



NEW Y-STRUCTURE KIT 1

for proximity binding assay - Spot 1

Dynamic Biosensors GmbH HK-NYS-1 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 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-1

Table 1. Contents and Storage Information

Material	Сар	Concentration	Amount	Buffer	Storage
New Y-structure Red Adapter strand with Ra	Red	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 (Ifs) - Red for binary binding in green	Yellow	500 nM	3 x 100 µL	TE40 ^[1]	-20°C
New Y-structure Ligand Free Strand (Ifs) - Green for binary binding in red	Yellow	500 nM	3 x 100 µL	TE40 ^[1]	-20°C
cAnchor strand 2	Black	100 nM	5 x 200 μL	TE40 [1]	-20°C

For research use only.

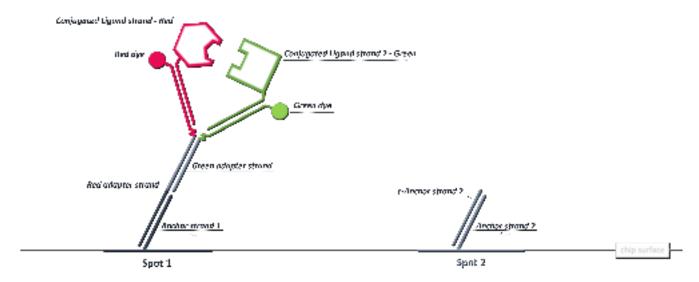
This product has a limited shelf life, please see expiry date on label.

To avoid many freeze thaw cycles please aliquot the nanolever.

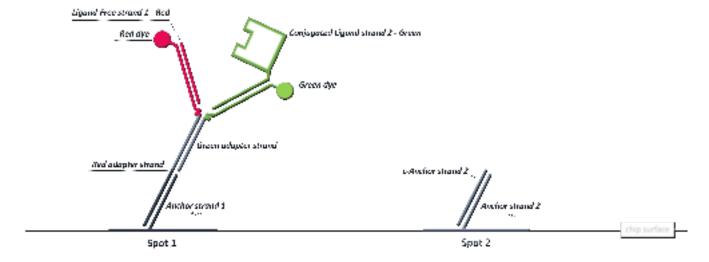


heliX® Adapter Chip Overview

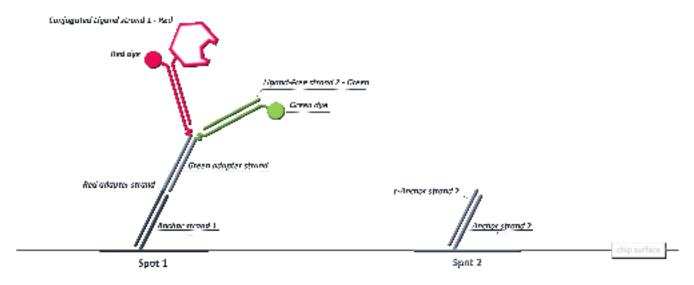
Ternary binding



Binary binding in green

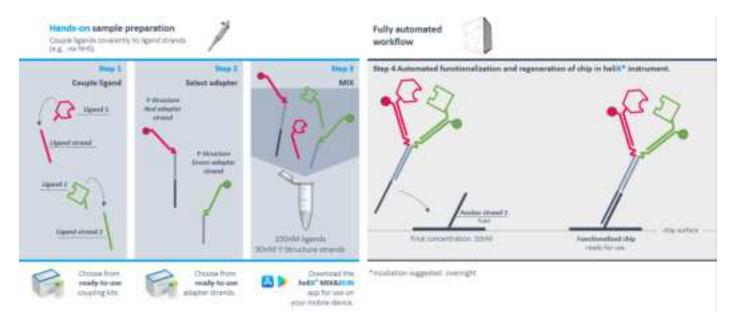


Binary binding in red





Preparation



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 **pro**FIRE® 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. For any questions, please contact the support team at support.dbs@bruker.com.

Step 2

For surface functionalization, the new **Y-structure** *Red Adapter strand* harboring the red dye **Ra** and the new **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**. In solution hybridization of **Y-structure** strands:

- i. Mix the new Y-structure Red Adapter strand with Ra (400 nM), the new Y-structure Green Adapter strand with Ga (400 nM), conjugated Y-structure Ligand strand 1 (500 nM) and conjugated Y-structure Ligand strand 2 (500 nM) at a 1:1 ratio (v/v).
- ii. 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 ii) and cAnchor strand 2 (100 nM) 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			
Y-structure Red Adapter strand (400 nM)	Y-structure Green Adapter strand (400 nM)	Conjugated <i>Ligand</i> strand 1 (500 nM)	Conjugated <i>Ligand</i> strand 2 (500 nM)	cAnchor strand 2 (100 nM)
4.5 µL	4.5 µL	4.5 μL	4.5 μL	18 μL

Assay Setup in heliOS

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

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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 \geq 1. However, do not forget to set it back to 0 when **FRET** interactions are under investigation.

For more details, please refer to the **heliX**⁺ guide available at this **link**.

Alternatevely, **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 2: for proximity binding assay Spot 2	400 nM x 250 μL	HK-NYS-2
New Y-structure His Capture Kit	500 nM x 200 μL	HK-NYS-NTA



Contact

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