

# User Guide



YOUSEQ

## NGS Library Clean-up kit

Magnetic bead-based cleaning kit for NGS libraries

Version 1.2

# Kit contents

## Bead clean pack

	Cap colour	Size
YouSeq beads	●	5 ml
Wash buffer*	●	80 ml*
Elution buffer	●	10 ml

\*Add 64 ml of molecular biology grade ethanol to the supplied diluent to make the wash buffer ready to use.

## Other items to be supplied by the user

### Essential

- Magnetic rack
- Pipettes and general laboratory equipment

## Introduction and kit principles

The YouSeq bead clean-up kit uses magnetic beads that bind to your DNA Library. A magnet then pulls the beads to one side so that all contaminants and artefacts can be washed away. A couple of washes completes the cleaning process then a simple elution step washes the DNA library off the beads in to solution.

The kit can be used to remove either small and/or large (or both) unwanted fragments.

# Size selection – Principles.

You can remove unwanted DNA fragments of a specific size by simply adding more or less beads. The ratio of beads vs volume of sample dictates the size of DNA that will be captured.

## Size selection – removing small fragments.

Ration beads /Sample	Your Library Size bp
0.45	600-900
0.5	400-600
0.6	300-400
0.65	200-300
0.9	150-200
1.2	100-150

## Size selection – removing large fragments.

You can remove unwanted large DNA fragments by capturing them smaller fragments on the beads and collect the supernatant fraction instead

Ration beads /Sample	Your Library Size bp
0.4	2000-4000
0.45	1000-2000
0.5	700-1000
0.55	500-700
0.6	450-500
0.65	350-450
0.7	300-350

e.g. if a 0.9x ratio is required on 20  $\mu$ l PCR reaction

- $0.9 \times 20 = 18$
- So, 18  $\mu$ l beads needs to be added during the clean-up

# Example protocol to remove small fragments

E.g. a 0.8x bead ratio was required.

*\*Bring the beads to room temperature by removing from fridge 30 minutes before use\**

## Bind

- Mix beads well until suspension is homogeneous
- Add **16  $\mu\text{L}$**  \* YouSeq NGS Beads to 20 $\mu\text{l}$  of your NGS Library and mix well
- Incubate for 3 mins at room temp
- Place on magnet and wait until liquid is clear
- Remove supernatant

## 1<sup>st</sup> wash

- Leave your tube on the magnet

**\*N.B.** ensure the wash buffer has had correct volume of ethanol added\*

- Add **200  $\mu\text{L}$**  wash buffer
- Incubate for 30s
- Remove and discard the supernatant carefully

## 2<sup>nd</sup> wash

- Repeat as per 1<sup>st</sup> wash
- Ensure all remainder ethanol has been removed

## Dry beads

- Leave until noticeably dry (cracking) on the magnet \* (should be around 10 minutes)

**\*N.B.** if you notice any residual buffer on the sides of the tubes, carefully remove with pipette.

## Elute

- Add **20  $\mu\text{L}$**  Elution buffer
- Remove your tube from the magnet and mix well
- Incubate for **5 min** at RT
- Place on magnet and wait until liquid is clear

DNA is in the solution.

*\*This example is for removing DNA fragments of around 250bp and smaller. See tables above for precise volumes/ratios to use for your particular experiment*

# Example protocol to remove large fragments

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E.g. a 0.45x bead ratio was required.

*\*Bring the beads to room temperature by removing from fridge 30 minutes before use\**

## Bind

- Mix beads well until suspension is homogeneous
- Add 9 $\mu$ L \*\* YouSeq NGS Beads to 20 $\mu$ l of your NGS Library and mix well
- Incubate for 10 mins at room temp
- Place on magnet and wait until liquid is clear
- Remove supernatant

This step can be repeated to improve the stringency of the clean-up if necessary

Your size selected Library is in the supernatant.

\*\* This example is for removing DNA fragments of around 1000bp and above. See tables above for precise volumes/ratios to use for your particular experiment

# Example protocol for Dual Size Selection

E.g. all products between 0.45x – 1.0x bead ratios

*\*Bring the beads to room temperature by removing from fridge 30 minutes before use\**

## Bind large fragments

- Mix beads well until suspension is homogeneous
- Add **9  $\mu\text{L}$**  YouSeq NGS Beads to 20  $\mu\text{L}$  of your NGS Library and mix well
- Incubate for 3 mins at room temp
- Place on magnet and wait until liquid is clear
- Transfer 25  $\mu\text{L}$  supernatant to fresh tube

## Bind small fragments

- Mix beads well until suspension is homogeneous
- Add **25  $\mu\text{L}$**  YouSeq NGS Beads to 25  $\mu\text{L}$  of your NGS Library and mix well
- Incubate for 3 mins at room temp
- Place on magnet and wait until liquid is clear
- Remove the supernatant

## 1<sup>st</sup> wash

- Leave your tube on the magnet

**\*N.B.** ensure the wash buffer has had correct volume of ethanol added\*

- Add **200  $\mu\text{L}$**  wash buffer
- Incubate for 30s
- Remove and discard the supernatant carefully

## 2<sup>nd</sup> wash

- Repeat as per 1<sup>st</sup> wash
- Ensure all remainder ethanol has been removed

## Dry beads

- Leave until noticeably dry (cracking) on the magnet \* (should be around 10 minutes)

**\*N.B.** if you notice any residual buffer on the sides of the tubes, carefully remove with pipette.

## Elute

- Add **20  $\mu\text{L}$**  Elution buffer
- Remove your tube from the magnet and mix well
- Incubate for **5 min** at RT
- Place on magnet and wait until liquid is clear

DNA is in the solution.