Certificate of Analysis (CoA) for induced Pluripotent Stem Cells



This product is for research only

| Cell Line Name BIONi010-C-15 Batch / Lot Number |
|---|
|---|

| Reprogramming Method Genetic Modification | Non-integrating episomal vector (POU5F1, SOX2, MYC, Lin28, shP53 and KLF4) TALEN Dox inducible NGN2-GFP expression | | |
|--|--|-----------------------------------|--------------------------------------|
| Passage Number | 32 | Cell number / vial | 1,5x10E6 |
| Culture Matrix | Matrigel™ | Culture Medium | Essential 8 [™] |
| O ₂ Concentration | 18% | CO ₂ Concentration | 5% |
| Passaging Method | EDTA | Additional Culture Information | Rho kinase inhibitor used at thaw |
| Cryopreservation Medium | Cryostor | | |
| Recommendation for thawing | Recommended thaw into 60mm plates 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 | 85% match to donor Sex match to donor | 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 |



In case of queries, please get in touch via Contact@EBiSC.org

BIONi010-C-15.M002.CoA.v2

Certificate of Analysis (CoA) for induced Pluripotent Stem Cells



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Cell Line Name BIONi010-C-15 Batch / Lot Number M002

| Test | Assay | Acceptance Criteria | Result |
|----------------------|---|--|---|
| | Flow Cytometry | SSEA-4 > 70% positive TRA-1-60 > 70% positive SSEA-1 < 10% positive POU5F1 > 70% positive | Pass |
| Genomic Stability | G-Banding (10 -20 successful karyotypes recorded) | Sex match to donor. | 46, XY |
| Genetic Modification | DOX induction, qPCR and microscopy | Match to reported modification | Rapid neuronal differentiation and inducible GFP expression |

Differentiation potential not performed for batch M002.

Please refer to historical data from the initial batch P001:

| Differentiation Potential | Trilineage differentiation and qPCR for trilineage markers | Up-regulation of germ layer markers | Endoderm: Pass Mesoderm: Pass Ectoderm: Pass |
|------------------------------|--|--|--|
|------------------------------|--|--|--|

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

Approved CoA

Signature Pfwau Date 14.06-2024



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BIONi010-C-15.M002.CoA.v2