



ISO 9001 : 2008 CERTIFIED LAB

# KOS Diagnostic Lab

(A Unit of KOS Healthcare)



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

**Dr. Yugam Chopra**  
MD (Pathology)  
CEO & Consultant Pathologist

## LABORATORY REPORT



Name : Mrs. JATINDER KAUR	Sex/Age : Female/ 32 Years	Case ID : 30100113379
Ref By :	Dis. Loc. :	Pt ID :
Bill. Loc. : Neuberger Diagnostics Pvt Ltd Delhi		Pt. Loc. :
Registration Date & Time : 12-Jan-2023 09:44	Sample Type : Heparin Whole Blood - Na	Ph # :
Sample Date & Time : 12-Jan-2023 09:44	Sample Coll. By :	Ref Id : NDPL - DELHI
Report Date & Time : 19-Jan-2023 19:27	Acc. Remarks :	Ref Id 2 : 2155147869

## Chromosome Analysis Report

<b>Clinical History</b>	--
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<b>Karyotype (ISCN-2020)</b>	46,XX
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<b>Interpretation</b>	Normal Karyotype
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<b>Banding Method</b> : GTG	<b>Culture type</b> : 72hrs PHA stimulated
<b>Banding Resolution</b> : 400	<b>Metaphases Counted</b> : 20
<b>Metaphases Analyzed</b> : 20	<b>Metaphases Karyotyped</b> : 5

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**Dr. Samarth S. Bhatt**  
Ph.D, EU Dip in  
Mol. Cytogenetics

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### NOTE:

***This Sample was outsourced***

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**LABORATORY REPORT**

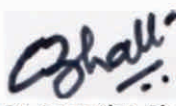


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<b>Proliferative Index</b>	: Average	<b>Quality of Metaphase</b>	: Average
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**Karyogram**

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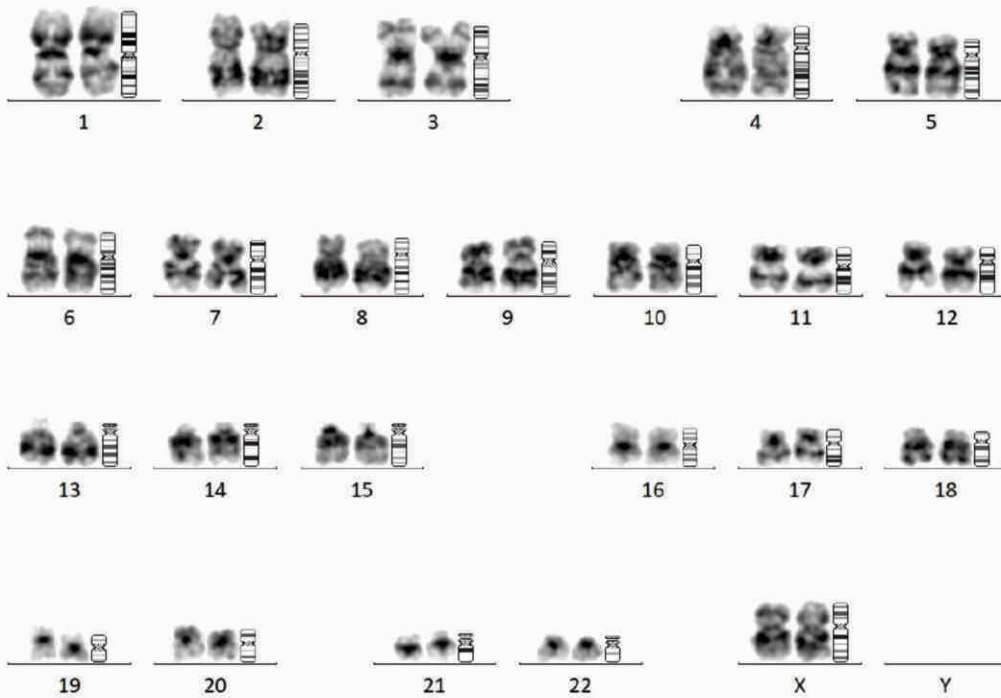
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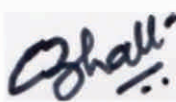
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**Metaphase**

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**Limitations**

The error rate of the test is 0.5%. The normal report does not rule out minor chromosomal anomalies, mosaicism, malformation, fragile X syndrome and other genetic disorders. The report should be interpreted in accordance with the counseling provided before the test and with the report.

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**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).

**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.

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

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