Certificate of Analysis (CoA) for induced Pluripotent Stem Cells



This product is for research only

ECACC Catalogue No: 66540329

Cell Line Name	UNEWi025-B	Batch Number	P001
Donor ID	F324		
Disease Association	Age-related Macular Degeneration	Phenotype of Donor	Unaffected control
Tissue of Origin	Dermal Fibroblast	Sex	Female
Reprogramming Method	Non-integrating Sendai virus (POU5F1, SOX2, KLF4, MYC)		
Passage Number	Passage 12	Cell number / vial	2 x 10 ⁶
Culture Matrix	Geltrex / Matrigel	Culture Medium	mTeSR [™] 1
O ₂ Concentration	20%	CO ₂ Concentration	5%
Passaging Method	EDTA	Additional Culture Information	N/A
Cryopreservation Medium	40% FBS*/ 50% medium / 10% DMSO *Serum of Zone 1 origin		
	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	PubMed ID: 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
	Inoculation for microbiological growth	Not Detected	Pass
Sterility	qPCR for Mycoplasma	Not Detected	Pass
	Virology (HBV, HCV, HIV1, HIV2)	Not Detected	Pass
Cell Line Identity	Short Tandem Repeat analysis using PCR	N/A	Allele data recorded and available upon request. 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 to 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 10-FEB-2017

Test	Assay	Result
Genetic Defect	Sequencing of CFH and HTRA1 genes	Medium risk
Phenotype	Flow Cytometry	TRA-1-60: 100%; NANOG: 96.8%; SSEA-1: 0.5%; SSEA-4: 96.8%
Karyology	CytoSNP	No significant imbalance detected
Cell Line Identity	Short Tandem Repeat analysis using PCR Match to donor	
Clearance of Reprogramming Factors	PCR for Sendai virus	Not detected
Differentiation Potential	Spontaneous differentiation to three germ layers	Endoderm : Detected Mesoderm : Detected Ectoderm : Detected

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 Jone en Spate 14 feb 2017

