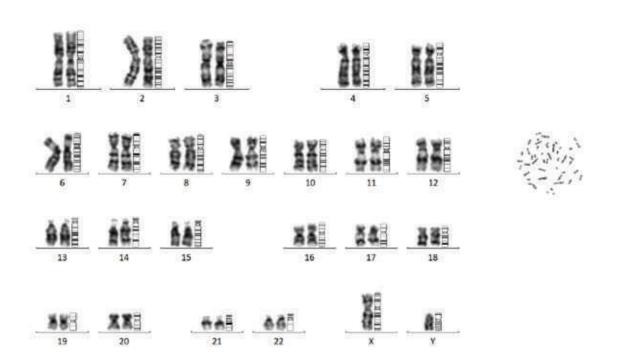


Dr. Vinay Chopra
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Chairman & Consultant Pathologist

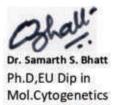
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## Karyogram and Metaphase



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	LABORATORY REPORT						
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#### 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.

### Disclaimer

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).

<u>Reference:</u> 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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Dr. Samarth S. Bhatt Ph.D,EU Dip in Mol.Cytogenetics

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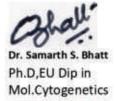


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----- End Of Report -----

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