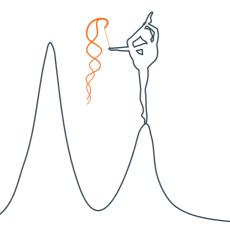
mRNA and other ncRNA rRNA

Ribosomal RNAs occupy up to 90% of RNA-Seq reads.

RIbosomal RNAs (rRNA) are highly abundant and limit detection of relevant coding and non-coding RNAs by Next Generation RNA Sequencing (RNA-Seq), microarray and other analysis. Their removal greatly improves and economizes RNA-Seq experiments.

riboPOOL™ for efficient and flexible ribosomal RNA depletion

riboPOOL uses siTOOLs' technology of high complexity pooling of optimally designed biotinylated oligos to deplete rRNAs with high efficiency and specificity. Available for any species or any RNA sequence, riboPOOLs provide a superbly flexible solution for scientists with demanding sequencing applications.



riboPOOL Key Benefits

Any Species

Deplete rRNAs from any species. Diverse species in-stock with Custom riboPOOLs on-demand.

Affordable

riboPOOLs are priced competitively and suitable for high-throughput sequencing.

Highly Efficient and Specific

Human and mouse riboPOOLs depleted > 95% cvtoplasmic and mitochondrial rRNAs.

HPLC Purified

Low risk of contaminants.

Applicable for Nonpolyadenylated RNA

Detects non-polyadenylated RNA (e.g. snoRNA, IncRNA, histones) with uniform transcript coverage.

Suitable for Metatranscriptomics

Pan-prokaryote and Combination riboPOOLs available for samples containing multiple species.

Deplete Custom RNA

Deplete any abundant RNA of choice. E.g. globin from whole blood samples

Fast Workflow

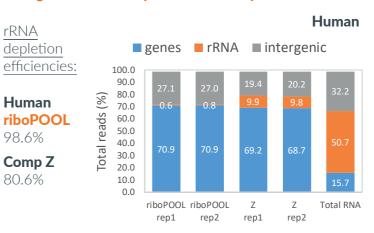
Workflow completed in ~70 min with column-based purification.

Broad Input Range

Efficient rRNA depletion between 100 ng and 5 μg input RNA

Data with riboPOOL

10% greater rRNA depletion efficiency with riboPOOL



Mouse

96.4%

Comp Z

riboPOOL

Mouse

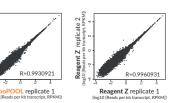
■ genes ■ rRNA ■ intergenic

rep2

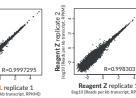
rep1

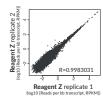
High reproducibility with both reagents





Mouse





Study performed by Core Unit Systems Medicine (University of Würzburg) and IMGM Laboratories GmbH with protocol developed by Kim et al. (University of Bayreuth)₁. RNA (1 μg, RIN ≥ 7.5) from HeLa cells or mouse embryonic fibroblasts subject to rRNA depletion with human or mouse riboPOOLs vs. human/mouse/rat rRNA depletion kit from Competitor Z and sequenced with Illumina TruSeg® Stranded Total RNA technology.

1 Kim, I et. al (2019) Efficient depletion of ribosomal RNA for RNA sequencing in planarians. bioRxiv (Preprint with riboPOOL)

riboPOOLs for Prokaryotes

Pseudomonas aeruginosa

Staphylococcus aureus

Salmonella enterica

Haloferax volcanii

Stenotrophomonas sp.

Caulobacter crescentus

Clostridium perfringens

Escherichia coli

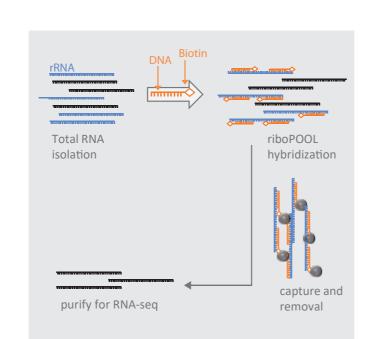
Bacillus subtilis

Pan-Prokaryote (Universal Microbe)

riboPOOL Workflow

Great for metatranscriptomics and microbial sequencing

For microbial sequencing, the Pan-Prokaryote riboPOOL can be broadly applied across many microbial species. Combination riboPOOLs containing up to four existing riboPOOLs enable rRNA depletion from samples containing multiple species such as pathogen infected tissue or metagenomic samples (e.g. soil, sludge).



Hybridization

riboPOOLs are resuspended and hybridized to DNA-free total RNA.



Capture & Removal

Streptavidin-coated magnetic beads separate riboPOOLbound rRNAs.



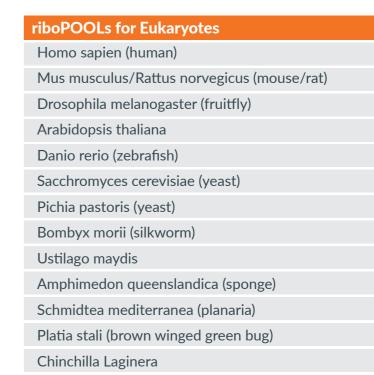




Remaining relevant RNAs are purified by ethanol, silica column or SPRI beads prior to downstream analysis.

*Time required dependant on clean-up method

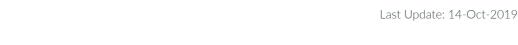
List of Ready-Made riboPOOLs



riboPOOLs for Abundant RNAs

Human Globin (mRNA)





BIOTECH

Special Applications with riboPOOL

Metatranscriptomics with Pan-Prokaryote riboPOOL

The Pan-Prokaryote riboPOOL, designed for universal microbial rRNA depletion, has been applied for metagenomic samples. Depletion efficiency however may vary between species and sample quality.

Depletion of abundant RNAs

Aside from rRNA, depletion of abundant tissue-specific mRNAs (e.g. globin) or other abundant RNAs found in sequencing data can be performed with riboPOOLs.

Ribosome profiling

riboPOOLs have been applied for ribosome profiling samples with depletion efficiencies of ~50-80%. Please contact us for more information.

Depletion of rRNAs from multiple species with Combination riboPOOLs

Combination riboPOOLs are mixtures of up to four riboPOOLs combined in custom ratios for use in samples containing multiple species. Please refer to **riboPOOL Application Note** on website for data on the Human/Mouse/Pan-Prokaryote Combination riboPOOL.

Product Properties

- 3'-biotinylated, single-stranded DNA
- HPLC-purified
- Provided as probes alone with nuclease-free water **OR** in kits that include buffers, streptavidin-magnetic beads and ethanol precipitation reagents. A protocol is provided for either format.
- Delivered as freeze-dried powder at room-temperature.
- Upon resuspension, riboPOOLs are stable up to 6 months when stored at or below -20°C. Store in aliquots to avoid multiple freeze-thaw cycles.

Ordering Information

riboPOOLs can be ordered through our **webshop** at www.sitoolsbiotech.com, by **email** at info@sitools.de or via your **local distributor** (www.sitoolsbiotech.com/distributors.php)

Please specify full species name, format (probes/kit), and reaction scale required.

Products available	Scales available
riboPOOL probes	20, 50 or 100 reactions
Combination riboPOOL probes	20, 50 or 100 reactions
Human/Mouse riboPOOL kit	6 (trial), 12, 24 or 96 reactions
Pan-Prokaryote/Globin/Other Species riboPOOL kit	6 (trial), 12, 24 or 96 reactions
Combination riboPOOL kit	12, 24 or 96 reactions
Custom riboPOOL set up service	(one-time purchase required for new species)

Contact us

info@sitools.de / +49 89 12501 4800

Learn more

www.sitoolsbiotech.com/riboPOOLs.php



