

Charm-Pure™ Medium-Scale Genomic DNA Purification from Saliva

Store at room temperature, store Proteinase K and RNase A at 4 °C Product Contents and Storage

The components included with the Charm-Pure Medium-Scale gDNA Purification from Saliva are listed below. Upon receipt store all components at room temperature with the exception of Proteinase K and RNase A, which need to be stored at 4 °C. Sufficient reagents are included to perform 50 or 250 genomic DNA extractions.

| Product Catalogue | QC-276-M | QC-276-L |
|---------------------------------------|----------|----------|
| Purification Scale | 50 | 250 |
| Binding Tube (BT3) | 50 | 250 |
| Spin-Column with Collection Tube(CC1) | 50 | 250 |
| Lysis Buffer (PKLB) | 5.5 ml | 27.5 ml |
| Binding Buffer (KB6) | 10.0 ml | 50.0 ml |
| Binding Buffer (GB5) | 11.0 ml | 55.0 ml |
| Washing Buffer I (WB1) | 5.5 ml | 27.5 ml |
| Washing Buffer II (WB2) | 11.0 ml | 55.0 ml |
| Elution Buffer (EB2) | 24 ml | 120 ml |
| Proteinase K (PK2) (20 mg/ml) | 1.1 ml | 5.5 ml |
| RNase A (RA2) (20 mg/ml) | 1.1 ml | 5.5 ml |

Product Description

The Charm-Pure TM Medium-Scale Genomic DNA Purification from Saliva kit is designed and optimized for easy and fast isolation of high-quality gDNA from fresh saliva, frozen saliva or saliva sample stored in the Charm-Protect Saliva Collection and Storage Vial or preservation solutions from other companies. Based on Charm Biotech developed Solid Surface Reversible Binding (SSRB) technology and special silica membrane technology, micro-tubes coated with proprietary turbo-binders and spin-column with modified silica membranes are used to selectively capture and efficiently bind genomic DNA from saliva cell lysates. High-yield and high-quality DNA can be easily obtained from 1 ML saliva mixtures. The purified gDNA can be eluted in Elution Buffer or water. The purified ready-to-use gDNA is ideal for downstream application such as restriction digestion, primer extension, SNP analysis, PCR, sequencing, whole genome amplification (WGA) and DNA methylation analysis.

Additional Materials Needed

- 100 % ethanol, Isopropanol
- 1.5 ml or 2.0 ml microtubes
- Bench-top centrifuge capable of spinning 1.5-2.0 ml microtubes, Vortex or Shaker, Incubator (air or water) or hot blocker

General Precautions

- This kit is for research use only. All due care and attention should be exercised in the handling of the kits.
- Wear a laboratory coat, disposable gloves, and eye protection when handling reagents and tubes. Avoid ingestion and
 inhalation of reagents. Avoid skin contact with reagents in the kit. In case of contact, wash thoroughly with water. See
 Material Safety Data Sheets (MSDS) for emergency procedures in case of accidental contact or ingestion. All MSDS
 information is available upon request.
- Always use proper aseptic techniques to avoid nuclease contamination when working with nucleic acids and use only sterile, new pipette tips to prevent cross contamination.

Preparation before Starting

• Prepare fresh working Binding Buffer prior to performing gDNA isolation from saliva lysates based on the number of samples processed. To make working Binding Buffer (KB6), mix 100 µl of Binding Buffer (KB6) with 400 µl 100 % Isopropanol and mix well. Prepare a master Binding Buffer solution based on (1) the number of samples processed and (2) any anticipated loss, generally 10 %, during dispensing. For working Binding Buffer (KB6), dispense 100 µl of

Binding Buffer containing Isopropanol per $100 \mu l$ cell lysates (1:1 ratio). Discard the unused Binding Buffer at the end of the day.

- For QC-276-M kit, add 22 ml of 100 % ethanol to Washing Buffer I (WB1) and mix well. For QC-276-L kit, add 110 ml of 100 % ethanol to Washing Buffer I (WB1) and mix well. (Mark bottle that ethanol has been added!) Store at room temperature and use WB1 containing ethanol within six (6) months.
- For QC-276-M kit, add 44 ml of 100 % ethanol to Washing Buffer II (WB2) and mix well. For QC-276-Lkit, add 220 ml of 100 % ethanol to Washing Buffer II (WB2) and mix well. (Mark bottle that ethanol has been added!) Store at room temperature and use WB2 containing ethanol within six (6) months.

Experimental Procedure

Preparing Lysates

The protocols listed below are for purification of gDNA from 1ml of saliva preservation mixture. Adjust volume accordingly for actual volume of saliva mixture.

Note: If the saliva samples are stored in the Charm-Protect Saliva Preservation Solution, please follow the procedure below to prepare the saliva lysates. (If you need to isolate gDNA from saliva samples without using Charm-Protect Saliva Preservation Solution, please check "Procedure for Non-Charm Protected saliva samples").

Procedure for Non-Charm Protected saliva samples:

- 1. Add 20 µl RNase A (RA2) into a 1.5 ml or 2.0 ml microtube.
- 2. Mix the saliva sample mixture stored in the Charm-Protect Saliva Preservation Solution in the Saliva Collection Vial by inverting and shaking 7-8 times. Let the vial or tube sit for 3-5 minutes.
- 3. Transfer 1 ml saliva preservation mixture from the Saliva Collection Vial into the microtube preloaded with RNase A (RA2) and mix well by pipetting the solution up and down 8-9 times or by inverting the microtube 7-8 times after closing the cap. (Note: Don't collect the particulate settlement at the bottom of the collection vial when aspirate saliva mixture.)
- 4. Add 20 µl Proteinase K (PK2) into the tube and mix well by inverting the tube 15-18 times after closing the cap..
- 5. Incubate the microtube at 62 °C in an air incubator or a hot-blocker for 45 minutes with occasional mixing by tapping or inverting the tube. (Optional: Incubate the sample overnight at 52 °C if prefer)
- 6. Centrifuge the microtube at maximum speed (about 13000 rpm or 16000 g) for 4 minutes to remove any particulate materials, such as food particles, undigested dead cells and fibers debris.

Procedure for Non-Charm Protected saliva samples:

If your saliva samples are fresh saliva, frozen saliva or saliva mixed with saliva preservation solution from other companies, please prepare saliva lysates as follows:

- 1. Add 20 µl RNase A (RA2) into a 1.5 ml or 2.0 ml microtube.
- 2. Add 100 µl Lysis Buffer (PKLB) into the tube (the volume ratio of PKLB vs. saliva sample is 1 to 9).
- 3. Mix the saliva sample by inverting the saliva container 7-8 times and then let the container sit for 3-5 minutes.
- 4. Transfer 900 μl saliva or saliva mixture into the microtube preloaded with RNase A (RA2) and PKLB, mix well by pipetting the solution up and down 8-9 times or by inverting the microtube 9-10 times after closing the cap. (Note: Don't collect the particulate settlement at the bottom of the container when aspirate saliva mixture.)
- 5. Add 20 μl Proteinase K (PK2) into the tube and mix well by inverting the tube 15 -18 times.
- 6. Incubate the microtube at 62 °C in an air incubator or a hot-blocker for 45 minutes with occasional mixing by tapping or inverting the tube. (Optional: Incubate the sample overnight at 52 °C if prefer)
- 7. Centrifuge the microtube at maximum speed (about 13000 rpm or 16000 g) for 4 minutes to remove any particulate materials, such as food particles, undigested dead cells and fibers debris.

Binding Products

- 1. Transfer 900 μl clarified saliva lysate into a Binding Tube (BT3) without collecting the debris pellet at the bottom of the microtube (Note: To avoid the debris pellet, you may leave about 150-250 μl solution in the micro-tube when doing aspiration).
- 2. Add 900 μl working Binding Buffer (KB6) containing isopropanol to the Binding Tube (BT3). Mix well with the saliva lysate by pipetting up and down the solution 10 times or by inverting the microtube 15-18 times after closing the cap to obtain a homogenous solution. (Note: It is crucial to mix working Binding Buffer with saliva lysate thoroughly.)
- 3. Centrifuge the Binding Tube (BT3) at maximum speed (about 13000 rpm or 16000 g) at room temperature for 4 minutes to bind the DNA. (Note: During centrifugation, always position micro-tube hinge pointed outward from the center of rotation. Majority of DNA will collect at the bottom along the hinge side of the microtube.)
- 4. Open the cap, remove the solution from the Binding Tube by flipping the Binding Tube over a waste container and shaking briskly. Let the tube stand upside down on a piece of clean absorbent paper for 1-5 minutes to drain as much liquid as possible.

- 5. Add 300 μl Elution Buffer (EB2) to the Binding Tube. Vortex for 1 minute and incubate the tube at 62 °C for 10-15 minute to dissolve the DNA pellet thoroughly.
- 6. Add 180 µl Binding Buffer (GB5) to the Binding Tube and mix well by pipetting the solution 8-9 times.
- 7. Add 180 µl ethanol to the mixture above and mix well by pipetting the solution 8-9 times.
- 8. Transfer all solution mixture (about 660μ l) from the step above to a Spin-Column with a Collection Tube (CC1). Centrifuge the Spin-Column at 8000 rpm (RCF = 5800 g) for one minute to allow all liquid pass through the membrane at room temperature.
- 9. Discard the filtrate in the Collection Tube and reinsert the Spin-Column back into the Collection Tube.

Washing Products

- 1. Add 500 μl of Wash Buffer I (WB1) with ethanol to the Column.
- 2. Centrifuge the Spin-Column at maximum speed (≥ 13000 rpm or 16000 g) for 1 minute at room temperature. Discard the filtrate
- 3. Add 500 µl of Wash Buffer II (WB2) with ethanol to the Column.
- Centrifuge the Spin-Column at maximum speed (≥ 13000 rpm or 16000 g) for 1 minute at room temperature. Discard the filtrate.
- 5. Repeat step 3 and 4 above once for a total of two washes with Wash Buffer II (WB2).
- 6. Centrifuge the Spin-Column at maximum speed for one minute to remove any residual wash buffer with ethanol. Discard collection tube.
- 7. Insert the Spin-Column into a new clean microtube (self-provide). (Note: When transferring the Spin-Column, be careful of preventing the residual wash buffer contacting the tip of the column)

Eluting Genomic DNA

- 1. Warm up the Elution Buffer (EB2) in a 1.5 ml microtube with a 62 °C hot block or water-bath.
- 2. Add 75 µl Elution Buffer (EB2) onto the center of the Spin-Column, incubate the spin-column with collection microtube at 62 °C in an air incubator or a hot-blocker for 5 minutes.
- 3. Centrifuge the column at maximum speed for 1 minute at room temperature. The microtube contains the filtrate which contains purified gDNA. (Note: You may perform a second elution with another 50 µl of Elution Buffer (EB2) if desired. This can increase the total gDNA yield by 15-30 %, but the final concentration of isolated DNA in the eluted sample is reduced because of the increased volume).
- 4. The eluted gDNA may be used immediately in downstream applications. Alternatively, the eluted gDNA may be stored in the microtube at 4 °C for short-term storage or -20 °C for long-term storage.

Electrophoresis and Downstream Application

Purified gDNA can be examined by agarose gel electrophoresis. Yield can be measured with a spectrophotometer or fluorescent DNA assays. The yield of purified gDNA varied between 4-25 μg from different saliva samples, $A_{260/280} \geqslant 1.7$ (the majority is between 1.8-2.0). Purified gDNA can be used immediately for downstream application, such as restriction enzyme digestion, primer extension, SNP analysis, PCR, STR analysis, DNA sequencing, whole genome amplification (WGA) and other molecular manipulations.

Troubleshooting

| Problem | Cause | Solution |
|----------------|--|---|
| Low DNA yield | Poor quality of starting material | Be sure the protocol for sample collection and preservation recommended by manuals is followed correctly. |
| | Incomplete lysis | If incomplete lysis is observed, extend the incubation time at 62 °C for an addition of 15 – 30 minutes. |
| | Variable DNA amount in different individual saliva samples | Increase the starting saliva sample volume for DNA isolation. Please call us for exact procedure if you want to process whole 4 ml saliva mixtures. |
| No PCR product | Missing Component in the PCR mixture | Be sure to add all components. Check positive control and negative control for PCR reaction. |

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Charm-Pure[™]精纯中量唾液 DNA 提纯试剂盒

室温运输和保存,Proteinase K和 RNase A保存在 4°C。

试剂盒组成及贮存

Charm-Pure[™] 精纯中量唾液DNA提纯试剂盒可从1 ml 唾液中提纯高质量的gDNA,其组成如下表所示,除了 Proteinase K 和 RNase A 需要保存在4 °C 之外,其余组分室温保存即可。可进行 50或者250 次唾液gDNA 提纯。

| Product Catalogue | QC-276-M | QC-276-L |
|---------------------------------------|----------|----------|
| Purification Scale | 50 次 | 250 次 |
| Binding Tube (BT3) | 50 个 | 250 个 |
| Spin-Column with Collection Tube(CC1) | 50 套 | 250 套 |
| Lysis Buffer (PKLB) | 5.5 ml | 27.5 ml |
| Binding Buffer (KB6) | 10.0 ml | 50.0 ml |
| Binding Buffer (GB5) | 11.0 ml | 55.0 ml |
| Washing Buffer I (WB1) | 5.5 ml | 27.5 ml |
| Washing Buffer II (WB2) | 11.0 ml | 55.0 ml |
| Elution Buffer (EB2) | 24 ml | 120 ml |
| Proteinase K (PK2) (20 mg/ml) | 1.1 ml | 5.5 ml |
| RNase A (RA2) (20 mg/ml) | 1.1 ml | 5.5 ml |

产品介绍

Charm-Pure ™ 精纯中量唾液 DNA 提纯试剂盒专为从新鲜唾液、冻存唾液、Charm-Protect™ 或其他公司唾液保存液中分离提纯高质量的唾液 gDNA 而设计,操作步骤简单迅速。昌美生物将固体表面可逆结合技术(SSRB)和传统的核酸硅膜技术相结合,采用特别的结合管和改进的硅膜来选择性地吸附核酸,能有效处理 1 ml 的唾液混合液,高效地提纯回收高纯度的唾液 gDNA。产量高,质量好!提纯的 gDNA 可直接用于下游各种实验,如限制性酶切反应、SNP 分析、PCR、序列分析、NGS、全基因组扩增(WGA)和 DNA 甲基化分析等。

亮点

操作简单: 可处理 1 ml 唾液样品,避免多次处理小量样品,操作步骤简单,不需要特别的仪器。

<u>质量可靠,回收率高</u>:采用独特的固体表面可逆结合技术和改进的硅膜核酸结合技术,能最大程度的回收唾液样品的 gDNA 不易被污染,质量高,重复性好。

所需的额外材料

- 无水乙醇、异丙醇
- 1.5 ml 或 2.0 ml 离心管
- 可离 1.5 ml 或 2.0 ml 离心管的离心机、涡旋振荡器、水浴锅或恒温金属浴。

常规防范

- 该试剂盒仅用于实验。
- 实验操作时需穿实验服,戴一次性手套。避免摄取和吸入试剂,避免与试剂盒中试剂进行直接接触,一旦接

触,用水彻底冲洗。

• 提纯核酸时,使用无菌新吸头避免污染。

实验前准备

- 在进行唾液提纯 gDNA 之前,需要根据提纯的样品数目制备新鲜的 Binding Buffer(KB6)工作液: 将 Binding Buffer(KB6)与异丙醇以 1: 4 体积比混匀,如配制 500 μl Binding Buffer(KB6)工作液需 将 100 μl Binding Buffer(KB6)与 400 μl 异丙醇混合均匀得到。需要制备 Binding Buffer(KB6)工作液的体积根据: (1)待提纯的样品数目; (2)预期的损失,在悬浮过程中,一般会有 10%的损失。每 100 μl 细胞裂解液需加 100 μl 含异丙醇的 Binding Buffer(KB6)工作液(1: 1 体积比)。剩余的 Binding Buffer(KB6)工作液当天予以丢弃。
- 制备 Washing Buffer I(WB1)工作液:如果是 QC-276-M 试剂盒,加 22 ml 无水乙醇到 Washing Buffer I(WB1)中,充分混匀。如果是 QC-276-L 试剂盒,加 110 ml 无水乙醇到 Washing Buffer I(WB1)中,充分混匀。在瓶上做好标记表示已加乙醇!室温保存,加乙醇的 WB1 在六个月之内使用。
- 制备 Washing Buffer II(WB2)工作液:如果是 QC-276-M 试剂盒,加 44 ml 无水乙醇到 Washing Buffer II(WB2)中,充分混匀。如果是 QC-276-L 试剂盒,加 220 ml 无水乙醇到 Washing Buffer II(WB2)中,充分混匀。在瓶上做好标记表示已加乙醇!室温保存,加乙醇的 WB2 在六个月之内使用。
- 该试剂盒可从 1 ml 唾液保存液中提纯 gDNA (0.5 ml 唾液 + 0.5 ml 唾液保存液) 或冻存的唾液中提纯 gDNA。如果需要处理 4 ml 保存的唾液 (2 ml 唾液 + 2 ml 唾液保存液) 或冻存的唾液, 请使用"昌美精纯大量唾液 DNA 提纯试剂盒"。在收集唾液时,为了避免 gDNA 降解,请遵守标准的 Charm-Protect™ 唾液 DNA 收集和保存程序。

操作指南

制备唾液细胞裂解液

注意:如果唾液样本是昌美生物 Charm-Protect™ 唾液保存液保存的唾液,请使用以下步骤制备唾液细胞裂解液(如果您需要提纯其它唾液样本,请见"非昌美生物 Charm-Protect™ 唾液保存液保存的唾液操作步骤"。)

昌美生物 Charm-Protect ™ 唾液保存液保存的唾液操作步骤:

- 1. 取 20 μl RNase A (RA2) 到一个 1.5 ml 或 2.0 ml 离心管中。
- 2. 将储存在 Charm-Protect 唾液收集瓶中的唾液样品与唾液保存液颠倒 7-8 次后静置 3-5 分钟,让颗粒杂物沉 淀到瓶底。
- 3. 从唾液收集瓶中吸取 1 ml 唾液与保存液的混合物到预先加了 RNase A (RA2)的离心管中,吸头吹打 8-9 次,或将离心管盖子盖上后颠倒 7-8 次,混合均匀。当吸取唾液混合物时,注意不要吸到收集瓶底颗粒状沉淀杂物。
- 4. 向管中加 20 μl Proteinase K (PK2), 将离心管盖紧后颠倒离心管 15-18 次, 充分混匀。
- 5. 将离心管置于 62 ℃ 水浴锅或金属浴中温育 45 分钟,进行 PK2 裂解,期间颠倒混合离心管。如有需要,也可将样品在 52 ℃ 温育过夜。(注意不是 62 °C 过夜)。
- 6. 将离心管以最大转速(13000 rpm 或 16000 g)离心 4 分钟来去除颗粒物质,如食物颗粒、未经消化的死细胞和纤维碎片。

非昌美生物 Charm-Protect M 唾液保存液保存的唾液操作步骤:

如果您的样本是新鲜唾液、冰冻的唾液,或者是使用其他公司的唾液保存液保存的唾液,请使用以下步骤制备唾液 细胞裂解液:

- 1. 取 20 μl RNase A (RA2) 到一个 1.5 ml 或 2.0 ml 离心管中。
- 2. 向管中加入 100 μl Lysis Buffer (PKLB) (PKLB 和唾液样品的体积比例是 1:9)。
- 3. 将唾液样品颠倒 7-8 次后静置 3-5 分钟, 让颗粒杂物沉淀。
- 4. 从样本中吸取 900 μl 唾液或者唾液混合物,加到预先加了 RNase A (RA2) 和 PKLB 的离心管中,吸头吹打 8-9 次,或将离心管盖子盖上后颠倒 9-10 次,充分混匀。(当吸取唾液混合物时,注意不要吸到底部颗粒状沉淀杂物。)
- 5. 向管中加 20 μl Proteinase K (PK2),将离心管盖紧后颠倒离心管 15-18 次,充分和唾液混合均匀。
- 6. 将离心管置于 62°C 水浴锅或金属浴中温育 45 分钟,进行 PK2 裂解,期间颠倒混合离心管。如有需要,也可将样品放在 52°C 温育过夜(注意不是 62°C 过夜)。
- 7. 将离心管以最大转速(13000 rpm 或 16000 g)离心 4 分钟来去除颗粒物质,如食物颗粒、未经消化的死细胞和纤维碎片。

结合 DNA 产物

- 1. 转移 900 μl 澄清的唾液裂解液到一个结合管 (BT3) 中,注意不要吸到离心管底的碎片沉淀。
- 2. 加 900 μ l 含有异丙醇的 Binding Buffer(KB6)工作液到结合管(BT3)中,将结合管(BT3)盖紧后颠倒结合管 15-18 次,充分混合均匀(注意:这一步非常重要)。
- 3. 将结合管(BT3)在室温下以最大转速(13000 rpm 或 16000 g)离心 4 分钟来结合 DNA 到管壁(注意:在离心过程中,离心管盖朝向转子方向放置,离心后 DNA 大部分聚集在背对管盖的管底一侧。)
- 4. 直接将结合管中的液体倾倒入废液收集杯中,然后将结合管倒放在干净的吸水纸上面1至5分钟。
- 5. 加 300 μl Elution Buffer (EB2) 溶液, 涡旋 1 分钟, 62 °C 温育 10-15 分钟, 使沉淀完全溶解。
- 6. 再加 180 μl Binding Buffer (GB5), 用移液器来回吹打溶液 8-9 次,混合均匀。
- 7. 向上述混合物中加 180 µl 乙醇,用移液器来回吹打溶液 8-9 次,混合均匀。
- 8. 将所得的唾液混合物(约 660 μ l)转移到配有收集管(Collection Tube)的离心柱(Spin-Column)中,室温下 8000 rpm(RCF = 5800 g)离心 1 分钟,让所有液体通过离心柱。
- 9. 取出 Spin-Column,将收集管中的滤液丢弃,将 Spin-Column 重新插回到收集管中。

清洗 DNA 产物

- 1. 向离心柱(Spin-Column)中加 500 μl 含有乙醇的 Wash Buffer I(WB1)。
- 将带有收集管的离心柱(Spin-Column)以最大转速(≥13000 rpm 或 16000 g)室温离心 1 分钟,倒掉收集管中滤液。
- 3. 向 Spin-Column 中加 500 μl 含有乙醇的 Wash Buffer II(WB2)。
- 4. 将 Spin-Column 以最大转速 (≥13000 rpm 或 16000 g) 室温离心 1 分钟, 倒掉收集管中滤液。
- 5. 重复步骤 3 和 4, 即用 Wash Buffer II (WB2) 洗两次。

6. 以最大转速将带有收集管的 Spin-Column 离心 1 分钟彻底去除含有乙醇的 WB2。丢掉收集管。将 Spin-Column 放入一个干净的 1.5 ml 离心管中。(在转移 Spin-Column 时,注意避免残留的 Wash buffer 接触到 Spin-Column 的尖端。)

gDNA 的洗脱

- 1. 根据需要洗脱的样品数量,取一定体积的 Elution Buffer (EB2) 到 一个 1.5 ml 离心管中,置于 62 °C 金属浴或水浴中加热。
- 2. 加 75 μl 预热的 Elution Buffer (EB2) 到 Spin-Column 的膜中央, 62°C 温育 5 分钟。
- 3. 室温下最大速度离心 1 分钟, 离心管中的滤液即为纯化的 gDNA。注意:如果有需要,可另加 50 μl Elution Buffer (EB2)进行再次洗脱。这一步可增加产量 15-30 %,但提纯的 gDNA 因为体积增大,终浓度会降低。
- 4. 洗脱到离心管中的 gDNA 可直接用于下游反应,或者置于 4°C 短期保存或 20°C 长期保存。

电泳及下游的应用

提纯的 gDNA 可用琼脂糖凝胶电泳进行质检,也可用分光光度计或荧光标记法测定产量。根据不同的唾液来源, gDNA 产量在 4-25 μ g 之间, $A_{260/280}$ 比值 \geq 1.7,大部分值在 1.8-2.0 之间。纯化的 gDNA 可直接用于下游反应,如限制 性酶切反应、SNP 分析、PCR、STR 分析、DNA 序列分析、全基因组扩增(WGA)及其它分子实验操作。

常见问题及解决方案

| 问题 | 原因 | 解决方案 |
|----------|--|---------------------------------|
| DNA 产量低 | 起始材料质量差。 | 确保正确遵守手册推荐的样品收集和保存程序。 |
| | 裂解不彻底 | 将 Proteinase K 消化的时间延长 15-30 分钟 |
| | 样品本身 DNA 量不同(唾液 DNA 含量,因不同个体体质而异,如小孩唾液 DNA 量比老年人少) | 增加要提纯唾液样品起始量 |
| PCR 没有产物 | PCR 反应混合物中漏加组分 | 确保加了所有的组分。核对 PCR 反应的阳性对照和阴性对照。 |

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