

Sansure Ultra HBV Manual Extraction and Amplification SOP

Notes before start:

□ Go through Sansure User's manual before operation and don't mix different batches of reagents.

1. Sample preparation:

- □ Collect more than 1.2mL sample so that it can repeat test for 3 times;
- □ Sample should be transported in a sealed frozen pitcher with ice or in a sealed foam box with ice;
- □ Sample can be stored at 4°C for 4 days, or stored at -20°C for 3 months, or stored below -70°C for long-term storage. Sample repeated freezing and thawing should be avoided;
- \Box Recover the sample to room temperature, <u>vortex for 5~10s and centrifuge at 3000rpm for 5s</u>.

2. Consumable preparation:

□ Magnetic separator, 1.5mL EP tube, 0.2mL PCR 8-tube strip.

3. Reagent preparation:

 \Box 3.1 Extraction Kit: S10012E-48, 48T/kit, stored at 2~8°C;

S10012E Extraction Solution 1
S10012E Extraction Solution 2
S10012E Extraction Solution 3
S10012E Extraction Solution 4
Elution Buffer

 \Box 3.2 Amplification Kit: S3118E-48 HBV Ultra, 48T/kit, stored at -20°C;

HBV PCR Mix	HBV Negative Control		
HBV Enzyme Mix	HBV Quantitative Reference A		
HBV Internal Control	HBV Quantitative Reference B		
HBV Strong Positive Control	HBV Quantitative Reference C		
HBV Weak Positive Control	HBV Quantitative Reference D		

□ 3.3 Sansure Kits Melting:

□ Take all components from extraction kit and equilibrate to room temperature until no ice inside, then gently shake and reverse them upside down several times to mix throughly, especially the **Extraction Solution 2** (Observe the Extraction Solution 2 tube bottom, make sure no magnetic beads accumulation).

 \Box Take all components from amplification kit and equilibrate to room temperature until no ice inside, then vortex them for 5~10s and centrifuge at 3000rpm for 5s. After centrifuge, no more reverse.

 \Box 3.4 Reagents allocation:

Prepare (n+1) tests, where n is the total sample number including controls and references.

□ Reverse the Extraction Solution 2 tube upside down several times to mix throughly again. Observe the tube bottom, make sure no magnetic beads accumulation.

 \Box Take an EP tube and prepare Extraction Solution 2 Mix: 100µL Extraction Solution 2 + 1µL HBV Internal Control / test for (n+1) tests, when adding the IC into the extraction solution 2, pipette several times to mix;

 \Box Cap and mark the tube, then <u>vortex for 5~10s and centrifuge at 3000rpm for 5s</u>.



4. Extraction:

- \Box Nucleic acid lysis:
 - \Box Take (n) 1.5ml centrifuge tubes, cap and mark them according to the samples;
 - □ Open the tubes lid, pipette 300µL Extraction Solution 1 + 400µL Sample into each tube;
 - □ Cap the tubes lid, vortex them for 5~10s and centrifuge at 3000rpm for 5s;
 - \square Open the tubes lid, pipette 100 μL Extraction Solution 2 Mix into centrifuge tubes;
 - \Box Cap the tubes lid, <u>vortex them for 5~10s</u>;
 - \square Place the tubes at room temperature for 10 mins.
- \Box Nucleic acid binding to the magnetic beads:
 - □ <u>Centrifuge the above (n) centrifuge tubes at 3000rpm for 5s;</u>
 - □ Put the centrifuge tubes onto the magnetic separator for 5 minutes, until the magnetic beads attached to the tube wall;
 - □ Open the tubes lid, gently aspirate the waste reagents out of the tubes, without touching the magnetic beads.

Slowly aspirate with <u>1000µL tips;</u>

- \Box Nucleic acid purified:
 - \Box Keep the above (n) centrifuge tubes on the separator, open the tubes lid, pipette 600µL Extraction Solution 3 + 200µL Extraction Solution 4 into each tube;
 - \Box Cap the tubes lid, take out them from the separator, then <u>vortex for 5~10s and centrifuge at</u> <u>3000rpm for 5s;</u>
 - □ Put the tubes onto the magnetic separator again for 3 mins, until the magnetic beads attached to the tube wall and the waste reagents being separated into two layers.

[If not two layers, vortex fully and centrifuge again, then repeat this step.]

- □ Open the tubes lid, gently aspirate the waste reagents out of the tubes, without touching the magnetic beads.
 - □ Firstly, slowly aspirate with <u>1000µL tips;</u>
 - □ Secondly, centrifuge at 3000rpm for 5s and back to the separator for 3mins, then aspirate with 200μ L and 10μ L tips for several times until no liquid inside.

[The waste liquid will affect the amplification process, so this step is essential!]

- $\hfill\square$ Nucleic acid eluted:
 - \Box Take the above (n) tubes out from the magnetic separator;
 - \Box Open the tubes lid, pipette 30µL Elution Buffer into each tube to wash down the magnetic beads to the tube bottom, and vortex for 5~10s to mix;
 - □ Cap the tubes lid and place them at room temperature for 10mins;
 - □ While waiting, prepare **PCR-Master-Mix**:
 - □ 27.5µL HBV PCR Mix + 2.5µL HBV Enzyme Mix / test for (n+1) tests;
 - □ Cap and mark the tube, vortex for 5~10s and centrifuge at 3000rpm for 5s;
 - \Box <u>Centrifuge the samples at 3000rpm for 5s</u>, put the tubes back to magnetic separator again for 3 minutes, until the magnetic beads attached to the tube wall.



 \Box Amplification preparation:

- □ Take certain number of PCR 8-tube strips according to the samples' amplification layout;
- D Pipette 30µL PCR-Master Mix into to each sample PCR well, mark them according to the samples;
- \square Pipette 20 μL samples eluted nucleic acid into each sample PCR well according to the marks;
- Cap the PCR tubes lid tightly, <u>vortex for 5~10s and centrifuge at 3000rpm for 5s</u>.
 [If no centrifuge for PCR tube, pipette several times when adding the samples.]
 [Make sure there are no bubbles in the tubes.]
- 5. Amplification:
- \Box Amplification channels setting:
 - □ FAM (Quencher: None): HBV-DNA;
 - □ HEX/VIC (Quencher: None): HBV Internal Control;
 - □ ROX: passive reference. [Only for instruments with ROX correction system]
- \Box Amplification volume: 50µL;
- \Box Cycle parameters:

	Steps	Temperature	Time	Cycles
1	Uracil-DNA Glycosylase reaction	50°C	2min	1
2	Uracil-DNA Glycosylase inactivation	94°C	5min	1
3	Denaturation	94°C	15sec	- 45
	Annealing, extension and fluorescence collection	57°C*	30sec	- 43

*Click on 'signal acquisition' in the 3.2 step.