# **Custom Amine Coupling Kit**

# for Proteins (> 5 kDa)

Functionalization of 48mer nanolevers via amines (-NH<sub>2</sub>)

# **Key Features**

- Coupling of biomolecules with primary amines (e.g. NH<sub>2</sub>-terminus, lysines) to modified nanolevers in a reaction tube
- Convenient standard chemistry
- Applicable for proteins (and peptides) (MW > 5 kDa)
- Compatible with all switchSENSE<sup>®</sup>
  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

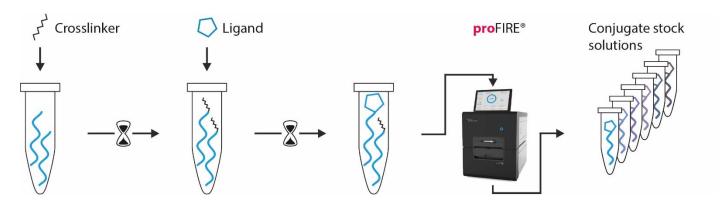
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- Includes reagents for five individual conjugation reactions (approx. 10-50 regenerations each; up to 500)
- Compatible with automated standard regeneration process
- proFIRE<sup>®</sup> purification for pure ligand-DNA conjugates

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# **Workflow Overview**

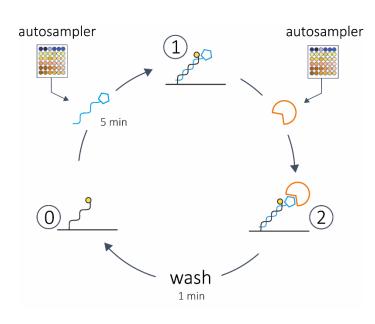
### 3-Step Conjugation Workflow (in-vitro)



1. DNA Modification	2. Ligand Conjugation	3. Purification	4. Ready-to-use
"complementary nanolevers" (cNL) are activated with amine reactive groups.	After incubation the excess linker is removed by a spin column. The protein/peptide (ligand) is added to the functionalized cNL and incubated for at least 1 h.	The ligand-cNL conjugate is purified using the <b>pro</b> FIRE® system. After buffer exchange the conjugates are aliquoted and stored.	The ligand-cNL conjugate stock solutions are ready for insertion in the DRX autosampler.

Time line: Hands on time < 1 h | Incubation ~ 2 h | Total ~ 3 h

# Measurement Workflow with ligand-cNL conjugates



#### (0)→(1)

Functionalization of the switchSENSE<sup>®</sup> biochip with ligands by automatic hybridization of ligand-cNL conjugate.

#### ①→②

Interaction measurement while flowing analyte solution (association) or buffer solution(dissociation) over the chip.

# (2)→(0)

The ligand-cNL analyte complex is washed away from the surface by DNA denaturation under basic pH conditions, which ensures a complete removal of the analyte.



# **Product Description**

Order Number

#### CK-NH2-CUS-1-B48 (nanolever sequence B48)

TABLE 1 | Contents and Storage Information

Material	Сар	Amount	Storage	Comment
cNL-B48-DBCO	blue	5 x	-20°C	
cNL-A48 (1 μM)	yellow	500 μL	-20°C	
Buffer C (50 mM Na <sub>2</sub> HPO <sub>4</sub> /NaH <sub>2</sub> PO <sub>4</sub> pH 8.0, 150 mM NaCl)	trans- parent	6 x 1.8 mL	-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	5 x 1.5 mL	-20°C	
ddH <sub>2</sub> O	trans- parent	1.5 mL	-20°C	
Crosslinker	brown	5 x	-20°C	
Purification spin column	red	10 x	2-8°C	
2.0 mL Reaction tubes for Purification spin column		10 x	r.t.	
Centrifugal filter unit (3 kDa MWCO) <sup>1</sup>		5 x	r.t.	
Centrifugation collection tube		10 x	r.t.	

For *in vitro* use only.

Please check date of expiry on the kit. Products are shipped at ambient temperature. The kit contains reagents sufficient for 5 conjugations of approx. 50-200  $\mu$ g biomolecule each. The resin slurry of the Purification spin column contains 0.02 % sodium azide.

<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<sup>®</sup> compatibility sheet (application area on 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 C. Buffer exchange is recommended prior to conjugation process<sup>1</sup>.
- Before you begin, briefly centrifuge all tubes with blue, brown, yellow and transparent caps 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 chromatographic column. For more information please email support@dynamic-biosensors.com.
- If the pl of the protein is < 6, it might be necessary to use CK-NH2-7-B48. For more information, please email support@dynamic-biosensors.com.

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



### 3-Step Conjugation of a Biomolecule to a 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.

#### Nanolever Modification

- Dissolve cNL-B48-DBCO (blue cap) in 40 μL Buffer C prior to use and vortex until solids are completely dissolved and spin down shortly.
- Dissolve the crosslinker (brown 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.
- 3. Add **10 μL** of the freshly prepared linker solution to one nanolever aliquot (cNL-B48-DBCO, blue cap). Discard the remaining linker solution from step 2.
- 4. Vortex the reactants for 10 sec, spin down and incubate them for **20 minutes** at room temperature. **IMPORTANT**: Do not exceed incubation time as the reaction yield will decrease.
- 5. Equilibrate in the meantime **two** purification spin columns (red cap) 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 Buffer C** 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.
- 6. Sample loading
  - a. Place columns from step 5 in new 1.5 mL reaction tubes.
  - b. Remove cap of spin column number 1 and apply the sample from step 4 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.



#### II Ligand Conjugation

Add approx. 100 μg (up to 200 μg) of the ligand (concentration approx. 0.5 – 50 mg/mL) to the sample from step 6. For optimal conditions use a volume of approx. 50 μL.
 EXAMPLE: Adjust protein concentration to 2 mg/mL and use 50 μL for conjugation.

*IMPORTANT: Be sure that the storage buffer of the ligand does not contain any primary amines, e.g. TRIS buffers, glycine (please see page 4, Important Notes).* 

#### Mix the reaction by pipetting up and down and let it react at room temperature for at least 1 hour.

**IMPORTANT**: Do not vortex. If necessary, the reaction can be carried out at 4 °C with a longer reaction time (e.g. overnight).

#### III **pro**FIRE<sup>®</sup> Purification

Please refer to the proFIRE® User Manual.

- 9. Perform a purification using the **pro**FIRE<sup>®</sup>. Please make sure that the sample volume is 160 μL.
  - $\circ~$  If the volume is less than 160  $\mu\text{L}$  , add Buffer C.
  - $\circ~$  If it exceeds 160  $\mu\text{L}$  , please perform two subsequent runs.
- 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.
- 11. Take the recommended fractions out of the fraction collector.

#### IV Buffer Exchange

- Add **500 μL** of the first fraction containing the ligand-DNA conjugate from the proFIRE<sup>®</sup> 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 **switch**SENSE<sup>®</sup> compatibility sheet (Application area on **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

13. Check ligand-DNA conjugate concentration after buffer exchange by using absorbance at 260 nm and the following equation:

c (ligand-DNA conjugate)=  $A_{260 \text{ nm}}/(490,000 \text{ L mol}^{-1} \text{ cm}^{-1} \text{ * d})$ 

d = optical path length

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

#### VI Aliquots and Storage

- 14. Adjust the concentration to **200 nM 1 \muM** with **PE40** (or TE40, HE40) buffer (including up to 10 % glycerol if needed) and prepare **20 \muL** aliquots.
- 15. 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.



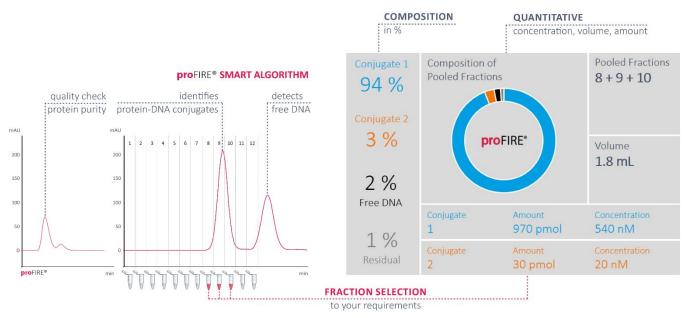
# **Additional Information**

### I proFIRE® Purification of a Ligand-cNLConjugate

- 1. To perform a size analysis during the measurement it has to be ensured that there is no free DNA on the chip and only 1:1 conjugates are present. Therefore crude ligand-cNL-B48 conjugates have to be purified by ion exchange chromatography. This quality control step gives you additional useful information about your sample purity.
- We recommend using the provided proFIRE® system equipped with an ion exchange column.
  For an example chromatogram, see figure below.

Prepare 250 mL Buffer A (50 mM  $Na_2HPO_4/NaH_2PO_4$  pH 7.2 and 150 mM NaCl)<sup>1</sup> and 250 mL Buffer B (50 mM  $Na_2HPO_4/NaH_2PO_4$  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 buffer of choice using a Centrifugal filter unit, as described in section II (Additional information, page 9).



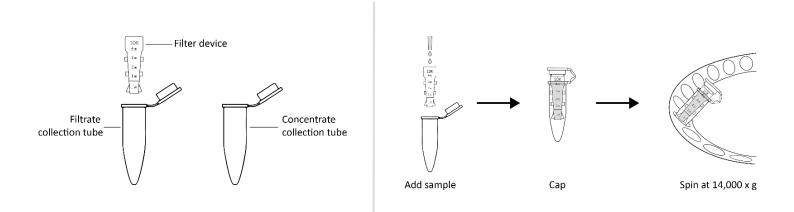
#### proFIRE® chromatogram of a ligand-cNL-B48 conjugate purification.

*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:* Chromatographic column. *Flow:* 1 mL/min. *Used program:* DNA length **48**, Type **1**.

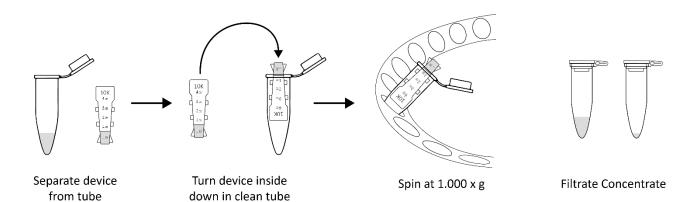
<sup>&</sup>lt;sup>1</sup> See page 10 for order no.

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#### II 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
Amine coupling kit 1 for proteins (>5 kDa); cNL-B48 and NHS modifier, sufficient for 5 conjugation series	CK-NH2-1-B48
Amine coupling kit 2 for proteins (>5 kDa); cNL-B48 and NHS modifier, sufficient for 5 conjugation series (spin column purification)	CK-NH2-2-B48
Amine coupling kit 7 for proteins (>5 kDa) with a low pl (< 6); cNL-B48 and NHS modifier, sufficient for 5 conjugation series	CK-NH2-7-B48
Centrifugal filter unit (3 kDa MWCO), 5 pcs.	CF-003-5
Centrifugal filter unit (10 kDa MWCO), 5 pcs.	CF-010-5
Chromatographic column	TB-CC-1-1
10x Buffer A pH 7.2 (50 mL of: 500 mM Na2HPO4/NaH2PO4, 1.5 M NaCl) Yields 0.5 L of: 50 mM Na2HPO4/NaH2PO4, 150 mM NaCl	BU-P-150-10
5x Buffer B pH 7.2 (50 mL of: 250 mM Na <sub>2</sub> HPO <sub>4</sub> /NaH <sub>2</sub> PO <sub>4</sub> , 5 M NaCl) Yields 0.25 L of: 50 mM Na <sub>2</sub> HPO <sub>4</sub> /NaH <sub>2</sub> PO <sub>4</sub> , 1 M NaCl	BU-P-1000-5
1x Buffer C pH 8.0 (12 mL of: 50 mM Na2HPO4/NaH2PO4, 150 mM NaCl)	BU-C-150-1



**My Notes** 



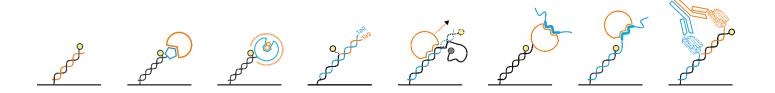
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switchSENSE<sup>®</sup> is a proprietary measurement technology by Dynamic Biosensors GmbH. Instruments and biochips are engineered and manufactured in Germany.

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