Certificate of Analysis (CoA) for induced Pluripotent Stem Cells



This product is for research only

ECACC Catalogue No: 66540493

Cell Line Name	UKKi025-C	Batch Number	P001
Donor ID	NP0135		
Disease Association	Hypertrophic Cardiomyopathy	Phenotype of Donor	Affected
Tissue of Origin	PBMC	Sex	Female
Reprogramming Method	Non-integrating Sendai virus (POU5F1, SOX2, KLF4, MYC)		
Passage Number	Passage 37	Cell number / vial	1.1x10 ⁶
Culture Matrix	Vitronectin	Culture Medium	Essential 8 TM /Essential 8 Flex TM
O ₂ Concentration	20%	CO ₂ Concentration	5%
Passaging Method	EDTA	Additional Culture Information	N/A
Cryopreservation Medium	90% medium / 10% DMSO		
Recommendation for thawing	Recommended thaw into 2 wells of a 6-well plate or per 10cm ²		
Recommendation for thawing	Refer to cell line user protocols for further guidance at www.EBiSC.org		
Additional Comments	Slow recovery after thaw, typical growth to confluency		
Associated Publications	N/A		

Please see www.EBiSC.org for further information on Quality Control applied to lines released by EBiSC. The following standard testing criteria have been determined within EBiSC, prior to release of this product:

Test	Assay	Acceptance Criteria	Result
Sterility -	Inoculation for microbiological growth	Not Detected	Pass
	qPCR for Mycoplasma	Not Detected	Pass
Cell Line Identity	Short Tandem Repeat analysis using PCR	N/A	Allele data recorded and available upon request. Gender match to donor
Viability	Visual Assessment	Growth to confluence post-thaw	Low, slow recovery
Phenotype	Continuous visual assessment of iPSC colony morphology	Recorded	Obvious iPSC colonies with medium differentiation levels



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Additional cell line characteristics have been determined by original reprogramming centres and have not been independently verified by EBiSC. Historical cell line data displayed here is accurate according to data provided by depositors on 30-MAY-2017

Test	Assay	Result
Phenotype	Flow Cytometry	Positive Expression of SSEA-1, SSEA- 4 and TRA-1-80
Тиспосурс	Immunocyto-chemistry	Positive expression of TRA-1-80, POU5F1, Nanog and SSEA-4
Karyotype	SNP Analysis (OmniExpress Exome Chip)	No larger chromosomal aberrations observed
Cell Line Identity	PowerPlex 16 STR Genotyping System	Match to donor profile
Clearance of Reprogramming Factors	PCR for Sendai virus	Not detected
Pluripotency	PCR	Pluripotency markers detected
Differentiation Potential	Trilineage differentiation	Differentiation to endoderm, ectoderm and mesoderm detected
Sterility	Virology (HBV, HCV, HIV1, HIV2) PCR	Not detected
Genetic Lesion	DNA sequencing Myh7 affected location	Mutation Myh7: c.2156G>A; Arg719Glu confirmed

The following guidance can be found in the Instructions for Use			
Intended use	Expiry Date		
Product Format	Recommended storage conditions		
Volume	Hazardous Information		

Approved CoA

Signature Company of the Company of

Date 26 feb 2018

