dynamic BIOSENSORS

Amine coupling kit 7 for proteins (> 5 kDa)

Functionalization of 96mer nanolevers via amines (-NH₂)

Key Features

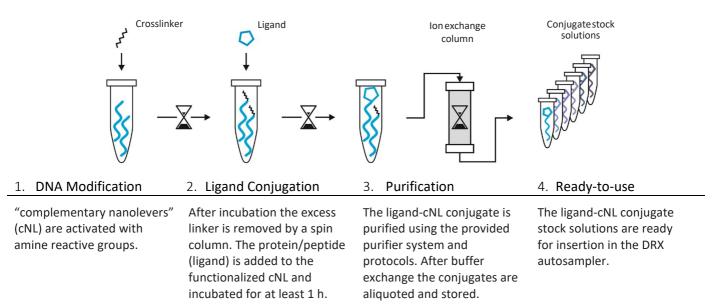
- Coupling of biomolecules with primary amines (e.g. NH₂-terminus, lysines) to modified nanolevers in a reaction tube
- Convenient standard chemistry
- Applicable for proteins (and peptides) (MW > 5 kDa) with a low pl (pl < 6)
- Longer nanolever for larger complexes
- Compatible with all <u>switchSENSE®</u> Multi-purpose biochips carrying sequence B96
- 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 yourproduct
- Includes reagents for five individual conjugation reactions (approx. 10-30 regenerations each; up to 200)
- Compatible with automated standard regeneration process



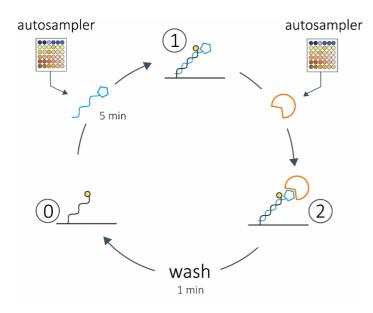
Workflow Overview

3-Step Conjugation Workflow (in-vitro)



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

Measurement Workflow with ligand-cNL conjugates



(0)→(1)

Functionalization of the **switch**SENSE[®] 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-NH2-7-B96 (nanolever sequence B96)

TABLE 1 | Contents and storage information

Material	Сар	Amount	Storage	Comment
cNL-B96-01	blue	5 x	-20°C	
cNL-A96 (1 μM)	yellow	500 μL	-20°C	
Buffer M (50 mM MES pH 6.5, 150 mM NaCl)	trans- parent	5 x 1.8 mL	-20°C	
Buffer A (50 mM Na ₂ HPO ₄ /NaH ₂ PO ₄ pH 7.2, 150 mM NaCl)	trans- parent	1.8 mL	-20°C	
Buffer PE40 (10 mM Na ₂ HPO ₄ /NaH ₂ PO ₄ 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 ₂ O	trans- parent	1.5 mL	-20°C	
Crosslinker	green	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 (10 kDa MWCO)		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 μ g biomolecule each. The resin slurry of the Purification spin column contains 0.02 % sodium azide.



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).
- 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 M. Buffer exchange is recommended prior to conjugation process¹.
- Before you begin, briefly centrifuge all tubes with blue, green, 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 IEX column. For more information please email support@dynamic-biosensors.com.

¹ See page 9 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

- 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 M** 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.
- 2. Dissolve cNL-B96-01 (blue cap) in **40 µL Buffer A** prior to use and vortex until solids are completely dissolved and spin down shortly.
- Dissolve the crosslinker (green cap) by adding 100 μL ddH₂O and vortex until solids are completely dissolved and spin down shortly. IMPORTANT: Always use fresh compounds.
- 4. Add **10 μL** of the freshly prepared linker solution to one nanolever aliquot (cNL-B96-01, blue cap). Discard the remaining linker solution from step 3.
- 5. 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.

- 6. 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 5 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. 1 – 50 mg/mL) to the sample from step 6. For optimal conditions use a volume of approx. 10 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).

IIIPurification & Concentration

- Perform a purification reaction using the ÄKTA Start with the provided column and program. If the volume exceeds 150 μL, please perform two subsequent runs. Please refer to the Manual Instruction ÄKTA Start.
- 10. Take the fractions with the ligand-cNL-B96 conjugate out of the fraction collector. On page 7 (*Additional Information section:* Purification of a ligand-cNL conjugate) an example chromatogram is shown.
- a. Add the first **500 μL** fraction from the ÄKTA Start 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 8: 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** and discard 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).
 - d. Add **350 μL of PE40** (or TE40, HE40) buffer and centrifuge at 13,000 x g for **15 minutes** and discard flow-through again.
 - e. To recover the 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.
- 12. Determine conjugate concentration by using absorbance at 260 nm and the following

equation: $c (conjugate) = A_{260 nm}/(1,044,000 L mol^{-1} cm^{-1} * d)$ d = optical path length (usually d = 1 cm, please check photometer manual for further information).

- 13. Adjust the concentration to **200nM 1μM** with **PE40** (or TE40, HE40) buffer (including up to 10 % glycerol if needed) and prepare **20 μL** aliquots.
- 14. Store between 8 °C and -86 °C as desired.
- 15. Prior to an interaction measurement: Please add the appropriate reference cNL (e.g. cNL-A96) to the conjugate solution.



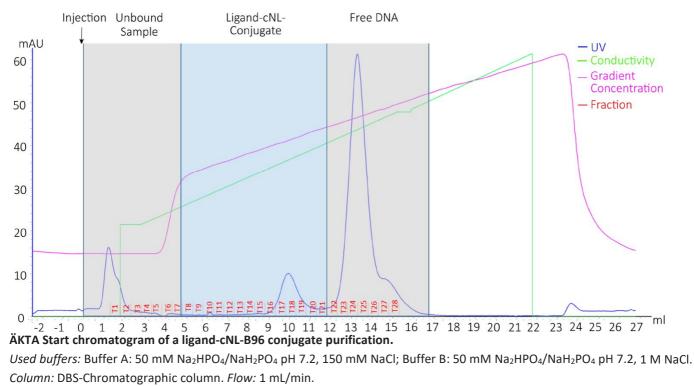
Additional Information

I Purification of a Ligand-cNLConjugate

- To perform a kinetic analysis 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-B96 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 ÄKTA Start equipped with an anion exchange column. For an example chromatogram see figure below. Prepare 250 mL Buffer A (50 mM Na₂HPO₄/ NaH₂PO₄ pH 7.2 and 150 mM NaCl)¹ and 250 mL Buffer B (50 mM Na₂HPO₄/NaH₂PO₄ pH 7.2 and 1 M NaCl)¹.
- Collect the ligand-cNL conjugate fraction (here: T16-T20), concentrate the conjugate and exchange buffer with your buffer of choice using a Centrifugal filter unit, as described in section II (Additional information, page 8). If you are not sure about your conjugate collect the fractions T6-T15 or email the chromatogram to:

support@dynamic-biosensors.com

Usually, free DNA elutes in fraction T22-T28 and should not be used.

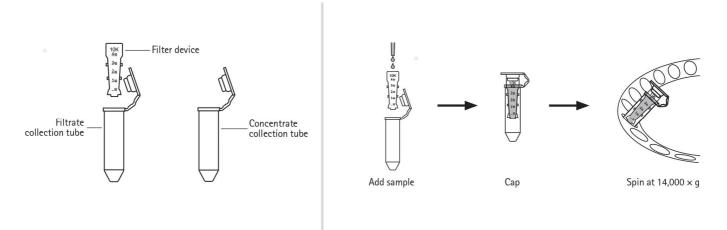


Used program: DBS_Purification run 96mer.

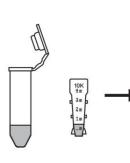
¹ See page 9 for order no.

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II Buffer Exchange and Concentration with Centrifugal Filter Units

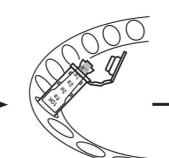


- 1. Take one centrifugal filter unit, add the appropriate volume of buffer in the filter device, and cap it.
- 2. Place capped filter device into the centrifuge rotor, aligning the cap strap toward the center of the rotor; counterbalance with a similar device.
- 3. Spin the device at 13,000 x g (or 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.



Separate device from tube

Turn device upside down in clean tube



Spin at 1,000 x g

Filtrate Concentrate

CK-NH2-7-B96_v6.0



Useful Order Numbers

TABLE 3 | Order numbers.

Product Name	Order Number	
Amine coupling kit 1 for proteins (>5 kDa); cNL-A96 and NHS modifier, sufficient for 5 conjugation series	CK-NH2-1-A96	
Amine coupling kit 2 for proteins (>5 kDa); cNL-B96 and NHS modifier, sufficient for 5 conjugation series (spin column purification)	СК-NH2-2-B96	
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 Na2HPO4/NaH2PO4, 5 M NaCl) Yields 0.25 L of: 50 mM Na2HPO4/NaH2PO4, 1 M NaCl	BU-P-1000-5	
1x Buffer M pH 6.5 (12 mL of: 50 mM MES, 150 mM NaCl)	BU-M-150-1	



My Notes



My Notes

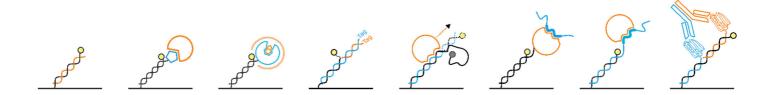


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switchSENSE[®] is a proprietary measurement technology by Dynamic Biosensors GmbH. Instruments and biochips are engineered and manufactured in Germany.

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