# CELL Quality Control Report: Karyotyping analysis

**Technical Form** 

### Ref of QCR: iPS W11P47\_QCR-K\_v0.1\_230503\_LK

## **1. EXPERIMENTAL INFORMATION**

- Team / Project title: Génomique et physiopathologie des maladies cardiovasculaires (U1166-ICAN) / Projet EHT-CAVD
- Provider name: BOBIN Pierre
- Comments: human iPS cell line karyotype integrity validation
- Fixation and hypotonic choc were performed by Lina El Kassar

#### 1.1. Cells information

I-Stem

HROUGH INNOVATION

Type of QC	Cell type	N° of	Passage	Derive	d from:	Date of QC	Investigator
Type of QC	een type	differentiation	Iussuge	Cell	Passage	(YYMMDD)	initials
Karyotype	iPS : W11		P47	iPS 3.3	P24	230503	LK

- Date of freezing: NA
- Date of thawing : NA
- If iPSC, reprogramming method: reprogramming method: Episomal vectors (Okita *et al*, 2011) used on dermal fibroblasts to generate the 3.3 line.
- Specie of cells: Human

#### **1.2.** Culture conditions

- Culture medium: mTeSR1®
- Matrix or feeder type: Matrigel®
- Dissociation method: Gentle Cell Dissociation Reagent®
- If relevant, seeding conditions (dilution, density, ...):
- Culture with or without antibiotic, if yes which one: Without
- If feeder free adapted hPSC culture, for how many passage pN+X: None
- If differentiated cells, give culture conditions and cells information of hPSC: None

### 1.3. Test conditions

Performed: Prior freezing If Not applicable, please comment here:

After thawing

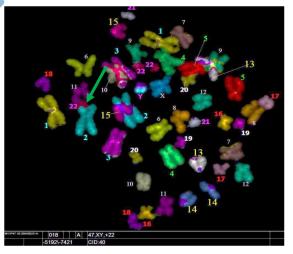
Not applicable

### 2. ANALYSIS

- Analysis conformity:
  - The karyotyping has been completed according to:
    - G-Banding (Instruction 4ITec0040) + R-Banding
      - mFISH (Instruction 4ITec0214)

<u>×</u>			Technical Form				
	Ref.: 4FTec0012v4EN			Application date: 2016.02.26			
	CEL	L Quality ( Karyotypi	Control Repo ng analysis	rt:			
Num Num	Yes otype conformity: ber of analyzed cells: ber of karyotyped cell lts (number of chromo	ls: 6	of cell line): 46,X	Y			
3. INTERP	RETATION						
The cell karyot	yping is therefore:	🔀 Norm	al 🗌 Abnorm	al			
<u> </u>		<u>.</u>	<u>×</u> •11 •21	Image: state Image: state<			
-6## •7## -	880 <u>990</u> <u>810</u>	·11== •12==	•688 ×788 4				
•13 • • • • • • • • • • • • • • • • • •	•15 •••••••••••••••••••••••••••••••••••	<b>8 6</b> 17 <b>1 1 1 1 1 1</b>	•13■ •14■ •				
• 19 • • 20 • • • 20		X	 	•21 •• • • • • • • • • • • • • • • • • •			
4. COMME	3D:44	46	wither 18 2042023-A 030 A 46,X -1416h-3226 CID				
4.1. Invest	tigator comments						
- On	e metaphase is 47,	XY,+22 (green	arrow). Chron	nosome 22 was checked in 49			

metaphases. This anomaly is not significant.





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- One metaphase is **46**, **XY**,**t**(**1**;**4**), (red arrows)). Chromosome 1 and chromosome 4 were checked in 49 metaphases. This anomaly is not significant.

