

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

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

### LABORATORY REPORT



Name : Mr. BALVIR	Sex/Age : Male/27 Years	Case ID : 30300129010
Ref By :	Dis. Loc. :	Pt ID :
Bill. Loc. : Neuberg Diagnostics Pvt Ltd Delhi		Pt. Loc. :
Registration Date & Time : 20-Mar-2023 16:45	Sample Type : Heparin Whole Blood - Na	Ph # :
Sample Date & Time : 20-Mar-2023 16:45	Sample Coll. By :	Ref Id : 2155147846
Report Date & Time : 30-Mar-2023 18:08	Acc. Remarks :	Ref Id 2 : DELHI

### Chromosome Analysis Report

#### Clinical History

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#### Karyotype (ISCN-2020)

46,XY

#### Interpretation

Normal Karyotype

Banding Method	: GTG	Culture type	: 72hrs PHA stimulated
Banding Resolution	: 550	Metaphases Counted	: 20
Metaphases Analyzed	: 20	Metaphases Karyotyped	: 5
Proliferative Index	: Good	Quality of Metaphase	: Good

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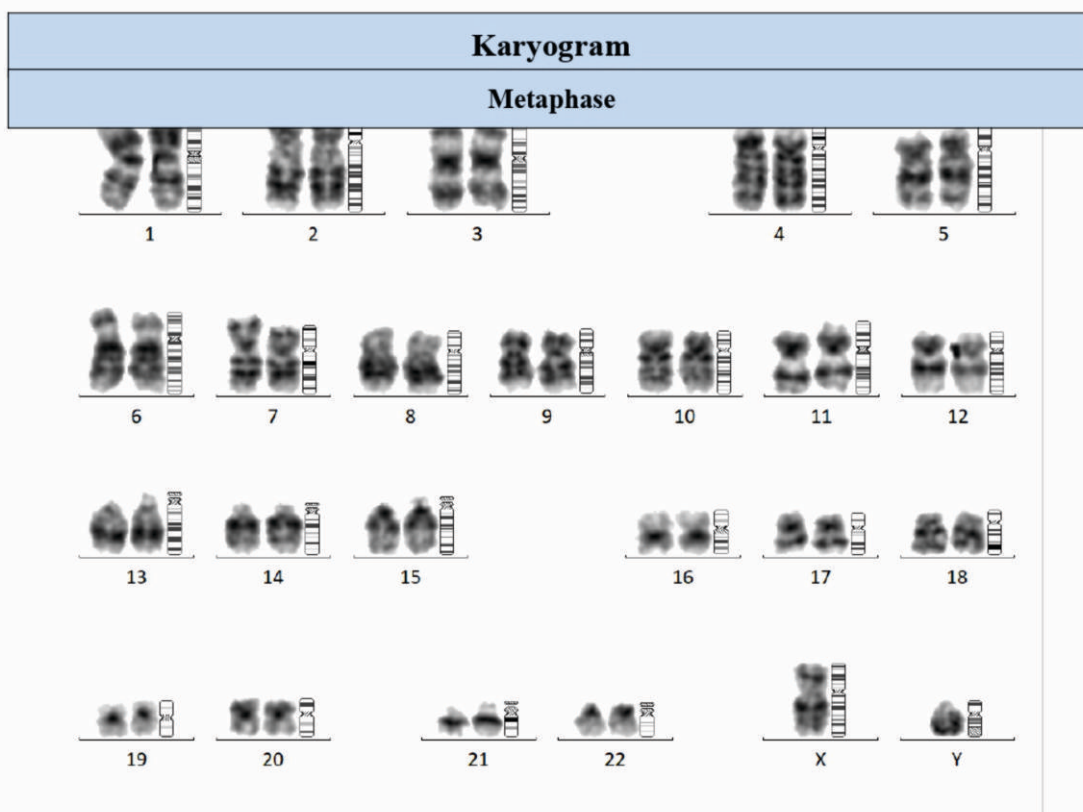
### NOTE:

***This Sample was outsourced***

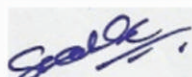
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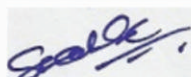
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### Limitations

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.

### Disclaimer

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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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