## **Certificate of Analysis (CoA) for induced Pluripotent Stem Cells**



This product is for research only

Cell Line Name	BIONi010-C-2	Batch / Lot Number	M001
Reprogramming Method	Episomal vector (POU5F1, SOX2, MYC, LIN28, SHP53 and KLF4)		
Genetic Modification	CRISPR-Cas9		
Passage Number	Passage 39	Cell number / vial	1.2 x 10 <sup>6</sup>
Culture Matrix	Matrigel <sup>™</sup> / Geltrex <sup>™</sup>	Culture Medium	E8-Flex
O <sub>2</sub> Concentration	21%	CO <sub>2</sub> Concentration	5%
Passaging Method	EDTA	Additional Culture Information	None
Cryopreservation Medium	40% FBS*/ 50% medium / 10% DMSO  *Serum of Zone 1 origin		
Recommendation for thawing	Recommended thaw into 2 wells of a 6-well plate or per 10cm <sup>2</sup> 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 <a href="www.ebisc.org/">www.ebisc.org/</a> 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
Sterility	Inoculation for microbiological growth	Not Detected	Pass
	Mycoplasma	Not Detected	Pass
	Virology (HBV, HCV, HIV1, HIV2)	Not Detected	Pass
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
Phenotype	Continuous visual assessment of iPSC colony morphology	Recorded	Typical PSC colonies with low differentiation levels
	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 : Fail Mesoderm : Pass Ectoderm : Pass



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Test	Assay	Acceptance Criteria	Result
Genomic Stability	G-Banding	Sex match to donor. 20 successful karyotypes recorded.	No chromosomal abnormalities detected.
Genetic Modification Sanger sequencing at locus		Match to reported modification	Hemizygous APOE3/E3, see PMID: 32971461.

Additional guidance on storage, safety and usage can be found in the <u>EBISC Technical Information</u> .				
Approved CoA	Signature	Date		

