

## Quick-Start Protocol

# PICO Protein Detection Trial Kit

The **PICO Protein Detection Trial Kit** (#PICO-000090) must be stored at 4°C and -20°C (see product labels for exact storage conditions). Find the date of expiry on the back side of the pouches.

### Further information:

- The PICO Protein Detection Trial Kit User Manual, the PICO Calculator, and the Safety Data Sheets are available at: [www.actome.de/resources/downloads](http://www.actome.de/resources/downloads)

### Notes before starting:

- Read the **PICO Protein Detection Trial Kit User Manual** for detailed description of the steps and to learn more about the scientific background of a **calibration curve (CLC) experiment** and **relative quantification (RQ)**.
- The cOplete Protease™ Inhibitor Cocktail tablet and the QIAGEN consumables (QIAcuity Probe PCR Mix, QIAcuity Nanoplate 26k 24-Well) are **not included** in the kit.
- Mix the dilutions in the 96-well plates by **gently pipetting up and down 30 times** while avoiding air bubbles.
- Perform **all centrifugation steps** with 1,000 rcf for 5 s at RT unless stated otherwise.
- The concentration of the pre-labeled antibodies is: **4.82 × 10<sup>9</sup> cp/μl**.

### Preparation of the dilution series and the binding reaction:

1. Prepare the following chemicals and buffers:

<b>Additive C (5x stock)</b> add 500 μl PBS	<b>BSA (5x stock)</b> add 400 μl PBS	<b>EDTA-free Protease Inhibitor Cocktail (PIC), (25x stock)</b> dissolve 1 tablet of cOplete Protease™ Inhibitor Cocktail in 2 ml PBS
<b>Lysis Buffer Stock (LB-Stock), (2x stock)</b> 200 μl Additive T 400 μl Additive C 80 μl PIC 200 μl Additive L 120 μl PBS	<b>Lysis Buffer (LB)</b> 300 μl LB-Stock 300 μl PBS	<b>Control Buffer (CB)</b> 250 μl LB-Stock 100 μl BSA 150 μl PBS

2. Thaw the vial of the supplied **recombinant human ErbB2/HER2 protein** and spin it down.
3. Prepare a dilution series of the protein in **LB** in a volume of at least **30 μl** each. Use the **PICO Calculator** (step 1 in the 'PICO Calculator tab') to plan the dilution series:  
**Stock**  $\xrightarrow{1 \text{ in } 100}$  **S#1** (100x)  $\xrightarrow{1 \text{ in } 60}$  **S#2** (6,000x)  $\xrightarrow{1 \text{ in } 2}$  **S#3** (12,000x)  $\xrightarrow{1 \text{ in } 2}$  **S#4** (24,000x)  $\xrightarrow{1 \text{ in } 10}$  **S#5** (240,000x)
4. Prepare the **antibody mix (ABX)** using the **PICO Calculator** (step 2 in the 'PICO Calculator tab'). We recommend using **100 μl** for the total volume of ABX.
5. Mix 2 μl sample with 2 μl ABX (**binding reaction**) in a 96-well PCR microplate following the pipetting scheme (**Figure 1**). For the **NTC** sample use **4 μl LB**. For the **ABC** sample mix **2 μl ABX** with **2 μl CB**.
6. Seal the plate, sonicate at full power for **1 min** and centrifuge the plate (~1,000 rcf, 30 s).
7. Incubate the plate at **4°C** for **12 - 24 h**.

**Pre-dilution and Digital PCR:**

8. Prepare the Master Mix, vortex for **10 s** and spin down.

697  $\mu$ l Ultrapure water  
284  $\mu$ l QIAcuity Probe Mix  
45  $\mu$ l PICO BL Probe  
45  $\mu$ l PICO P8 Probe  
36  $\mu$ l Coupling dPCR Mix

9. Calculate the required dilutions to target a **lambda of 0.15** using the **PICO Calculator** (step 3 in the 'PICO Calculator tab').
10. Prepare a new 96-well plate for the dilution steps. Add the calculated amount of **PBS** to columns 1 - 3 and 5 - 7 and **41  $\mu$ l Master Mix** to columns 10 - 12 (**Figure 2**).
11. Remove the adhesive foil from the plate containing the binding reaction and add **36  $\mu$ l PBS** to all wells (reflects the '10x pre-dilution of binding reaction' in **Figure 2**). Mix by pipetting.
12. Transfer the predetermined volume (usually 1  $\mu$ l) from each sample to the corresponding wells of the dilution plate and mix by pipetting (**DS 1**). Repeat the dilution once more (**DS 2**) and finally transfer **1  $\mu$ l** to the wells containing the **Master Mix** (**Figure 2**). Mix by pipetting.
13. Transfer **40  $\mu$ l Master Mix**, containing the diluted sample, into a **QIAcuity Nanoplate 26k 24-well**. Seal the plate according to the QIAcuity user manual and run a dPCR using the following parameters:

**Priming** - QIAGEN Standard Priming Profile

**PCR conditions**

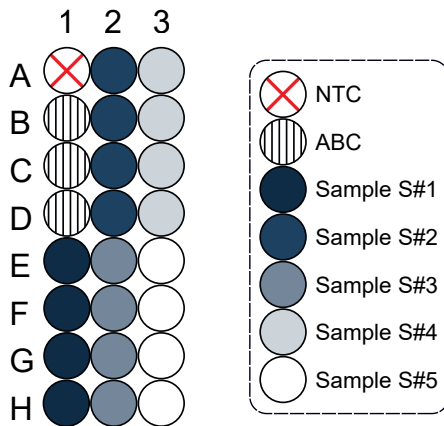
Hot-start	95°C for 2 min
Cycling	40 times
Denaturing	95°C for 15 sec
Annealing	58°C for 30 sec

**Imaging conditions**

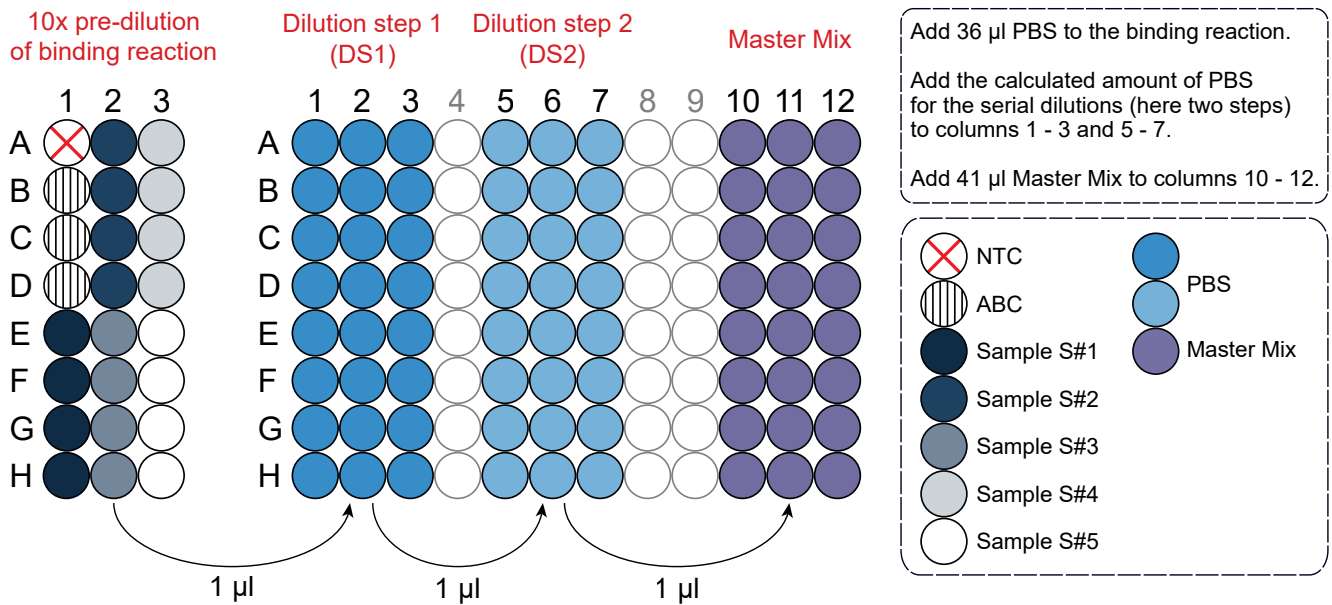
*PICO BL Probe* - HEX yellow channel, 400 ms integration time, gain 6  
*PICO P8 Probe* - FAM green channel, 500 ms integration time, gain 6

14. Instruction for evaluation of raw complex calculation with **AMULATOR** and **Relative Quantification** can be found in the user manual.

### Binding reaction



**Figure 1:** Plate layout of the binding reaction. Five dilution steps, with four technical replicates each, were made. For the ABC three technical replicates are sufficient. ABC - antibody control; NTC - non template control.



**Figure 2:** Plate setup for dPCR pre-dilution. The '10x pre-dilution of binding reaction' is prepared in the plate containing the binding reaction (see **Figure 1**). The two step dilution is prepared in an additional 96-well plate. ABC - antibody control; NTC - non template control.

Scan the QR code for the user manual:

