Certificate of Analysis (CoA) for induced Pluripotent Stem Cells



This product is for research only

ECACC Catalogue No: 66540618

Cell Line Name	UKBi011-A-2	Batch / Lot Number	M001		
Reprogramming Method	Sendai CytoTune™ 2.0 (POU5F1, SOX2, cMYC, and KLF4)				
Genetic Modification	CRISPR-Cas9				
Passage Number	Passage 29	Cell number / vial	1,8 ×10 ⁶		
Culture Matrix	Matrigel [™] / Geltrex [™]	Culture Medium	mTeSR [™] -1		
O ₂ Concentration	21%	CO ₂ Concentration	5%		
Passaging Method	EDTA	Additional Culture Rho kinase inhibito used at thaw			
Cryopreservation Medium	Cryostor®				
Recommendation for thawing	Recommended thaw into a 50mm cell culture plate. Refer to cell line user protocols for further guidance at www.EBiSC.org				
Additional Comments	Typical recovery after thaw, typical growth to confluency				

Please see https://cells.ebisc.org/ for further information on Quality Control and characterisation 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	
	Inoculation for microbiological growth	Not Detected	Pass	
Sterility	Mycoplasma	Not Detected	Pass	
	Virology (HBV, HCV, HIV1, HIV2) Not Detected		Confirmed Pass by depositor	
Cell Line Identity	STR / Fingerprinting N/A		Allele data recorded and available upon request. Match to donor.	
Viability	Visual Assessment	Growth to confluence post-thaw	Acceptable	
	Continuous visual assessment of iPSC colony morphology	Recorded	Typical PSC colonies with low differentiation levels	
Phenotype	Flow Cytometry	SSEA-4 > 70% + TRA-1-60 > 70% + SSEA-1 < 10% + POU5F1 > 70% +	Pass	
Differentiation Potential	Spontaneous EB differentiation and qPCR for trilineage markers	Up-regulation of germ layer markers	Endoderm Pass Mesoderm : Pass Ectoderm : Pass	



In case of queries, please contact <u>culturecollections.technical@phe.gov.uk</u>. European Collection of Authenticated Cell Cultures (ECACC), Culture Collections, Public Health England, Tel: +44 (0) 1980 612684

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Test	Assay	Acceptance Criteria	Result	
Genomic Stability	Genomic Stability G-Banding		No chromosomal abnormalities detected	
Genetic Modification Sanger sequencing at locus		Match to reported modification	ApoE2/E2 NM_000041.4:c.388C>T NM_000041.3:c.526C>T	

Additional guidance on storage, safety and usage can be found in the **EBISC Technical Information**.

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Approved CoA	Signature_	Miss	Date	28.05.	2021
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