

Document Code:	FORM-0068A	Document Type:	FORM
Revision:	3.0	Date Effective:	08-27-21

TITLE: KROMATID CHROMOSOME ANALYSIS REPORT

I. ASSAY INFORMATION

Project Quote #	Q200401
Specimen Type	iPSCs
Body Site	N/A
Sample ID	13-F06 (S008319)
	Male
Cell Line Gender	
Passage number (or N/A)	N/A
Study Objective	The purpose of this study is to characterize iPS cells grown <i>in vitro</i> , designated for cytogenetic analysis.

II. CELL MAINTENANCE

Culture vessel	N/A
Media	N/A
Density (estimated)	N/A
Culture atmosphere	N/A
Culture maintenance	N/A



U	137 304 10			
	Document Code:	FORM-0068A	Document Type:	FORM
	Revision:	3.0	Date Effective:	08-27-21

Culture Maintenance Process Description	N/A
	Analyst Initial/Date: N/A

III. CULTURE HARVEST

Culture Harvest Process Description	N/A
	Analyst Initial/Date: N/A

Material	Usage information	
Harvest materials (trypsin, EDTA, etc.)	Type: N/A LN/ Exp. Date: N/A	
Colcemid	LN/ Exp. Date: N/A Concentration: 0.1 µg/mL (10 µL/mL) Incubation time: N/a	
Hypotonic	LN/ Exp. Date: N/A Solution: N/A Incubation time:	
Fixative	Prepared Fresh, day-of-use	



•	107 004 10			
	Document Code:	FORM-0068A	Document Type:	FORM
	Revision:	3.0	Date Effective:	08-27-21

IV. STAINING

Solution Type	Lot#	Exp. Date	Solution Type	Lot#	Exp. Date
Isoton II Diluent	4710610	07/12/22	Wright Stain	210817-Wright	08/17/22
Pancreatin	SLCD9444	01/11/23	Gurr Buffer	220426-Gurr	05/26/22
FBS	20J481	01/06/24	Permount	210201-01	02/1/23

Description	A sample of fixed cells 13-F06 (KromaTiD Sample ID S008319) was received at KromaTiD on 4-20-22. The fixed cells were washed twice with fixative (prepared fresh day-of-use) and the O.D. was adjusted. Drops of the final cell suspension were placed on clean slides and aged for 60 minutes at 90°C. Slides were digested in a pancreatin solution with Isoton II diluent. The enzymatic reaction was then stopped by rinsing with FBS, followed by application of a stain solution (3:1 Wright/Gurr buffer) which was poured on the slides so that it covered the entire surface. After staining for up to 1 minute, slides were washed with deionized water for 1-5 seconds and air dried. The mounting medium Permount was applied to the slides, a coverslip was placed on the slide and the slides were scanned on the microscope.
	Analyst Initial/Date: MV 5/10/22

TEST DESCRIPTION:

G-banding with trypsin treatment and Giemsa stain (GTG-banding) is used in cytogenetics to produce a visible karyotype by staining metaphase chromosomes. This technique allows each chromosome to be distinguished by its characteristic banding pattern. G-banding is useful in assessing structural abnormalities in individual chromosomes, as well as extra or missing chromosomes within a cell. Industry-standard protocols for scoring and describing results were used (ISCN 2016: An International System for Human Cytogenomic Nomenclature).



٠,	37 304 10			
	Document Code:	FORM-0068A	Document Type:	FORM
	Revision:	3.0	Date Effective:	08-27-21

V. RESULTS

Cells Counted	20	Total Karyograms	2
Cells Analyzed	20	Average Band Resolution	425
Image File Location		Jax Gbanding_S008319	

5.1 CHROMOSOME COUNT PER 20 METAPHASES

Of the 20 cells counted, 19 contained 46 chromosomes (95%). *Cells containing greater than 57 chromosomes are recorded as polyploid.* The polyploid frequency was 0%, based on the metaphases counted.

5.2 CHROMOSOME ABERRATION DATA

The chromosome aberration data via G-band for the 20 metaphases examined is summarized in attached case report cell list. 0 chromosome aberrations were found in the 20 cells analyzed with 0% of the cells aberrant.

*Note: Cells with an euploidy gain/loss were found to be non-clonal, and therefore not included in the aberration data below.

Tech		Additional Comments	
Summary			
Karyotype	46,XY[20]	Normal Male Karyotype	
Cells Analyzed	20		
Normal Cells	20	Random loss/gain cells normalized	
Abnormal Cells	0		
Aberration	N/A	Non-clonal	
Type			
Aberration %	0%		



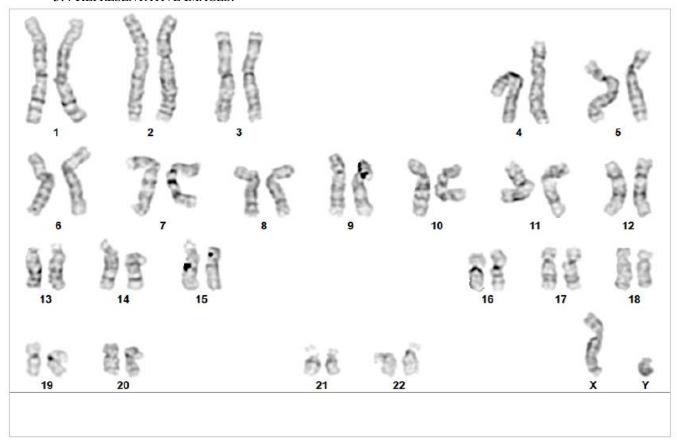
137 904 10			
Document Code:	FORM-0068A	Document Type:	FORM
Revision:	3.0	Date Effective:	08-27-21

5.3. INTERPRETATION/ SIGNIFICANCE:

G-banded chromosome analysis of metaphase cells designated 13-F06 (KromaTiD Sample ID S008319) shows a normal male karyotype 46,XY[20].

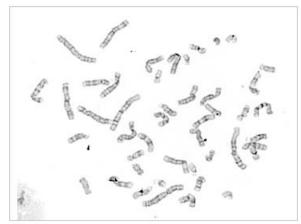
The other abnormalities/aberrations detected were non-clonal and were designated as low-level mosaicism or random gain/loss.

5.4 REPRESENTATIVE IMAGES:



Cell Results: Karyotyped: 46,XY

Cell Notes:

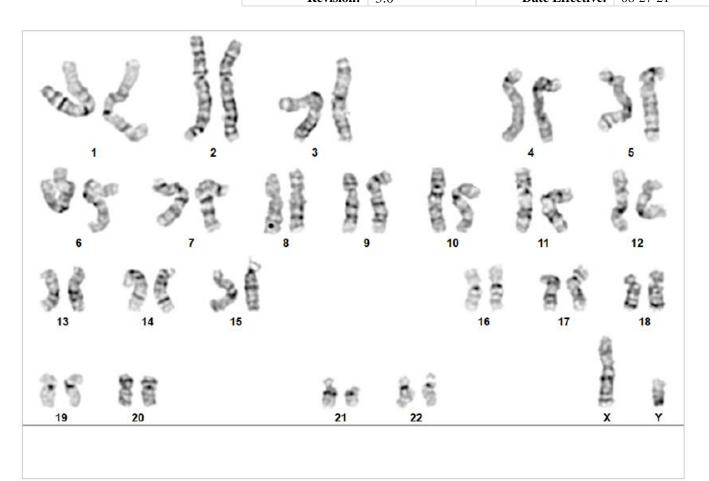


Label - Slide/Cell: \$008319 - 1/10 X,Y: 12.8 , 14.9



Document Code: FORM-0068A Document Type: FORM

Revision: 3.0 Date Effective: 08-27-21



Cell Results: Karyotyped: 46,XY

Cell Notes:



Label - Slide/Cell: S008319 - 1/15 X,Y: 14.9 , 18.2

Report Date: Tuesday, May 31, 2022



Document Code:	FORM-0068A	Document Type:	FORM
Revision:	3.0	Date Effective:	08-27-21

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:

— Docusigned by:

Michael Vernich

5/3

5/31/2022

Michael Vernich Cytogenetics Supervisor

Approved By/Date:

Docusigned by:

Gry Husar

5/31/2022

Gregory Husar Operations Manager



Document Code: FORM-0068A

Document Type:

FORM

Revision: 3.0

Date Effective: 08-27-21

				Kevision. 3.0	Date Effective.	00-27-21			
Table 1: Chromosome Analysis for sample 13-F06 (S008319). 20 cells were analyzed.									
#	Slide	Cell	Coordinates	Results	Analysis State	State By			
Slide: Name: 1 Label: S008319									
1	1	4	13.83 X 9.91	Karyotyped: 46,XY	Karyotyped	kstone			
2	1	9	11.46 X 13.46	Karyotyped: 46,XY	Karyotyped	kstone			
3	1	10	12.75 X 14.88	Karyotyped: 46,XY	Karyotyped	kstone			
4	1	12	14.24 X 15.71	Karyotyped: 46,XY	Karyotyped	kstone			
5	1	15	14.95 X 18.24	Karyotyped: 46,XY	Karyotyped	kstone			
6	1	16	14.40 X 18.33	Karyotyped: 46,XY	Karyotyped	kstone			
7	1	20	14.68 X 20.04	Karyotyped: 46,XY	Karyotyped	kstone			
8	1	27	14.88 X 22.71	Karyotyped: 46,XY	Karyotyped	kstone			
9	1	29	14.77 X 23.47	Karyotyped: 45,X, −Y	Karyotyped	kstone			
10	1	30	15.07 X 24.71	Karyotyped: 46,XY	Karyotyped	kstone			
11	1	32	6.65 X 25.07	Karyotyped: 46,XY	Karyotyped	kstone			
12	1	33	7.33 X 25.10	Karyotyped: 46,XY	Karyotyped	kstone			
13	1	35	14.62 X 27.62	Karyotyped: 46,XY	Karyotyped	kstone			
14	1	39	4.70 X 26.45	Karyotyped: 46,XY	Karyotyped	kstone			
15	1	40	5.94 X 27.72	Karyotyped: 46,XY	Karyotyped	kstone			
16	1	44	16.07 X 28.04	Karyotyped: 46,XY	Karyotyped	kstone			
17	1	48	9.64 X 29.81	Karyotyped: 46,XY	Karyotyped	kstone			
18	1	51	5.40 X 30.18	Karyotyped: 46,XY	Karyotyped	kstone			
Slide: Name: 2 Label: S008319									
19	2	4	7.92 X 12.63	Karyotyped: 46,XY	Karyotyped	kstone			
Slide: Name: 3 Label: S008319									
20	3	2	17.24 X 15.46	Karyotyped: 46,XY	Karyotyped	kstone			