

# iPure kit v2

# Magnetic DNA Purification kit for epigenetic applications

Cat. No. C03010014 (x24) Cat. No. C03010015 (x100)





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## Introduction

Diagenode's **IPure kit** is the only DNA purification kit using magnetic beads, that is specifically optimized for extracting DNA from **ChIP**, **MeDIP** and **CUT&Tag** (Chromatin Immunoprecipitation, Methylated DNA Immunoprecipitation and Cleavage Under Targets & Tagmentation).

It's a simple and straightforward protocol that delivers pure DNA ready for any downstream application (e.g. next generation sequencing). This approach guarantees a minimal loss of DNA and reaches significantly higher yields than a column purification (see results pag). Comparing to phenol-chloroform extraction, the IPure technology has the advantage of being nontoxic and much easier to be carried out on multiple samples. The use of the magnetic beads allows for a clear separation of DNA and increases therefore the reproducibility of your DNA purification.

#### The kit IPure can be used with the Diagenode's DiaMag02 magnetic rack:



DiaMag02 - magnetic rack Cat. No. B04000001

- Holds 16x standard 0.2 ml tubes
- Working volume: 25-200 µl

### Kit Method Overview



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### Workflow description

### IPure after ChIP

**Step 1:** Chromatin is decrosslinked and eluted from beads (magnetic or agarose) which are discarded. Magnetic beads for purification are added.

**Step 2:** Magnetic beads acquire positive charge to bind the negatively charged phosphate backbone of DNA. DNA-bead complex is separated using a magnet.

Step 3: Proteins and remaining buffer are washed away.

**Step 4:** DNA is eluted from magnetic beads, which are discarded. Purified DNA is ready for any downstream application (NGS, qPCR, amplification, microarray).

### **IPure after MeDIP**

**Step 1:** DNA is eluted from beads (magnetic or agarose) which are discarded. Magnetic beads for purification are added.

**Step 2:** Magnetic beads acquire positive charge to bind the negatively charged phosphate backbone of DNA. DNA-bead complex is separated using a magnet.

Step 3: Remaining buffer are washed away.

**Step 4:** DNA is eluted from magnetic beads, which are discarded. Purified DNA is ready for any downstream application (NGS, qPCR, amplification, microarray).

### IPure after CUT&Tag

**Step 1:** pA-Tn5 is inactivated and DNA released from the cells. Magnetic beads for purification are added.

**Step 2:** Magnetic beads acquire positive charge to bind the negatively charged phosphate backbone of DNA. DNA-bead complex is separated using a magnet.

Step 3: Proteins and remaining buffer are washed away.

**Step 4:** DNA is eluted from magnetic beads, which are discarded. Purified DNA is ready for any downstream application (NGS, qPCR, amplification, microarray).

# Kit Content

The kit content is sufficient to perform 24 (C03010014) or 100 reactions (C03010015).

IPure Kit			
Description	Quantity (x24)	Quantity (x100)	Storage
200 µl tube strips (8 tubes/strip) + cap strips	4 pc	15 pc	Room temperature
Buffer A	2.8 ml	10 ml	4°C
Buffer B	115 µl	460 µl	4°C
Wash buffer 1 w/o isopropanol	1.5 ml	9 ml	4°C
Wash buffer 2 w/o isopropanol	1.5 ml	9 ml	4°C
Buffer C	1.2 ml	9 ml	4°C
IPure Beads v3	240 µl	1 ml	4°C
Carrier*	48 µl	200 µl	-20°C

\*This product is shipped at 4°C. Store it at -20°C upon arrival. This is an optimized buffer (NOT CARRIER DNA).

# **Required Materials Not Provided**

- DiaMag02 magnetic rack (B04000001)
- Microcentrifuge for 0.2 ml tubes or for 1.5 ml tubes with corresponding adaptor.
- 100% isopropanol
- Sodium Acetate
- 70% ethanol
- 100% ethanol
- DNase-free water
- For CUT&Tag:
  - 0.5M EDTA
  - 10% SDS
  - Thermolabile proteinase K (eg New England Biolabs P8111S)



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#### IPure after ChIP

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# STEP 1A Chromatin Reverse Crosslinking and Elution

After the last wash of immunoprecipitated material, discard the last traces of wash buffer and use the pellet of beads (8-tube strip) for STEP 1.1.

### 1.1 Prepare the Elution Buffer by mixing Buffer A and B as follows:

Elution Buffer	1 rxn
Buffer A	115.4 µl
Buffer B	4.6 µl
Total Volume	120 μl

- Place Buffer A at 25°C during 30 minutes before use.
- 100 µl of Elution Buffer are needed per IPure reaction (20 µl excess).
- 1 IPure reaction corresponds to the purification of 1 ChIP or 1 input sample.

**NOTE:** Make sure when working with Buffer A, that there are no crystals left in solution. Otherwise heat up gently and mix until complete disappearence of such crystals.

**1.2** Add **100 µl of Elution Buffer** to the bead pellets (tube strip).

**1.3** Thaw your input sample (1.5 ml tube), and perform a pulse spin. Add **90 μl of Elution Buffer** and **10 μl of input sample** to a new 200 μl tube (8 tube-strip).

**NOTE:** Input sample corresponds to 10% of the IP sample.

- **1.4** Incubate samples and input for 4 hours (or overnight) at 65°C on a thermomixer, with continuous shaking.
- **1.5** Spin the 8-tube strip and place it into the **DiaMag02** magnetic rack. After 1 minute, transfer the supernatants to a new labelled 8-tube strip. Keep the samples on ice.



### **IPure after ChIP**

# STEP 2A DNA Binding

- **2.1** Add **2 μl of carrier** to each IP and input sample. Vortex briefly and perform a short spin.
- **2.2** Add **100 µl of 100% isopropanol** to each IP and input sample. Vortex briefly and perform a short spin.

**NOTE:** Following the addition of isopropanol the solution may become cloudy. This is not detrimental to your experiment and will not influence the quality or quantity of your purified DNA.

- **2.3** Resuspend the provided **Magnetic beads** and transfer 10  $\mu$ l to each IP and input sample.
  - Keep Magnetic beads in liquid suspension during storage at 4°C and at all handling steps, as drying will result in reduced performance.
  - The final volume is now 212 µl per IPure reaction.
- **2.4** Incubate IP and input samples for 10 minutes at room temperature on a rotating wheel (40 rpm).

### **IPure after MeDIP**

# STEP 1B DNA Elution

After the last wash of immunoprecipitated material, discard the last traces of wash buffer and use the pellet of beads (8-tube strip) for STEP 1.1.

1.1 Prepare the Elution Buffer by mixing Buffer A and B as follows:

Elution Buffer	1 rxn
Buffer A	115.4 µl
Buffer B	4.6 µl
Total Volume	120 µl

- Place Buffer A at 25°C during 30 minutes before use.
- 100 µl of Elution Buffer are needed per IPure reaction (20 µl excess).
- 1 IPure reaction corresponds to the purification of 1 MeDIP or 1 input sample.

**NOTE:** Make sure when working with Buffer A, that there are no crystals left in solution. Otherwise heat up gently and mix until complete disappearence of such crystals.

- **1.2** Add **50 µl of Elution Buffer** to the bead pellets (tube strip).
- 1.3 Thaw your input sample (1.5 ml tube), and perform a pulse spin. Add
  92.5 μl of Elution Buffer and 7.5 μl of input sample to a new 200 μl tube (8-tube strip).

**NOTE:** Input sample corresponds to 10% of the IP sample.



- **1.4** Incubate samples and input DNA for 15 minutes at room temperature on a rotating wheel (40 rpm).
- 1.5 Spin the 8-tube strip and place it into the DiaMag02 magnetic rack. After 1 minute, transfer the supernatants to a new labelled 8-tube strip.
- **1.6** Repeat the incubation of the bead pellets for 15 minutes at room temperature on a rotating wheel (40 rpm) in **50 µl Elution Buffer**.
  - For input DNA samples: 1 elution in 100 µl.
  - For MeDIP samples: 2 elutions in 50 µl (total volume 100 µl).
- 1.7 Spin the 8-tube strip. Place the 8-tube strips into the DiaMag02 containing the 50 µl IP samples, wait 1 minute and transfer the supernatants to the new labelled 8-tube strip to pool with the corresponding IP samples.
  - $\bullet$  Elutions of IP and input samples are now completed in 100  $\mu l$  and are in the same 8-tube strip.

**IPure after MeDIP** 

# STEP 2B DNA Binding

- **2.5** Add **2 μl of carrier** to each IP and input sample. Vortex briefly and perform a short spin.
- **2.6** Add **100 µl of 100% isopropanol** to each IP and input sample. Vortex briefly and perform a short spin.

**NOTE:** Following the addition of isopropanol the solution may become cloudy. This is not detrimental to your experiment and will not influence the quality or quantity of your purified DNA.

- **2.7** Resuspend the provided **Magnetic beads** and transfer 10 μl to each IP and input sample.
  - Keep Magnetic beads in liquid suspension during storage at 4°C and at all handling steps, as drying will result in reduced performance.
  - The final volume is now 212 µl per IPure reaction.
- **2.8** Incubate IP and input samples for 10 minutes at room temperature on a rotating wheel (40 rpm).

# STEP 1C

### pA-Tn5 Inactivation and DNA Release

**NOTE**: the protocol below is compatible with CUT&TAG version described by Kaya-Okur, H.S., Wu, S.J., Codomo, C.A. et al. CUT&TAG for efficient epigenomic profiling of small samples and single cells. Nat Commun 10, 1930 (2019). https://doi.org/10.1038/s41467-019-09982-5

- 1.1 After completion of the tagmentation reaction, add 10 μl of 0.5 M EDTA, 3 μl of 10% SDS and 1 μl of thermolabile proteinase K directly to each sample and mix by full-speed vortexing for 2 seconds. Do not remove the Tagmentation Buffer and Concavalin A beads.
- **1.2** Incubate at 37°C for 15 minutes.
- **1.3** Heat samples at 55°C for 10 minutes to inactivate thermolabile proteinase K.

# STEP 2C

### **DNA** Binding

**NOTE**: an equival volume of isopropanol should be added. If the tagmentation reaction volume is different from 300  $\mu$ l, adjust the required volume of isopropanol.

2.1 Add 300 µl of 100% isopropanol to each sample. Vortex briefly and perform a short spin. Do not remove Concavalin A beads. Following the addition of isopropanol the solution may become cloudy. This is not detrimental to your experiment and will not influence the quality or quantity of your purified DNA.

**NOTE**: an equival volume of isopropanol should be added. If the tagmentation reaction volume differs from 300  $\mu$ l, adjust the required volume of isopropanol.

- **2.2** Re-suspend the provided Magnetic beads and transfer 10 μl to each sample.
  - Keep Magnetic beads in liquid suspension during storage at 4°C and at all handling steps, as drying will result in reduced performance.
  - $\bullet$  The final volume is now 624  $\mu l$  per IPure reaction.
- **2.3** Incubate samples for 10 minutes at room temperature on a rotating wheel (40 rpm).

# STEP 3

### Washes (ChIP, MeDIP and CUT&TAG)

**3.1** Prepare the **Wash Buffer 1** containing 50% isopropanol for 100 reactions:

Wash Buffer 1	24 rxns	100 rxns
Wash Buffer 1 w/o Isopropanol	1.5 ml	9 ml
Isopropanol (100%)	1.5 ml	9 ml
Total Volume	3 ml	18 ml

- Never leave the bottle open to avoid evaporation.
- Briefly spin the tubes, place in the DiaMag02, wait 1 minute and discard the buffer. Keep the captured beads and add per tube, 100 μl Wash Buffer 1. Close the tubes, invert the 8-tube strip to resuspend the beads and incubate for 5 minutes at room temperature on a rotating wheel (40 rpm).
  - Do not disturb the captured beads attached to the tube wall.
  - Always briefly spin the tubes to bring down liquid caught in the lid prior to positioning into the Diagenode Magnetic Rack.

### **3.3** Prepare the **Wash Buffer 2** containing 50% isopropanol as follows:

Wash Buffer 2	24 rxns	100 rxns
Wash Buffer 2 w/o Isopropanol	1.5 ml	9 ml
Isopropanol (100%)	1.5 ml	9 ml
Total Volume	3 ml	18 ml

• Never leave the bottle open to avoid evaporation.

- **3.4** Wash the IP and input samples with the **Wash Buffer 2** as follows. Briefly spin the tubes, place into the **DiaMag02**, wait 1 minute and discard the buffer. Keep the captured beads and add per tube, 100 μl Wash Buffer 2. Close the tubes, invert the 8-tube strip to resuspend the beads and incubate for 5 minutes at room temperature on a rotating wheel (40 rpm).
  - Do not disturb the captured beads attached to the tube wall.
  - Always briefly spin the tubes to bring down liquid caught in the lid prior to positioning in the **DiaMag02**.

# **STEP 4** DNA Elution (ChIP, MeDIP and CUT&TAG)

**NOTE**: This elution buffer (Buffer C) is suitable for direct qPCR analysis, whole genome amplification, chip hybridization and next generation sequencing.

- 4.1 Briefly spin the tubes and place them into the DiaMag02, wait 1 minute and discard the buffer. Keep the captured beads and add per tube, 25 µl of Buffer C (alternatively, a higher volume can be used for the elution if necessary). Close the tubes, invert the 8-tube strip to resuspend the beads and incubate for 15 minutes at room temperature on a rotating wheel (40 rpm). Resuspend the pelleted beads using the pipet and make sure that you drop them on the bottom of the tube.
- **4.2** Spin the 8-tube strips and place it into the **DiaMag02**, wait 1 minute and transfer the supernatants into a new labelled 1.5 ml tube. Keep the bead pellets on ice.
- **4.3** Place the DNA on ice and proceed to any desired downstream applications, or store it at -20°C or -80°C until further use.

## Results



### DNA recovery after purification of ChIP samples using IPure technology

ChIP assays were perfomed using different amounts of U2OS cells, the LowCell# ChIP kit and the H3K9me3 antibody (Cat. No. C15410056; 2  $\mu$ g/IP). The purified DNA was eluted in 50  $\mu$ l of water and quantified with a Nanodrop.



### **Related Products**

Description	Cat. No.	Format
DiaMag02 - magnetic rack	B04000001	1 pc
200 µl tube strips (8 tubes/strip) + cap strips	C30020002	120 рс
iDeal ChIP-seq kit for Histones x10	C01010050	10 rxns
iDeal ChIP-seq kit for Transcription Factors x10	C01010054	10 rxns
MagMeDIP kit x10	C02010020	10 rxns
MagMeDIP kit x48	C02010021	48 rxns
pA-Tn5 loaded	C01070001	15 µl/30 µl
pA-Tn5 unloaded	C01070002	10 µl/30 µl

### **Revision history**

Version	Date of modification	Description of modifications
V5 08_2023	August 2023	Correction of small error page 17, step 2.2.
V5 02_2023	February 2023	- Replacement of the IPure beads v2 by IPure beads v3 - Removal of obsolete reference (DiaMag1.5)

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