

## proFIRE® Amine Coupling Kit 2 for His-tagged Proteins (> 5 kDa)

Functionalization of DNA via amines (-NH<sub>2</sub>)

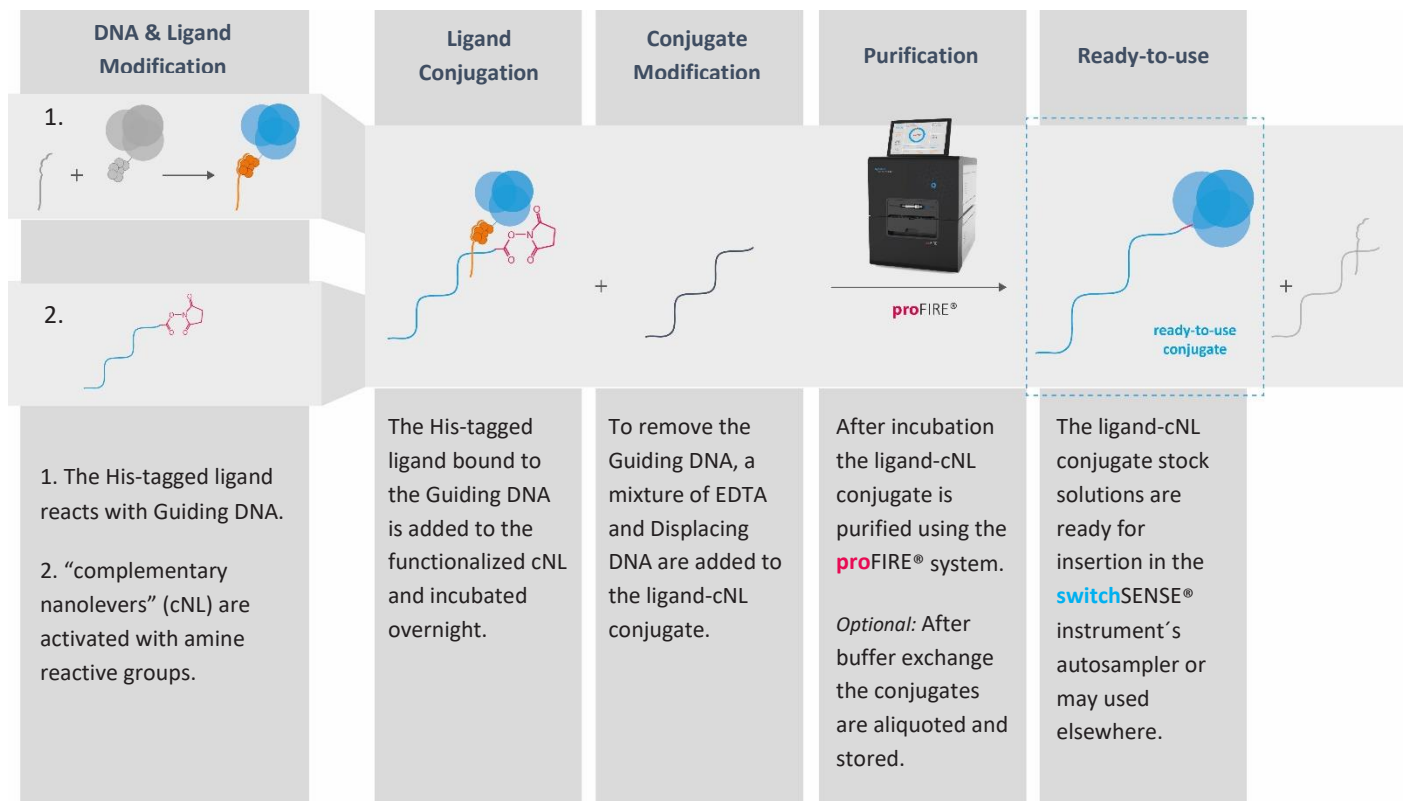


### Key Features

- Coupling of biomolecules with primary amines (e.g. NH<sub>2</sub>-terminus, lysines) and a His-tag to DNA in a reaction tube
- Oriented coupling in the proximity of the His-tag
- Convenient standard chemistry
- Applicable for proteins (and peptides) (MW > 5 kDa)
- Compatible with all **switchSENSE®** *Multi-purpose biochips* carrying sequence B48
- Suitable for parallel measurements via DNA encoded addressing
- Coupling of multiple ligands can be performed simultaneously
- Yields >95 % pure ligand-DNA conjugate with controlled quality of your product
- Includes reagents for three individual conjugation reactions (approx. 10-30 regenerations each; up to 500)
- Compatible with automated standard regeneration process
- **proFIRE®** purification for pure ligand-DNA conjugates

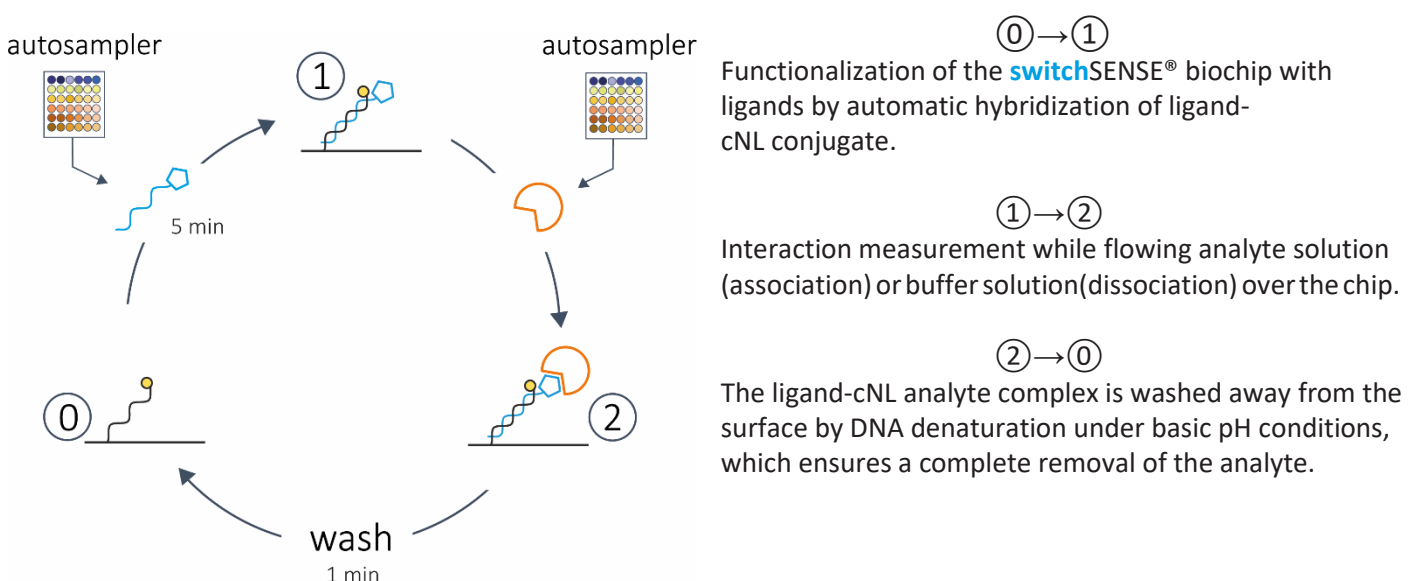
## Workflow Overview

### Conjugation Workflow



Timeline: Hands on time < 1 h | Incubation 12-16 h

## Measurement Workflow with ligand-cNL conjugates



## Product Description

Order Number **PF-NH2-2-B48**

TABLE 1 | Contents and Storage Information

Material	Cap	Amount	Storage	Comment
cNL-B48-08	blue	3 x	-20°C	
cNL-A48 (1 µM)	yellow	500 µL	-20°C	
Guiding DNA	orange	3 x 18 µL	-20°C	
Displacing DNA	purple	3 x 6 µL	-20°C	
Buffer H	trans- parent	3 x 1.8 mL	-20°C	
Loading solution (500 µM)	trans- parent	50 µL	-20°C	
EDTA solution (500 mM)	trans- parent	100 µL	-20°C	
Buffer PE40 (10 mM Na <sub>2</sub> HPO <sub>4</sub> /NaH <sub>2</sub> PO <sub>4</sub> pH 7.4, 40 mM NaCl, 0.05 % Tween, 50 µM EDTA, 50 µM EGTA)	trans- parent	3 x 1.5 mL	-20°C	
ddH <sub>2</sub> O	trans- parent	1.5 mL	-20°C	
Crosslinker	green	3 x	-20°C	
Purification spin column	red	6 x	2-8°C	
2.0 mL Reaction tubes for Purification spin column		6 x	r.t.	
Centrifugal filter unit (3 kDa MWCO) <sup>1</sup>		3 x	r.t.	
Centrifugation collection tube		6 x	r.t.	

For *in vitro* use only.

Please check date of expiry on the kit. Products are shipped at ambient temperature or frozen. The kit contains reagents sufficient for 3 conjugations of approx. 20-150 µg biomolecule each. The resin slurry of the Purification spin column contains 0.02 % sodium azide. The Loading solution contains 500 µM NiCl<sub>2</sub>.

<sup>1</sup> For conjugation of proteins with a molecular weight higher than 20 kDa: Centrifugal filter units with a MWCO of 10 kDa can be ordered for a faster concentration process (Please see page 10 for order number).

## Additional Materials Required

TABLE 2 | Additional Materials.

Material	Comment
Benchtop microcentrifuge	Capable between 1,000 x g and 13,000 x g
Vortexer	
1.5 mL reaction tubes	
UV-Vis spectroscopy (e.g. Nanodrop)	Concentration determination of the conjugate

All necessary solutions and buffers are included in the kit.

## Important Notes

- Do not use any buffer containing primary amines (i.e. Tris, glycine) during conjugation process.
- Dithiothreitol (DTT) can be used up to 1 mM during the conjugation process. Do not use 2-Mercaptoethanol or other thiol-based reducing agents during conjugation process. If a reducing agent is necessary, TCEP is recommended up to 1 mM. For reducing agents during interaction measurement, please refer to the [switchSENSE®](#) compatibility sheet (application area on [www.dynamic-biosensors.com/switchsense](http://www.dynamic-biosensors.com/switchsense)).
- Avoid using partially purified protein samples or protein samples containing carriers (e.g. BSA).
- To get highest reaction yields, the ligand should be dissolved in Buffer H. Buffer exchange is recommended prior to conjugation process. The ligand solution must not contain EDTA.
- Before you begin, briefly centrifuge all tubes to ensure that all material is at the bottom of the tubes.
- For molecules with a molecular weight around or lower than 5 kDa, special care during purification process shall be taken. A few peptides may not give a proper purification using the provided IEX column. For more information please email [support@dynamic-biosensors.com](mailto:support@dynamic-biosensors.com).

## Conjugation of a Biomolecule to a DNA-Nanolever in a Reaction Tube

Please read the entire protocol before starting and **perform conjugation without interruption**.

TIP: the protocol can be performed simultaneously for multiple coupling reactions.

### Spin Column Equilibration

1. Equilibrate **two** purification spin columns for one coupling reaction:
  - a. Remove column's bottom closure and loosen cap (do not remove cap).
  - b. Place column in a 2.0 mL reaction tube.
  - c. Centrifuge at 1,500 × g for 1 minute to remove the storage solution.
  - d. Add **400 µL of Buffer H** on top of column's resin bed. Centrifuge at 1,500 × g for 1 minute to remove buffer.
  - e. Repeat step d once, discard buffer from the reaction tube. The Purification spin column should be in a dry state now.

### Ligand Modification

2. Prepare **20-150 µg** of your His-tagged ligand (concentration approx. 0.5 – 25 mg/mL). For optimal conditions dissolve / dilute ligand in Buffer H.  
**EXAMPLE:** Adjust protein concentration to 2 - 5 mg/mL and use 10 - 25 µL for conjugation.  
*IMPORTANT: Be sure that the storage buffer of the ligand does not contain any primary amines, e.g. Tris buffers, glycine or EDTA (please see page 4, Important Notes).*
3. Add **18 µL Guiding DNA** and **5 µL Loading solution** to your His-tagged ligand. Mix the reaction by pipetting up and down and let it react at room temperature for **15 minutes**.  
**IMPORTANT:** During this reaction time continue with DNA-Nanolever Modification.

### DNA-Nanolever Modification

4. Dissolve cNL-B48-08 (blue cap) in **40 µL Buffer H** prior to use and vortex until solids are completely dissolved and spin down shortly.
5. Dissolve the crosslinker (green cap) by adding **100 µL ddH<sub>2</sub>O** and vortex until solids are completely dissolved and spin down shortly. **IMPORTANT:** Always use fresh compounds.
6. Add **2.5 µL** of the freshly prepared linker solution to one nanolever aliquot (cNL-B48-08, blue cap). Discard the remaining linker solution from step 5.
7. Vortex the reactants for 10 sec, spin down and incubate them for **5 minutes** at room temperature.  
**IMPORTANT:** Do not exceed incubation time as the reaction yield will decrease.

8. Sample loading
  - a. Place columns from step 1 in new 1.5 mL reaction tubes.
  - b. Remove cap of spin column number 1 and apply the sample from step 7 to the top of the resin bed.
  - c. Centrifuge at 1,500 x g for 2 min to collect the sample (flow-through).  
Discard Purification spin column after use.
  - d. Remove cap of spin column number 2 and apply the sample from step c on top of the resin bed.
  - e. Centrifuge at 1,500 x g for 2 min to collect the sample (flow-through).  
Discard Purification spin column after use.

### Ligand Conjugation

9. Add the ligand mix from step 3 to the sample from step 8. Mix the reaction by pipetting up and down and let it react at room temperature for **at least 1 hour (or overnight)**.

**IMPORTANT:** Do not vortex. For higher yields, a further incubation overnight (e.g. at 2-8 °C) is recommended.

### Conjugate Modification

10. Add **6 µL of Displacing DNA** and **10 µL of EDTA solution** to your conjugate from step 9. Mix the reaction by pipetting up and down and let it react at room temperature for **5 minutes**.

**IMPORTANT:** Please make sure, that the **proFIRE®** is ready to use.

### proFIRE® Purification

Please refer to the **proFIRE®** User Manual.

11. Perform a purification using the **proFIRE®**. Please make sure that the sample volume is 160 µL.
  - If the volume is less than 160 µL, add Buffer H.
  - If it exceeds 160 µL, please perform two subsequent runs.
12. Use the Data Viewer software of the **proFIRE®** to identify which fractions contains pure conjugate. On page 8 (Additional Information section: **proFIRE®** Purification of a Ligand-cNL Conjugate) an example chromatogram is shown.
13. Take the recommended fractions out of the fraction collector.

## Buffer Exchange

**NOTE:** If you don't want to perform a **switchSENSE®** measurement, the following protocol can be used as a guideline to conduct a buffer exchange with your desired buffer.

14. Add **500 µL** of the first fraction containing the ligand-DNA conjugate from the **proFIRE®** to the centrifugal filter unit.  
Centrifuge at 13,000 x g (up to 14,000 x g) for **10 minutes** and discard flow-through.
  - b. Add the remaining fractions in the same filter unit and repeat the centrifugation step in order to collect all samples in one tube (Please check on page 9: Additional information for the right use of centrifugal filter unit).
  - c. Add **350 µL of PE40** (or TE40, HE40) buffer and centrifuge at 13,000 x g for **10 minutes**. Discard the flow-through again.  
  
If the protein is not stable in **PE40** (or TE40, HE40), please check buffer compatibility with the **switchSENSE®** compatibility sheet (Application area on [www.dynamic-biosensors.com/switchsense](http://www.dynamic-biosensors.com/switchsense)).
  - d. Add **350 µL of PE40** (or TE40, HE40) buffer and centrifuge at 13,000 x g for **15 minutes**. Discard the flow-through again.
  - e. To recover the ligand-DNA conjugate, place the centrifugal filter unit upside down in a **new** centrifugal collection tube (provided in the kit).  
Spin for **2 minutes** at 1,000 x g to transfer the sample to the tube.

## Optional: Concentration

15. Check ligand-DNA conjugate concentration after buffer exchange by using absorbance at 260 nm and the following equation:  
$$c (\text{ligand-DNA conjugate}) = A_{260 \text{ nm}} / (490,000 \text{ L mol}^{-1} \text{ cm}^{-1} * d)$$

d = optical path length  
(usually d = 1 cm, please check photometer manual for further information).

## Aliquots and Storage

16. Adjust the concentration to **200 nM – 1 µM** with **PE40** (or TE40, HE40) buffer (including up to 10 % glycerol if needed) and prepare **20 µL** aliquots.
17. Store between 8 °C and -86 °C as desired.

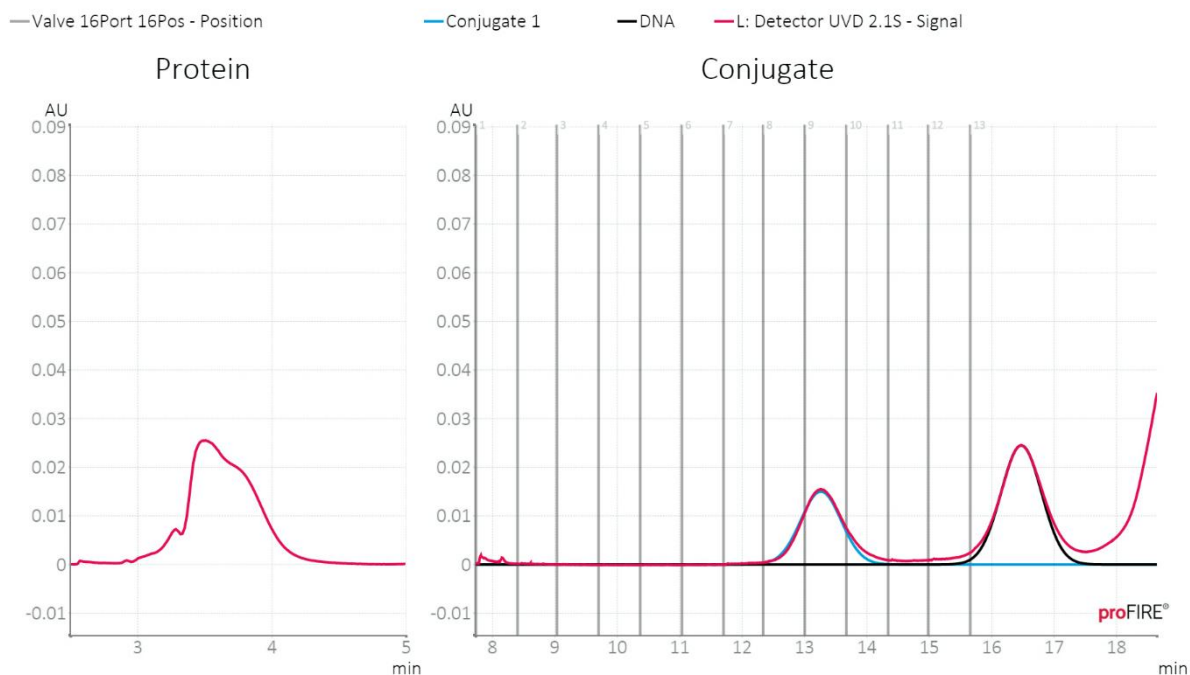
### IMPORTANT:

Please add prior to an interaction measurement the appropriate reference cNL (e.g. cNL-A48) to the conjugate solution.

## Additional Information

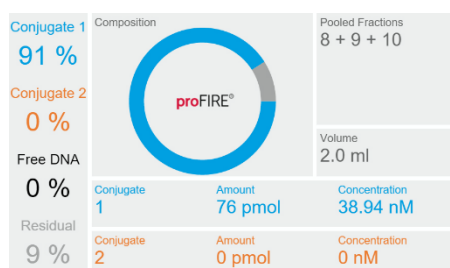
### proFIRE® Purification of a Ligand-cNL Conjugate

- To ensure highest quality of your results, we recommend to use highly purified protein-DNA conjugates. Therefore ligand-cNL-B48 conjugates have to be purified by ion exchange chromatography. Furthermore, this quality control step gives you additional useful information about your sample purity.
- We recommend using the provided proFIRE® system and the proFIRE® column. For an example chromatogram, see figure below.  
Prepare 250 mL Buffer A (50 mM Na<sub>2</sub>HPO<sub>4</sub>/ NaH<sub>2</sub>PO<sub>4</sub> pH 7.2 and 150 mM NaCl)<sup>1</sup> and 250 mL Buffer B (50 mM Na<sub>2</sub>HPO<sub>4</sub>/NaH<sub>2</sub>PO<sub>4</sub> pH 7.2 and 1 M NaCl)<sup>1</sup>.
- Collect the ligand-cNL conjugate fraction (here: 8-10), concentrate the conjugate and exchange buffer with your desired buffer using a Centrifugal filter unit, as described on page 9.



#### proFIRE® chromatogram of a ligand-cNL-B48 conjugate purification.

Used buffers: Buffer A: 50 mM Na<sub>2</sub>HPO<sub>4</sub>/NaH<sub>2</sub>PO<sub>4</sub> pH 7.2, 150 mM NaCl/ Buffer B: 50 mM Na<sub>2</sub>HPO<sub>4</sub>/NaH<sub>2</sub>PO<sub>4</sub> pH 7.2, 1 M NaCl. Column: proFIRE® column. Flow: 1 mL/min. Used program: DNA length 48, Type 1.

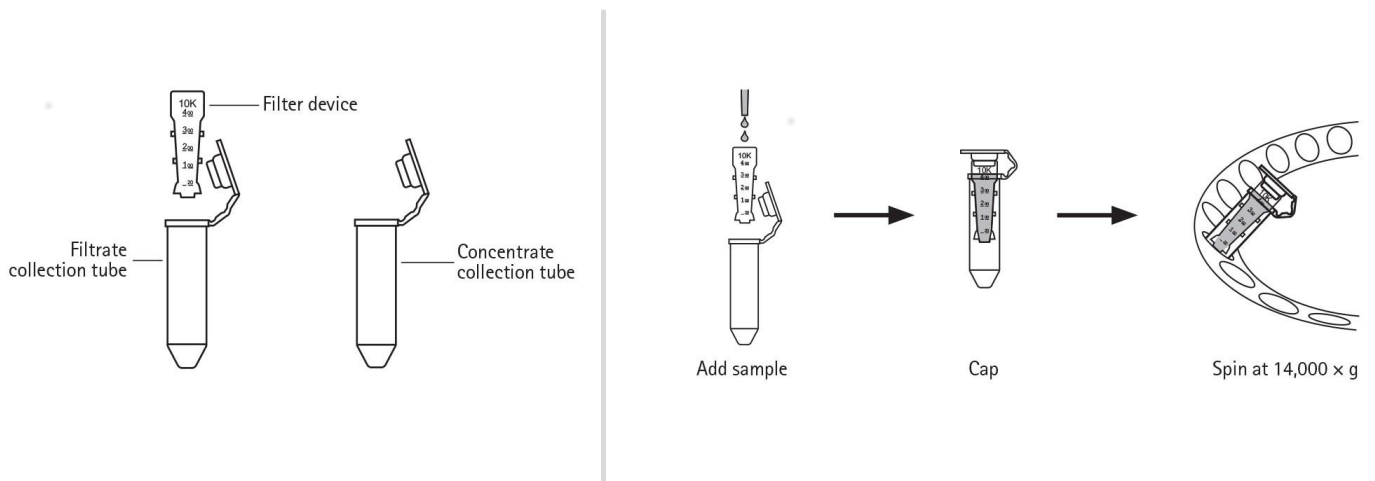


proFIRE® report gives you an automated conjugate analysis with the most important facts about your conjugate, e.g. concentration, amount and purity.

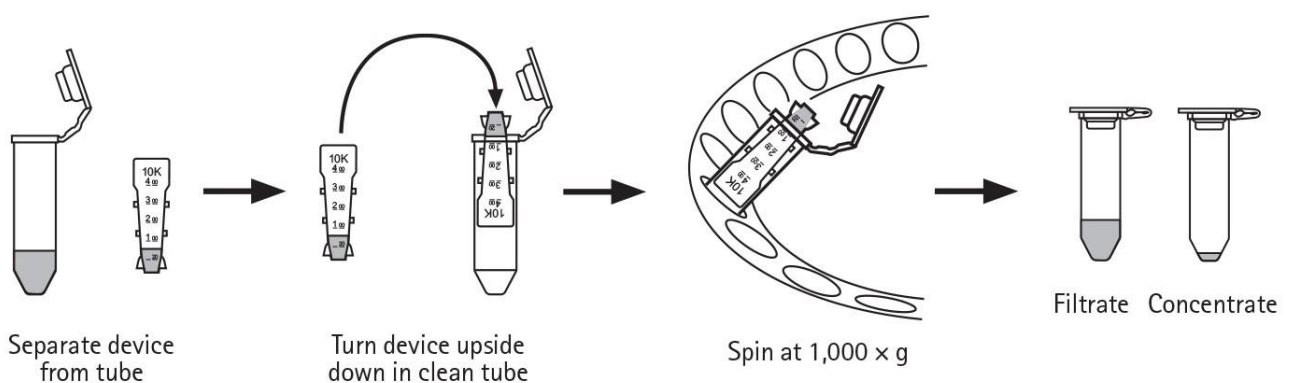
<sup>1</sup> See page 10 for order no.



## Buffer Exchange and Concentration with Centrifugal Filter Units



1. Take one centrifugal filter unit, add the appropriate volume of buffer in the filter device, and cap it.
2. Place capped filter device into the centrifuge rotor, aligning the cap strap toward the center of the rotor; counterbalance with a similar device.
3. Spin the device at 13,000 x g (or 14,000 x g) for the given time.
4. Remove the flowthrough and repeat the steps 1-3.
5. Remove the assembled device from the centrifuge and separate the filter device from the microcentrifuge tube.
6. To recover the conjugate, place the filter device upside down in a clean centrifugal tube, aligning open cap towards the center of the rotor; counterbalance with a similar device. Spin for 2 minutes at 1,000 x g to transfer the sample from the device to the tube.



## Useful Order Numbers

TABLE 3 | Order Numbers.

Product name	Order Number
proFIRE® Amine Coupling Kit 1 for proteins (>5 kDa); sufficient for 5 conjugation series	PF-NH2-1
proFIRE® Thiol Coupling Kit 1 for proteins (>5 kDa); sufficient for 5 conjugation series	PF-SH-1
Centrifugal filter unit (3 kDa MWCO), 5 pcs.	CF-003-5
Centrifugal filter unit (10 kDa MWCO), 5 pcs.	CF-010-5
proFIRE® column	PF-CC-1
10x proFIRE® Buffer A (50 mL)	PF-BU-A-10
5x proFIRE® Buffer B (50 mL)	PF-BU-B-5

**My Notes**

## Contact

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