

Complete your workflow with the **riboPOOL™ cleanUP** module

The final step of the riboPOOL kits before library preparation

Why use the riboPOOL cleanUP module?

RNA samples that have been subject to rRNA depletion must be purified before sequencing library preparation to remove salts and buffer concentrates. The riboPOOL cleanUP module utilizes magnetic beads and buffers to efficiently purify your rRNA depleted RNA sample prior to downstream applications

How does the riboPOOL cleanUP module work?

The **riboPOOL™ cleanUP module** uses Nucleic Acid Binding M-PVA magnetic beads, that hybridize to RNA. The final RNA product is eluted from the beads for further processing.

✓ Single-product solution

With the addition of the **riboPOOL™ cleanUP module**, you have everything you need to prepare your RNA samples. There is no need to buy extra reagents or beads for the RNA clean-up step anymore.

✓ Faster and more convenient

Our easy-to-follow protocol has been optimized for faster results and can be completed within ~30-45 minutes.

✓ Fully automatable and scalable

The clean-up beads' high magnetite content allows for fast magnetic separation, even in large sample volumes making it easy to automate the workflow.

Available Formats:

riboPOOLS kits with cleanUP module

12 rx Catalog-No. dp-P012	24 rx Catalog-No. dp-P024	96 rx Catalog-No. dp-P096
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For further questions contact us at:

info@sitools.de

Simple & Fast Workflow

The workflow can be completed within 30-45 minutes. The kit includes magnetic clean-up beads (CB), binding buffer (BB), wash buffer (WB) and nuclease-free water. Below is a protocol snapshot for the **riboPOOL™ cleanUP module**.

