DocuSign Envelope ID: 5BBF8D7B-3C9E-4FC9-8359-87A03CB8EB19 Case: Jax G-banding_S008770

Project Quote#:	200401	Specimen:	iPSC
LIMS ID:	S008770	Received Date:	5/19/2022
Customer Sample ID:	26.1_R_A5	Completed Date:	7/29/2022
Gender:	Male	Band Resolution:	400
Total Counted:	20		
Total Analyzed:	20		
Final Karyotype:	46,XY[20]		
Case Notes:	G-banded chromosome analysis of metaphase cells designated 26.1_R_A5(KromaTiD Sample ID S008770) shows a normal male karyotype. The other abnormalities/aberrations detected were non-clonal and were designated as low-level mosaicism or random gain/loss.		

Karyotype Summary:

Karyotype:	# Cells
46,XY	14
45,XY,-17	1
45,XY,-18	1
44,X,-Y,-12	1
44,XY,-7,-12	1
44,XY,-3,-3	1
45,XY,-19	1

Cells Images:

Karyotyped: 46,XY 1-61	Karyotyped: 46,XY 3-3
1 2 3 4 5 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 X Y	1 2 3 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 X Y

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Limitations: This assay allows for microscopic visualization of numerical and structural chromosome abnormalities. The size of structural abnormality that can be detected is >3-10Mb, dependent upon the G-band resolution obtained from this specimen. For the purposes of this report, band level is defined as the number of G-bands per haploid genome. Detection of heterogeneity of clonal cell populations in this specimen is limited by the number of metaphase cells analyzed, documented above as "number of cells counted". Results are for Research Use Only and should not be used for clinical purposes.

Completed By/Date: Michael Vernich Cytogenetics Supervisor

DocuSigned by: 7/30/2022 Michael Vernich B510035B47034EE...

DocuSigned by:

Approved By/Date: Gregory Husar Operations Manager

7/30/2022 Greg Husar 8836BEA4EF644E7...

Report Date: Friday, July 29, 2022