Certificate of Analysis (CoA) for induced Pluripotent Stem Cells



This product is for research only

ECACC Catalogue No: 66540481

Cell Line Name	UKKi019-B	Batch Number	P001
Donor ID	NP0081		
Disease Association	Familial Long QT Syndrome	Phenotype of Donor	Affected
Tissue of Origin	РВМС	Sex	Female
Reprogramming Method	Non-integrating Sendai virus (POU5F1, SOX2, KLF4, MYC)		
Passage Number	Passage 41	Cell number / vial	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	Typical 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	Acceptable
Phenotype	Continuous visual assessment of iPSC colony morphology	Recorded	Typical iPSC colonies with low 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 23-MAY-2017

Test	Assay	Result
Phonoismo	Flow Cytometry	Positive Expression of CD90, SSEA-1, SSEA-4 and TRA-1-80
Phenotype	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 hERG affected location	Mutation hERG: c.1280A>G; Tyr427Cys 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 Cue Cub

Date 23 Job 2018

