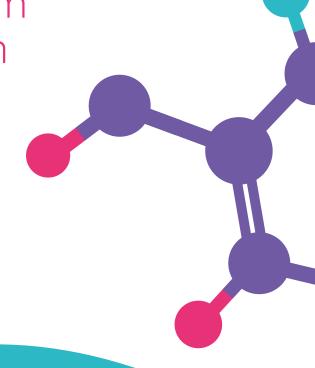


# Ubiquitin Peptides from Cambridge Research Biochemicals

Ubiquitinylation is a post-translational modification involving the covalent binding of the small, highly conserved protein ubiquitin to a target substrate to regulate protein functions.





This set of high quality AQUA peptides will allow you to absolutely quantify all ubiquitin chain linkages in a sample using targeted SRM/PRM mass spectrometry approaches. As there are no good antibodies for many chain linkages, it is indeed the only way to quantify all ubiquitin chain linkages.

# **Professor Matthias Trost**

Professor of Proteomics Institute for Cell and Molecular Biosciences Newcastle University



The ligation of the ubiquitin C-terminus to a target protein is facilitated by a cascade of ubiquitin activating and ligating enzymes and reversed by deubiquitylating enzymes (DUBs).

Ubiquitin possesses seven lysine residues and an N-terminus methionine (M1, K6, K11, K27, K29, K33, K48 and K63), any one of which can be ubiquitinylated itself leading to the assembly of polyubiquitin chains.

Any of these chains lead to different biological outcomes. For example, while K48 chains lead to proteasomal degradation, K63 chains are linked to cell signalling, trafficking and lysosomal degradation.

Ubiquitin signalling is integral to almost all cellular processes in eukaryotes and thus, disorders or mutations in ubiquitin pathways often result in cancer, autoimmune diseases and neurodegenerative diseases such as Parkinson's, Alzheimer's and Huntington's.

Mass spectrometry (MS) based proteomics has become a powerful method for the analysis of ubiquitinylation in the last 15 years. Particularly, targeted MS approaches using heavy isotope labelled K-GG peptides help to identify and quantify ubiquitin chain topologies.

Heap, R.E., Gant, M.S., Lamoliatte, F., Peltier, J. and Trost, M., Mass spectrometry techniques for studying the ubiquitin system (2017), Biochem Soc Transactions, Oct 15;45(5):1137-1148.

### Unlabelled Peptides

Ubiquitin M1 Light crb1000835 GGMQIFVK-acid 25nmol £85

Ubiquitin K6 Light crb1000805 MQIFV-[K(GG)]-TLTGK-acid 25nmol £150

Ubiquitin K11 Light crb1000807 TLTG-[K(GG)]-TITLEVEPSDTIENVK-acid 25nmol £150

Ubiquitin K27 Light crb1000809
TITLEVEPSDTIENV-[K(GG)]-AK-acid 25nmol £150

Ubiquitin K29 Light crb1000811 A-[K(GG)]-IQDK-acid 25nmol £150

Ubiquitin K33 Light crb1000813 IQD-[K(GG)]-EGIPPDQQR-acid 25nmol £150

Ubiquitin K48 Light crb1000815 LIFAG-[K(GG)]-QLEDGR-acid 25nmol £150

Ubiquitin K63 Light crb1000817 TLSDYNIQ-[K(GG)]-ESTLHLVLR-acid 25nmol £150

## Controls

Ubiquitin (12-27) Light crb1001282
TITLEVEPSDTIENVK-acid 25nmol £85

Ubiquitin (12-27) Heavy crb1301283
TITLEVEPSDTIENV[U-13C<sub>6</sub>, <sup>15</sup>N<sub>2</sub>-Lys]-acid
25nmol £220

Ubiquitin (64-72) Light crb1001284 ESTLHLVLR-acid 25nmol £85

**Ubiquitin (64-72)** Heavy crb1301285 ESTLHLVL-[U-<sup>13</sup>C<sub>6</sub>, <sup>15</sup>N<sub>4</sub>-Arg]-acid **25nmol £220** 

# Stable-labelled Peptides

**Ubiquitin M1** Heavy crb1300804 GGMQIFV-[U-<sup>13</sup>C<sub>6</sub>, <sup>15</sup>N<sub>2</sub>-Lys]-acid **25nmol £220** 

Ubiquitin K6 Heavy crb1300806 MQIFV-[K(GG)]-TLTG-[U-13C<sub>6</sub>,15N<sub>2</sub>-Lys]-acid 25nmol £275

Ubiquitin K11 Heavy crb1300808 TLTG-[K(GG)]-TITLEVEPSDTIENV-[U-13C<sub>6</sub>, 15N<sub>2</sub>-Lys]-acid 25nmol £275

Ubiquitin K27 Heavy crb1300810 TITLEVEPSDTIENV-[K(GG)]-A-[U-13C<sub>6</sub>, 15N<sub>2</sub>-Lys]-acid 25nmol £275

**Ubiquitin K29** Heavy crb1300812 A-[K(GG)]-IQD-[U-<sup>13</sup>C<sub>6</sub>, <sup>15</sup>N<sub>2</sub>-Lys]-acid **25nmol £275** 

Ubiquitin K33 Heavy crb1300814 IQD-[K(GG)]-EGIP-[U-<sup>13</sup>C<sub>5</sub>, <sup>15</sup>N-Pro]-DQQ-[U-<sup>13</sup>C<sub>6</sub>, <sup>15</sup>N<sub>4</sub>-Arg]-acid 25nmol £320

Ubiquitin K48 Heavy crb1300816 LIFAG-[K(GG)]-QLEDG-[U-13C<sub>6</sub>, 15N<sub>4</sub>-Arg]-acid 25nmol £350

Ubiquitin K63 Heavy crb1300818 TLSDYNIQ-[K(GG)]-ESTLHLVL-[U-13C<sub>6</sub>, 15N<sub>4</sub>-Arg]-acid 25nmol £275

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