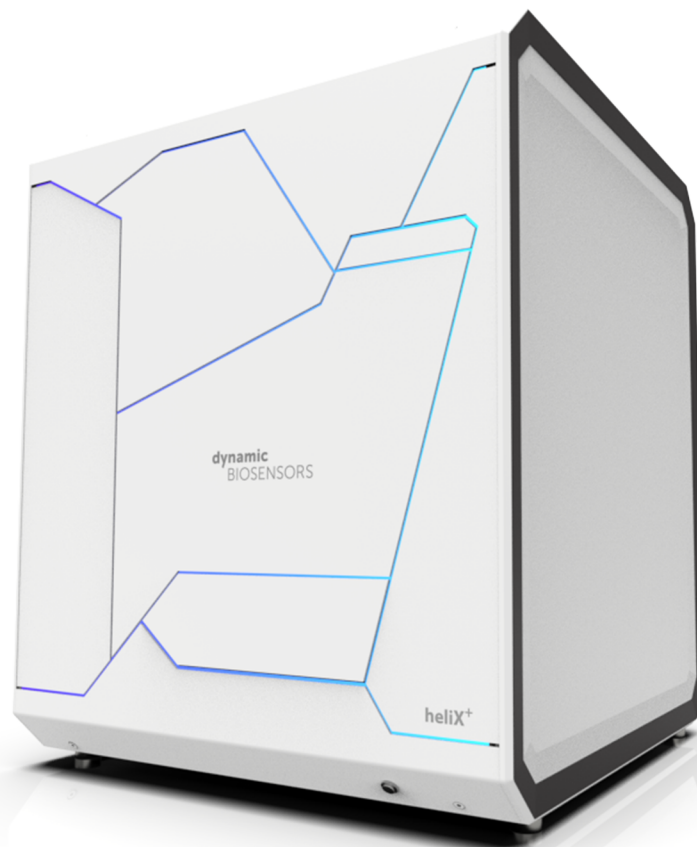


NEW Y-STRUCTURE KIT 2

for proximity binding assay - **Spot 2**

Dynamic Biosensors GmbH
HK-NYS-2 v1.0



Key Features

- Ideal for studying **ternary complex formation** upon binding of bispecific small molecules (e.g., **PROTACs**, molecular glues), homo-dimerization and bispecific antibodies with weak affinities.
- The new optimized **Y-structure** design allows higher **FRET** sensitivity.
- Compatible with **heliX[®] Adapter Chip**.
- The **Y-structure Red Adapter** strand (specific to Spot 2) carries a moderately hydrophilic red dye (**Ra**) with a single positive net charge.
- The **Y-structure Green Adapter** strand carries a hydrophilic green dye (**Ga**) with a single negative net charge.
- Green and red dyes can detect binding on each arm via **Fluorescence Proximity Sensing (FPS)**, or report if the structure is OPEN or CLOSED via sensitive **FRET**.
- Homo-/hetero-proteins can be coupled easily to the arms via exchangeable ligand strands.
- The flexible hinge region confines two proteins to a small volume and defined distance.

Product Description

Order Number: **HK-NYS-2**

Table 1. Contents and Storage Information

Material	Cap	Concentration	Amount	Buffer	Storage
New Y-structure Red Adapter strand 2 with Ra	Orange	400 nM	5 x 50 µL	TE40 ^[1]	-20°C
New Y-structure Green Adapter strand with Ga	Green	400 nM	5 x 50 µL	TE40 ^[1]	-20°C
New Y-structure Ligand Free Strand (lfs) - Red for binary binding in green	Yellow	500 nM	3 x 100 µL	TE40 ^[1]	-20°C
New Y-structure Ligand Free Strand (lfs) - Green for binary binding in red	Yellow	500 nM	3 x 100 µL	TE40 ^[1]	-20°C

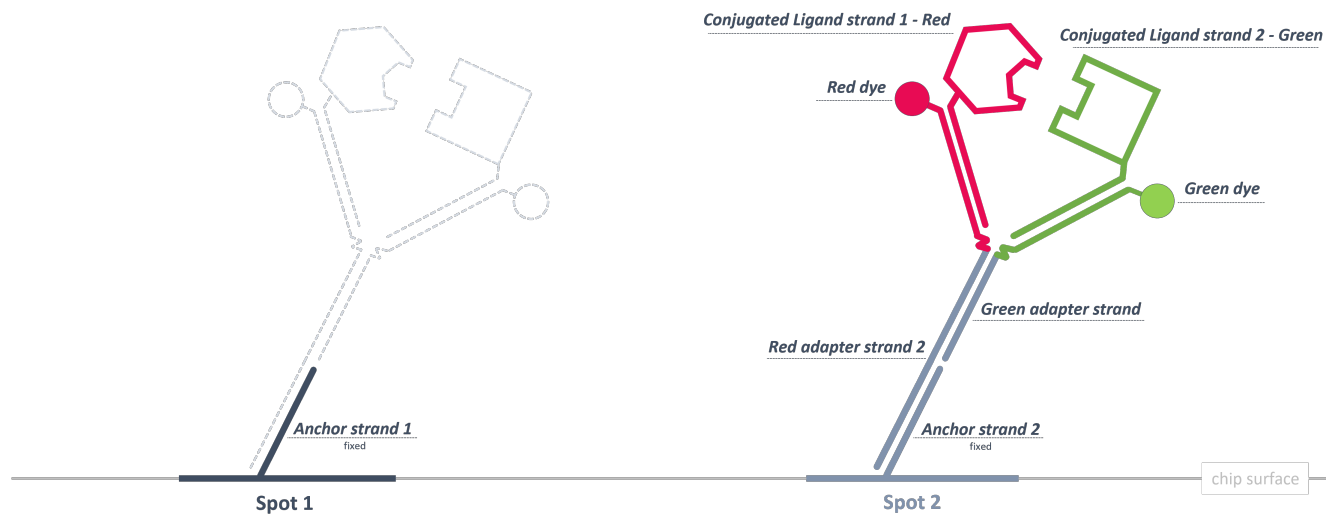
For research use only.

This product has a limited shelf life, please see expiry date on label.

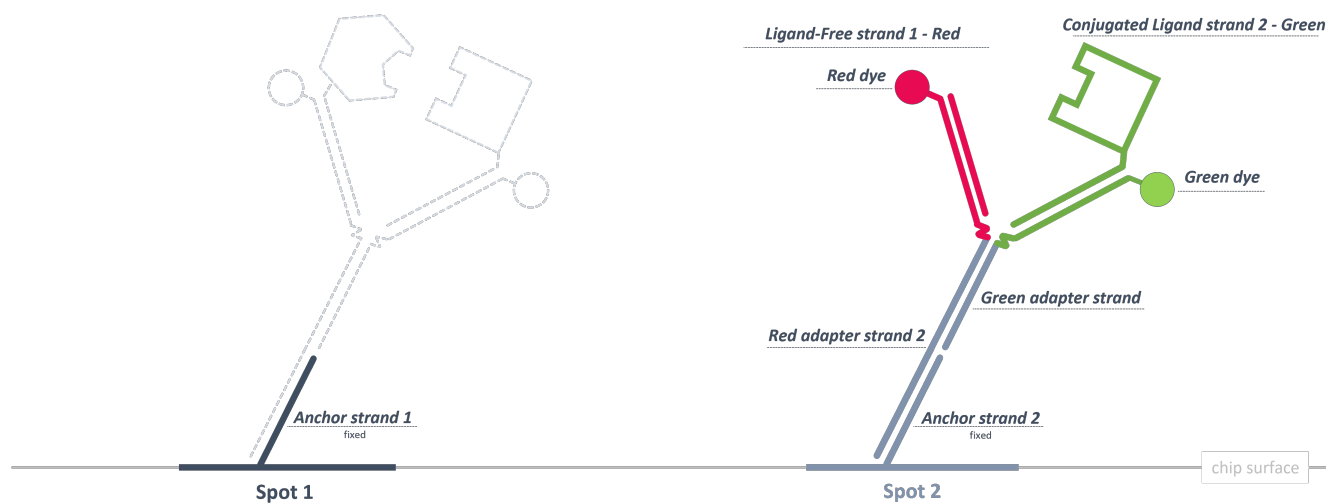
To avoid many freeze thaw cycles please aliquot the nanolever.

heliX[®] Adapter Chip Overview

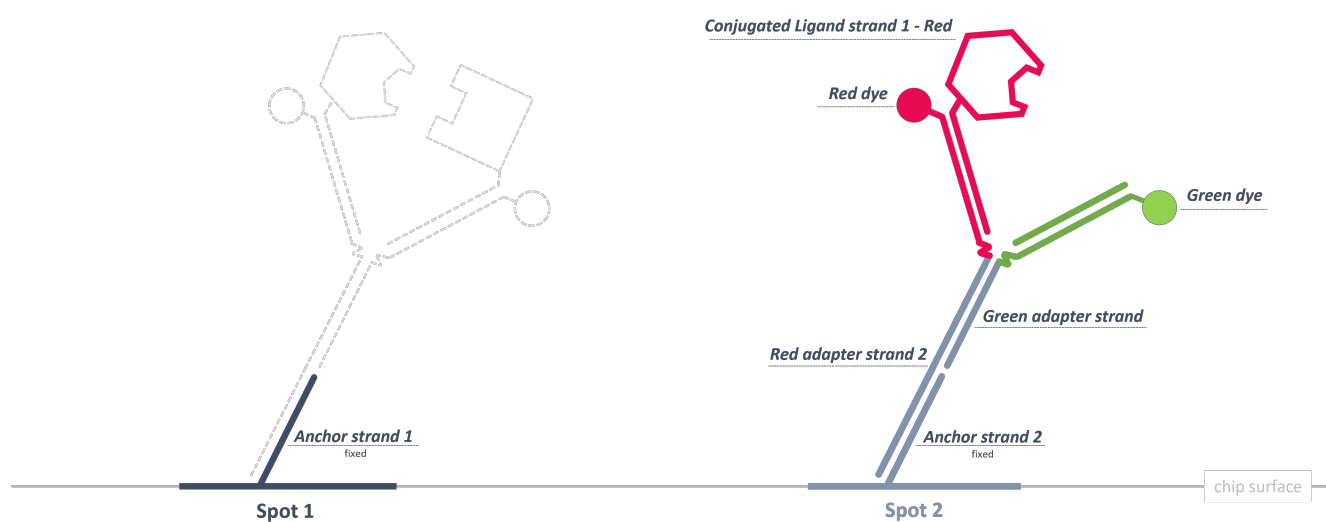
Ternary binding



Binary binding in green



Binary binding in red



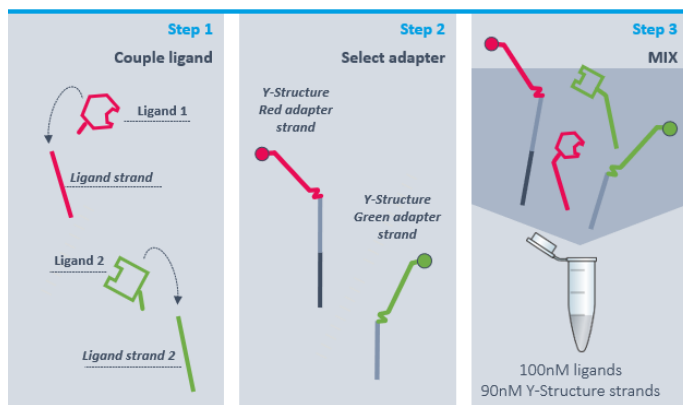
Preparation

Hands-on sample preparation

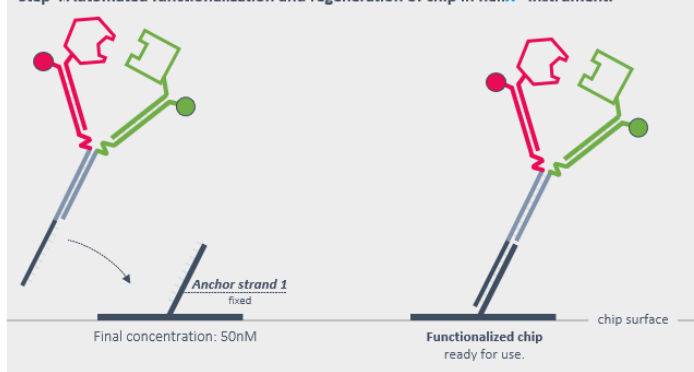
Couple ligands covalently to ligand strands (e.g., via NHS).



Fully automated workflow



Step 4 Automated functionalization and regeneration of chip in helix® instrument.



Choose from ready-to-use coupling kits.



Choose from ready-to-use adapter strands.



Download the helix® MIX&RUN app for use on your mobile device.

*Incubation suggested: overnight

Step 1

The *Ligand strands* can be extended at either the 5' (**Ligand strand 1** - Red arm) or 3' (**Ligand strand 2** - Green arm) end with any DNA/RNA sequence. Additionally, they can be crosslinked to a protein of interest through amine coupling using the specialized **helix®** Amine Coupling Kits (**HK-NYS-NHS-1**, which hybridizes to the red arm, and **HK-NYS-NHS-2**, which hybridizes to the green arm). It is highly recommended to purify the conjugation product with **proFIRE®** before conducting kinetic studies.

TIP

For higher FRET quality, the covalent coupling of the ligands is recommended. + Alternatively, the protein can be captured via His-tag using the HK-NYS-NTA kit.

Step 2

For surface functionalization, the **Y-structure Red Adapter strand 2** harboring the red dye **Ra** and the **Y-structure Green Adapter strand** harboring the green dye **Ga** need to be pre-hybridized with the **Ligand-strands** in order to build the **Y-structure**.

Example of sample preparation for measuring **ternary binding on Spot 1** and **binary binding in green on Spot 2**:

Vial 1 | In solution hybridization of **Y-structure** strands and ligand strands for **Spot 1** (not included in this kit, refer to HK-NYS-1, HK-NYS-NHS-1 and HK-NYS-NHS-2).

- Mix **Y-structure Red Adapter strand** (400 nM), **Green Adapter strand** (400 nM) conjugated **Ligand strand 1** (500 nM) and conjugated **Ligand strand 2** (500 nM) at a 1:1 ratio (v/v).
- Incubate the solution of step i) at **RT for 2 hours** to ensure complete hybridization. Overnight incubation at 4°C is also possible, but it depends on the stability of the conjugated protein.

Vial 2 | In solution hybridization of **Y-structure** strands and ligand strands for **Spot 2**.

- Mix **Y-structure Red Adapter strand 2** (400 nM), **Green Adapter strand** (400 nM), **Ligand-free strand 1 - Red** (500 nM) and conjugated **Ligand strand 2 - Green** (500 nM) at a 1:1 ratio (v/v).
- Incubate the solution of step i) at **RT for 2 hours** to ensure complete hybridization. Overnight incubation at 4°C is also possible, but it depends on the stability of the conjugated protein.

Step 3

Mix solution of step 2 at 1:1 ratio (v/v).

Step 4

Solution is ready to use for **heliX® Adapter Chip** functionalization.

Example

Required volume for 1 functionalization: **35 µL** with a final concentration of **50 nM**.

Vial 1		Vial 2	
New Y-structure Red Adapter strand & Green Adapter strand (400 nM)	Conjugated Ligand strand 1 & Conjugated Ligand strand 2 (500 nM)	New Y-structure Red Adapter strand 2 & Green Adapter strand (400 nM)	Ligand-free strand 1 - Red & Conjugated Ligand strand 2 (500 nM)
4.5 µL each > 9 µL tot	4.5 µL each > 9 µL tot	4.5 µL each > 9 µL tot	4.5 µL each > 9 µL tot

After incubation time, mix vial 1 and vial 2 to obtain a ready-to-use DNA solution.

Assay Setup in heliOS

For studying **ternary complex formation** upon binding of bispecific small molecules (e.g., **PROTACs**, **molecular glues**).

Go to **heliOS** > create a **New Assay Workflow** > add **Custom Assay** > load **Y-Structure FRET Kinetics** > modify the parameters based on your needs and run the assay.

Suggested assay parameters (e.g., flow rates, functionalization time, LED power, etc.) are within the **heliOS** assay.

IMPORTANT

For binary interaction in red, please set LED red ≥ 1 . However, do not forget to set it back to 0 when **FRET** interaction are under investigation.
For more details, please refer to the **heliX®** guide available [this link](#).

Alternatively, **Y-Structure FRET Kinetics - auto LED** assay can be utilized, where the **heliX®** system automatically adjusts the LED power to optimize the fluorescence signal for a better signal-to-noise ratio. This approach is highly recommended for weak binders or screening applications.

For studying bispecific antibodies with weak affinities (e.g., **Hemlibra** binding to Factor X and IX)

Go to **heliOS** > create a **New Assay Workflow** > add simply **Kinetics with Functionalization** from the **Custom Assay** list > modify the parameters based on your needs and run the assay.

TIP

Antibodies are big proteins which do not allow to bring the two dyes in close proximity, therefore **FRET** cannot be recorded. This is the reason why classic kinetics workflow and **Fluorescence Proximity Sensing (FPS)** is used for detecting binding.

For further questions, please contact the support team at support.dbs@bruker.com.

Useful Order Numbers

Table 2. Order Numbers

Product Name	Amount	Order No
Y-structure Amine coupling kit 1 - Red	3 conjugations	HK-NYS-NHS-1
Y-structure Amine coupling kit 2 - Green	3 conjugations	HK-NYS-NHS-2
New Y-structure Kit 1: for proximity binding assay Spot 1	400 nM x 250 µL	HK-NYS-1
New Y-structure His Capture Kit	500 nM x 200 µL	HK-NYS-NTA

Contact

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www.dynamic-biosensors.com

Instruments and chips are engineered and manufactured in Germany.

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[1] TE40: 10 mM Tris, 40 mM NaCl, 0.05 % Tween20, 50 µM EDTA, 50 µM EGTA