DocuSign Envelope ID: A039B54B-017E-4328-A6FC-B9336BDE10DA Case: Jax G-banding_S008337

Project Quote#:	200401	Specimen:	iPSC	
LIMS ID:	S008337	Received Date:	4/27/2022	
Customer Sample ID:	22_R-E05	Completed Date:	7/6/2022	
Gender:	Male	Band Resolution:	425	
Total Counted:	20			
Total Analyzed:	20			
Final Karyotype:	46,XY[20]			
Case Notes:	G-banded chromosome analysis of metaphase cells designated 22_R-E05 (KromaTiD Sample ID S008337) 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	17
46,XY,dup(14q)	1
45,XY,-15	1
47,XY,+M	1

Cells Images:

Karyotyped: 46,XY 4-56		Karyotyped: 46,XY 4-71	
71 15 36	ik ji	$\left\{ \sum_{i} \right\}_{2} \in \left\{ e \right\}_{3}$	XX
		75 12 78 18 16	
N N	17 18	10 10 10 10 10 13 14 15 16 16 88 86 86 86 36);
19 20 21 22	X Y	19 20 21 22	X Y

DocuSign Envelope ID: A039B54B-017E-4328-A6FC-B9336BDE10DA

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: 7/8/2022 Michael Vernich B510035B47034EE...

Approved By/Date: Gregory Husar Operations Manager — DocuSigned by: 7/8/2022 Grig Husar — 8836BEA4EF644E7...