

Ub-Ub(22-35)-FP K29 linked (human sequence, synthetic)

: UbiQ-046 UbiQ code Batch # : B01012013-001

: 25 ug, 0.84 uL of 2.89 mM stock in DMSO Amount

Purity : ≥95% by RP-HPLC

Mol. Weight: 10.55 kDa

: upon arrival, store solution at -80°C. Store dark and avoid multiple freeze/thaw cycles. Storage

Productsheet

Background. UbiQ-046 is a fluorescence polarization reagent based on the peptide sequence 22–35 of ubiquitin. The peptide is modified on the N-terminus with a 5-carboxytetramethylrhodamine (TAMRA) and conjugated at Lys29 to Ub via a native isopeptide bond.

sequence

TAMRA-TIENVKAK(Ub)IQDKEG

important: sample preparation

- dilute DMSO stock with milliQ and mix well (e.g., 29x to a 100 µM stock; this can be aliquoted and stored)
- For assays this 100 μM stock can be diluted for example 1000× in buffer affording a final assay solution of 100 nM. The DMSO concentration during the assay is now 0.01 vol%.
- the concentration can be verified by comparing the fluorescence intensity with that of a known concentration of TAMRA.
- for full experimental details please see reference 5.

Literature. (1) Tirat et al. Analytical Biochem. 2005, 343, 244. (2) Huang et al. Methods in Mol Biol 2009, 565, 127. (3) Levine et al. Analytical Biochem. 1997, 247, 83. (4) Faesen et al. Chem. Biol. 2011, 18, 1550. (5) Geurink et al. ChemBiochem 2012, 13, 293.



General experimental procedures for fluorescence polarization assays

FP assays were performed on a PerkinElmer Wallac EnVision 2010 Multilabel Reader with a 531 nm excitation filter and two 579 nm emission filters. Fluorescence intensities were measured in the S (parallel) and P (perpendicular) direction. FP values are given in mP (millipolarization) and calculated using the following formula (1):

Polarization (mP) =
$$\frac{S-(G \cdot P)}{S+(G \cdot P)} \cdot 1000$$
 (1)

The confocal optics were adjusted with the average P and S values for TAMRA-Lys-Gly and the grating factor (G) was determined using a polarization value (L) for TAMRA of 50 mP using the following formula (2):

$$G = \frac{\text{average S}}{\text{average P}} \cdot \frac{1 - (\frac{L}{1000})}{1 + (\frac{L}{1000})}$$
(2)

The assays were performed in "non-binding surface flat bottom low flange" black 384-well plates (Corning) at room temperature in a buffer containing 20 mM Tris·HCl, pH 7.5, 5 mM DTT, 100 mM NaCl, 1 mg/mL 3-[(3-cholamidopropyl) dimethylammonio] propanesulfonic acid (CHAPS) and 0.5 mg/mL bovine gamma globulin (BGG). Each well had a volume of 20 μ L. Buffer and enzyme were predispensed and the reaction was started by the addition of substrate. Kinetic data was collected in intervals of 2.5 or 3 min. From the obtained polarization values (Pt) the amount of processed substrate (St) was calculated with to the following formula (3):

$$S_{t} = S_{0} - S_{0} \times \left(\frac{P_{t} - P_{\min}}{P_{\max} - P_{\min}}\right)$$
(3)

Where P_t is the polarization measured (in mP); P_{max} is the polarization of 100% unprocessed substrate (determined for every reagent at all used substrate concentrations); P_{min} is the polarization of 100% processed substrate; S_0 is the amount of substrate added to the reaction.

From the obtained P_t values the values for initial velocities (v_i) were calculated. The v_i values are used to determine the Michaelis-Menten constants (K_m , V_{max} and k_{cat}) by fitting the data according to formula (4) (where $k_{cat} = V_{max}/[E]$). All experimental data was processed using Ms Excel and Prism 4.03 (GraphPad Software, Inc.)

$$v_{i} = \frac{V_{\text{max}} \times S_{0}}{K_{\text{m}} + S_{0}} \tag{4}$$