

Cell Line Details

| HEP-002 | |
|--|--|
| iPSC-derived Human Hepatocytes: Patient-derived Compound Heterozygous Glycogen Storage Disease Type 1 Alpha; G222R/Q347X | |
| XXXXXX | |
| Store at less than -130°C | |
| XXXXXX | |
| | |

Cell Quality Controls

| Test | Method | Specification | Result |
|---|---------------------------|-------------------------|--------------|
| Virus test for original iPSC clone (HIV1, HIV2, Hepatitis A, HBV, HCV, HTLV-1, HTLV-2) | PCR | Not detected | Pass |
| Post thaw viability | Automated cell counter | ≥ 70% viable | Pass |
| Viable cells per vial | Automated cell counter | ≥ 5.0 x 10 ⁶ | Pass |
| Cell morphology | Visual check | N/A | Pass (Fig.1) |
| Key hepatocyte maturity markers (<i>ALB, A1AT,</i> <i>HNF4</i> a) | qPCR | Present | Pass (Fig.2) |
| Disease confirmation | Sanger sequencing | Mutation present | Pass (Fig.3) |

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CERTIFICATE OF ANALYSIS

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Appendix

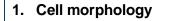
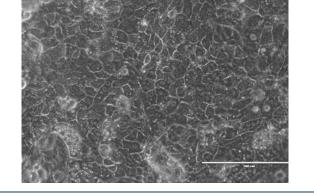


Figure 1. Morphology of cryopreserved hepatocyte-like cells, 14 days post-thaw. Brightfield picture, magnification: 200x.

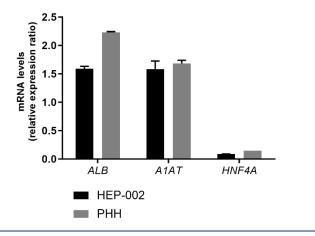


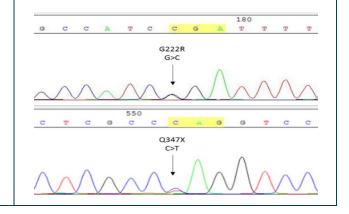
2. Detection of hepatocyte maturity markers via qPCR

Figure 2. mRNA expression of the key hepatocyte maturity markers *ALB* (Albumin), *A1AT* (Alpha-1 Antitrypsin) and *HNF4A* (Hepatocyte Nuclear Factor-4) in cryopreserved hepatocyte-like cells (black bars) and primary human hepatocytes (PHH, grey bars), 14 days post-thaw. mRNA data are normalized to endogenous *PPIA* expression, and are presented as mean±SD of n=3 technical replicates.

3. Disease confirmation via Sanger sequencing

Figure 3. Sanger sequencing showing heterozygous G222R mutation (GGA > CGA) and heterozygous Q347X mutation (CAG > TAG) in the *G6PC* gene. The codon change is highlighted in yellow.





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Checked by,

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