
Auto-Mag[®] HMW Size Select

Version 2.1

Magnetic Beads-Based Reagent for short DNA fragment depletion and size selection of High Molecular Weight (HMW) DNA sample for Long-Read Sequencing (LRS)

Catalog Number: S005-01, S005-02

Contents

• Disclaimers and Safety Information.....	1
• Product Introduction.....	2
• Kit Contents and Storage.....	2
• Preparation of Reagents.....	2
• Additional Information.....	3
• Auto-Mag [®] MHW Size Select Protocols.....	4
Protocol for Short DNA Fragments Depletion and HMW DNA Size Selection	5
Supplemental Protocol for Recovery Unbinding Short DNA Fragments.....	7
Customized Protocol and Programmed Procedure for Automation Purification.....	8
• Troubleshooting Guide.....	8
• Ordering Information.....	9

Disclaimers and Safety Information

This kit is designed for research use only. All biological samples are considered potentially infectious. When working with chemicals, always wear a suitable lab coat, disposable gloves, and protective goggles. For more information, please consult the appropriate material safety data sheets (MSDSs). MSDS can be downloaded from the “Product Documents” tab when viewing the product kit. Download MSDS at www.amdbiotech.com. Information in this document is subject to change without notice.

Product Introduction

The Auto-Mag® HMW Size Select (Mega Select) is an innovative and truly practical reagent that utilizes advanced nanomagnetic bead technology to efficiently sort, purify, and recover high molecular weight (HMW) DNA fragments. Its user-friendly operation allows precise adjustment of the reagent-to-sample volume ratio, enabling the rapid and effective removal of short DNA fragments from HMW genomic DNA samples. This process selectively retains large DNA fragments—such as those exceeding 5 kb, 10 kb, 20 kb, or 30 kb, meeting the stringent requirements of third generation long-read sequencing platforms like PacBio SMRT and Oxford Nanopore.

Third-generation sequencing technologies demand high-quality, long-length DNA templates to ensure accurate and reliable results. However, genomic DNA samples are often contaminated with short fragments due to factors such as suboptimal sample quality or partial degradation during extraction or storage. These short fragments can compromise sequencing accuracy and performance, making their removal critical. Auto-Mag® HMW Size Select effectively addresses this issue by selective recovery of high-quality HMW DNA while removing unwanted short fragments. By enriching large DNA fragments, it enhances sample quality and improves sequencing performance, yielding longer read lengths and greater accuracy.

Auto-Mag® HMW Size Select offers exceptional adaptability, delivering optimal performance with DNA sample concentrations ranging from 10 to 150 ng/μl. Its advanced technology eliminates the need for centrifugation or filtration, simplifying workflows and enhancing efficiency. Designed with both flexibility and convenience in mind, it supports manual operation as well as integration with automated liquid handling systems. This makes it a reliable, efficient, and powerful solution for researchers seeking high-quality HMW DNA to meet the demands of high-throughput long-read sequencing workflows.

Features:

- Works with HMW DNA, fragmented DNA, sheared DNA, and large PCR amplicons
- Removal of short DNA fragments, impurities, and unwanted reaction components
- Effective purification and recovery of HMW DNA, dsDNA fragments 5kb or longer.
- Improvement in the read length of long-read sequencing.
- Expectant recover Yield: Up to 90%
- Compatible with manual and automated processing and cost effective.

Kit Contents

Product Number	S005-01	S005-02
Auto-Mag® HMW Size Select	10 ml	100 ml
Elution Buffer	10 ml	100 ml

Storage and Stability

Auto-Mag® HMW Size Select is shipped at ambient temperature and is stable for at least 12 months after delivery when stored at 2-8°C. Contents of the kit should never be frozen at any time.

Preparation of Reagents

1. Prepare 80% Ethanol wash buffer. (Prepare from absolute ethanol. Do not use denatured alcohol).

Ethanol is hygroscopic. When opened the ethanol will both evaporate and absorb water over time. Fresh prepare 80% ethanol then keep cover tight and use within one week.

Additional Information

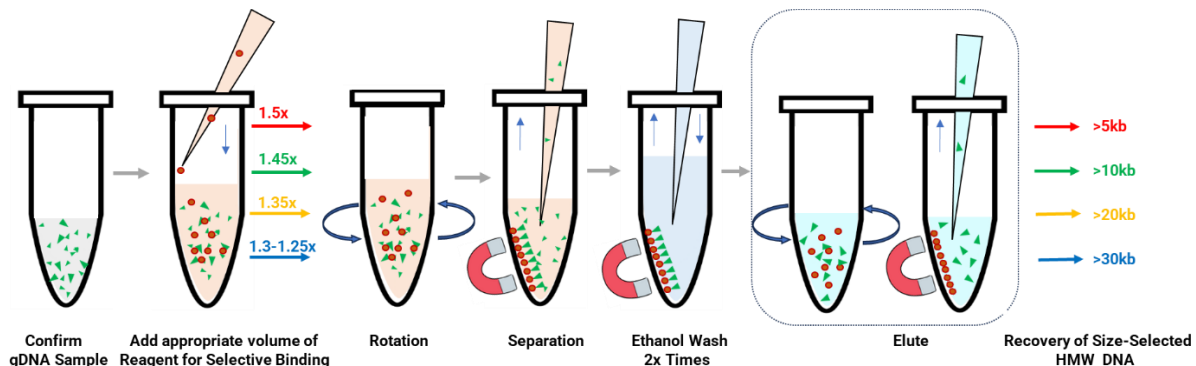
1. Heterogeneity and viscosity

The size selection accuracy and recovery efficiency of the Auto-Mag® MHW Size Select kit depend in part on the input DNA being homogeneous and fully in solution. HMW gDNA can sometimes be difficult to re-solubilize after extraction and results in inhomogeneous or contains sticky jellies. Such samples may result in low yields and short DNA carryover. If the HMW gDNA sample is inhomogeneous or contains sticky jellies, we recommend shearing sample with a 26G needle 2-5 times before beginning size selection. Sample homogeneity can be evaluated by performing triplicate concentration measurements and verifying that the concentration CV is <10%.

2. Pipetting

When adding, mixing sample, use wide bore pipette tips. Do not disturb beads when removing liquid from the microcentrifuge tube or well of plate. Carefully insert the pipette tip against the wall to dispense, likewise, remove liquid by pipetting from the liquid surface.

HMW DNA cleanup and size selection process workflow



1. Confirm the Quality and volume of input HMW DNA samples
2. Add Auto-Mag® MHW Size Select reagent and incubation at room temperature (15-25°C)
3. Bind sample to paramagnetic beads and discard supernatant
4. Wash beads with 80% ethanol to remove contaminants
5. Elute and recovery size selected HMW DNA
6. Transfer eluate to new vessel and ready for next applications.

Auto-Mag[®] MHW Size Select Protocols

Required Materials and Equipment

- Single-tube format: 1.5 ml Eppendorf DNA LoBind tube and Magnetic Rack Separator for tube.
- 96-well format: 96 well thermal cycling plate, or 500µl round bottom microtiter plate, and appropriate magnetic separation device.
- 200µl wide bore pipette tips
- Well calibrated pipettor and Disposable pipette tips.
- Polypropylene reservoirs
- DNA quantification equipment. (Thermo Fisher Scientific NanoDrop 2000. Thermo Qubit 3.0, Qubit dsDNA BR Assay Kit, etc.)
- Laboratory mixer, vortex, or equivalent.
- 80% ethanol (Prepare from absolute ethanol. Do not use denatured alcohol).

Before Starting

- Prepare 80% Ethanol for DNA wash steps.
- Bring the Auto-Mag[®] MHW Size Select reagent to room temperature for at least 30 minutes before use.
- Complete suspending Auto-Mag[®] MHW Size Select reagent by vortex.
- Check the pipettor and confirm it is well calibrated.

Sample Quality Control

Long-read sequencing requires high-quality, high molecular weight (HMW) DNA. Before performing DNA cleanup, it is necessary to evaluate the quantity and size distribution of input DNA to determine whether it is suitable for the protocol.

- To isolate HMW gDNA with a qualify HMW DNA isolation kit and 50% of HMW DNA should be >30 kb.
- DNA concentration should be measured using a Qubit dsDNA Broad Range Assay or equivalent and concentration is between 5–150 ng/µl. If the DNA concentration is over 150ng/µl, dilute sample using TE buffer (pH 8), Elution Buffer, or water.
- DNA samples: including HMW DNA, fragmented DNA, sheared DNA, or partially degraded genomic DNA, should be in DNA Elution buffer, TE buffer (pH 8.0) or water and concentration is between 10–150 ng/µl. If you are unsure of the buffer composition or the DNA is not in elution buffer, purify with 1X Auto-Mag[®] X-Pure Size Select or AMPure XP beads and elute in elution buffer or an equivalent low-salt buffer (i.e., TE buffer, 10 mM Tris-HCl, pH 8.0, or water).
- Sample volume: Although the starting sample volume can be any volume greater than 10ul. However, the size range of recovered gDNA is sensitive to changes in the volume ratio of reagent to sample. Using a smaller sample volume may increase the chance of size shift of recovered DNA by sampling error. It is recommended to use 50µl of sample. If a smaller sample volume must be used, please refer to the volume ratio of reagent to sample corresponding to the size range of recovered DNA and use a more accurate pipette.

Protocol for Short DNA Fragments Depletion and HMW DNA Size Selection.

This procedure describes the workflow to remove short DNA molecules from unsheared high molecular weight (HMW) genomic DNA, sheared genomic DNA fragments, or partially degraded genomic DNA samples. By adjusting the ratio of reagents to samples, the desired size ranges of HMW DNA fragments can be recovered. The recovered HMW DNA is also cleaned during the process which could help remove contaminants from samples.

1. Completely float the Auto-Mag® MHW Size Select reagent until it appears homogeneous in color.
2. Pulse vortex or pipette mix each sample to homogenize the DNA in solution. Adjust the DNA concentration between 10–150 ng/μl.

Note: Quick spin each sample to collect liquid if necessary.

3. Pipette 50μl of the sample into a 1.5 ml Eppendorf DNA LoBind tube, or the well of 96 plate (96 well format)
4. According to the table below, add the appropriate volume of Auto-Mag® MHW Size Select reagent to 50ul of sample for selection of desired size DNA fragments and depletion of the smaller DNA fragments.

For example: for selection of fragments >10 kb or for depletion of fragments <10 kb, add 72.5ul (1.45x) of Auto-Mag® MHW Size Select reagent to the 50ul sample.

Table 1. recommended ratio of Auto-Mag® MHW Size Select to the gDNA sample for depletion of short DNA fragments and size selection

Size of short DNA to be removed	Ratio of Auto-Mag® MHW Size Select / Sample	Auto-Mag® MHW Size Select Needed (μl) (For 50μl sample)
< 5 kb	1.5 x	75μl
< 10 kb	1.45 x	72.5μl
<20kb	1.35x	67.5μl
<30kb	1.3-1.25x*	65-62.5μl*

* When selecting DNA fragments above 30kd, the concentration and content of DNA above 30kd in the sample may affect the selection results and recovery rate. Sometimes, the ratio of reagent to sample may need to be adjusted slightly to harvest the desired DNA.

5. Mix the sample by gently pipetting up and down 15 times with wide bore pipette tips.
- Note: It is important to mix well.*
6. Incubate sample at room temperature for 15 minutes at least for maximum recovery.
 7. Place the sample tubes or plate on a compatible magnetic separation device for 5 minutes or until the solution clears. Beads will pull to the side of the well.

Note: The magnetic separation time may relate to the strength of the magnetic separation device and the sample volume. If necessary, gently pipette the liquid up and down twice to speed up magnetic bead separation. Do not touch the beads.

8. With the sample still on the magnet, carefully remove the supernatant by pipetting the liquid out which contains the unwanted smaller DNA fragments. **Do not disturb the beads while pipetting out the supernatant.**

Note: The supernatants have unbinding short DNA fragments. If unbound DNA needs to be recovered, transfer supernatant to a new tube or 96-well plate. Reference the protocol of unbinding short DNA fragments to recover the small DNA fragments.

9. With the samples still on the magnet, slowly add 200µl of 80% ethanol to each sample and gently pipet the liquid up and down twice. After incubating 30 seconds, slowly pipet out and discard all of liquid. Do not touch the beads.

Note: 80% ethanol is hygroscopic and should be prepared FRESH to achieve optimal results.

10. Repeat steps 9 two more times.

11. Keep the sample on the magnet, and air dry the magnetic beads at room temperature for 3 minutes. Remove any residue liquid with a pipettor.

Note: It is critical to completely remove all traces of alcohol but take caution in not over drying the beads as this will significantly decrease elution efficiency.

12. Remove the sample from the magnet. Add 20–100 µl of DNA Elution Buffer, ensuring that the beads clumped against the wall are completely submerged in the buffer. Do not suspend or mix the sample at this stage. Allow the sample to stand at room temperature for 5 minutes.

13. After 5 minutes incubation, gently pipetting up and down elution buffer 15 times with wide-bore pipette tips to ensure the beads are properly re-suspended in the elution buffer. Incubate sample again at room temperature for an additional 5 minutes.

Note: When resuspending magnetic beads, the volume of the pipettes should be adjusted to be smaller than the volume of Elution buffer to prevent bubbles generated when suspending magnetic beads.

14. Place the sample back on the magnet for 5 minutes or until the magnetic beads are completely cleared from solution.

15. Transfer the eluate to an appropriate storage vessel and analyze the recovery and purity of the DNA using a NanoDrop and Qubit system.

16. DNA can be stored in Elution Buffer at 4°C for several months. Keep at -20°C for long term storage. Avoid freeze/thaw cycles since this can degrade high molecular weight DNA.

Supplemental protocol for recovery unbinding short DNA fragments

The following protocol is for recovery all unbinding DNA fragments from supernatant of removing short DNA fragments and size selection procedure by using Auto-Mag® X-Pure select or compatible reagents.

Additional Materials and Equipment:

- Auto-Mag® PCR-Pure (Cat: AMD-S002) or compatible reagents.

Before Starting

- Prepare 80% ethanol for wash steps. A minimum of 0.5 ml is required per sample.
- Complete suspending Auto-Mag® PCR-Pure by vortex.

Procedure

1. Confirm the volume of supernatant from Step 8 of removing short DNA fragments and size selection procedure.
2. Add 1x volume of Auto-Mag® PCR-Pure reagents to the sample.
3. Mix thoroughly the Auto-Mag® PCR-Pure reagent and sample by pipetting up and down 15 times. Incubate at room temperature for 10 minutes.
4. Place the samples on the magnetic separation device for 3-5 minutes or until the Auto-Mag® X-Pure Select beads are completely cleared from solution. Carefully remove and discard the supernatant.
5. Keep the sample on the magnetic separation device, add 200µl of 80% Ethanol to each sample and incubate for 30 seconds at room temperature. Carefully remove and discard the supernatant.

Note: It is not necessary to resuspend the Auto-Mag® PCR-Pure

6. Repeat step 5 for second 80% ethanol wash.
7. Keep the sample on the magnet, and air dry the magnetic beads at room temperature for 5 minutes. Remove any residue liquid with a pipettor.
8. Remove the sample from the magnet. Add 20-50µl of elution buffer and pipette mixing 20 times. Ensure beads are no longer attached to the side of the well.
9. Incubate the sample at room temperature for 5 minutes.
10. Place the sample back on the magnetic separation device and wait 5 minutes or until the magnetic beads are completely cleared from the solution.
11. Transfer the eluate containing the recovered DNA to an appropriate storage vessel and keep at -20°C for long term storage.

Customized Protocol and Programmed Procedure for Automation Purification

To obtain a custom protocol for a specific fragment size selection; or automating this procedure on a liquid handler or a magnetic processor, please contact AMD Biotech for instrument-specific instructions or additional processing procedures.

Phone: 404-290-5063 (in US), Email: support@amdbiotech.com

Troubleshooting Guide

Please use this guide to troubleshoot any problems that may arise. For further assistance, please contact technical support via Phone: 1-404-290-5063 (in US), Email: support@amdbiotech.com

Observation	Possible Causes	Comments
Low yield/ Incorrect recovery	Highly fragmented gDNA.	Recovery will be low if DNA is not HMW. Verify the 50% of input DNA should be ≥ 30 kb
	Low concentration of input DNA	Verify the input gDNA concentration using Qubit dsDNA broad range assay or equivalent. Try increasing concentration of input DNA up to the maximum of 150 ng/ μ l.
	Insufficient Mixing	Mixing thoroughly during the initial bind mix and elution mix is critical. Ensure the beads get sufficiently resuspended.
	Bead Loss	If beads get aspirated into tips during supernatant removal, the nucleic acid bound to these beads will also be lost. Aspirate slowly and remove as much of the first supernatant as possible without disturbing the bead.
Fragment size incompatible	Handling error.	Ensure that proper care is taken with pipetting steps.
	Insufficient ratio	Use volume ratios outlined in this manual.
	Insufficient ethanol concentration used for washing step	Use freshly prepared 80% ethanol. Over time ethanol becomes more diluted through evaporation and absorption of atmospheric water. Therefore, parts of the DNA pellet go into solution and DNA fragments are washed away.
	Elution buffer volume insufficient	Bead pellet must be covered completely with elution buffer
	Beads over dried	Do not dry beads for longer than 15 minutes at room temperature. Over drying of beads may result in lower elution efficiencies.
Downstream applications are unsuccessful	Carry-over of ethanol from washing step	Be sure to remove all the ethanol after the final wash step. Dry beads 3-10 minutes at room temperature.
Carry-over of beads	Time for magnetic separation too short	Increase separation time to allow the beads to be attracted to the magnetic pins completely

Ordering Information

Product Description	Catalog No.	Size
Auto-Mag® Mega Select	S005-01	10 ml
	S005-02	100 ml
DNA Elution Buffer	B228-01	50 ml
	B228-02	250 ml
	B228-03	500 ml
Auto-Mag® PCR-Pure	S002-01	5 ml
	S002-02	50 ml
	S002-03	250 ml
	S002-04	500 ml

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