

http://www.clickchemistrytools.com E: inquiries@clickchemistrytools.com

T: 480 584 3340 F: 866 717 2037

DBCO-Maleimide

Important Product Information

- Molecules to be reacted with maleimide compounds must have free (reduced) sulfhydryls. Reduce peptide disulfide bonds with disulfide reducing reagents such as Immobilized TCEP Disulfide Reducing Gel (Pierce Biotechnology). Reduce disulfide bonds in high molecular weight proteins using 5 mM TCEP (1:100 dilution) for 30 minutes at room temperature, followed by TCEP removal using a desalting column. Proteins (e.g., antibodies) can be inactivated by complete reduction of their disulfide bonds. Selective reduction of hingeregion disulfide bonds in IgG can be accomplished with 2-Mercaptoethylamine•HCl (2-MEA). Sulfhydryls can be added to molecules using N-succinimidyl S-acetylthioacetate (SATA) or 2-iminothiolane•HCl (Traut's Reagent), which modify primary amines.
- Do not use buffers that contain sulfhydryl-containing components (e.g., DTT).
- Avoid buffers that contain azides, which can react with DBCO.
- The maleimide group reacts predominantly with free sulfhydryls at pH 6.5-7.5, forming stable thioether bonds. At pH values > 7.5, reactivity toward primary amines and hydrolysis of the maleimide groups can occur. At pH 7, the maleimide group is ~1,000 times more reactive toward a free sulfhydryl than to an amine.

Procedure for Sample Labeling

Additional Materials Required

- Water-miscible organic solvent such as dimethyl sulfoxide (DMSO) or dimethyl formamide (DMF)
- Reducing reagents such as Immobilized TCEP Disulfide Reducing Gel (Pierce Biotechnology)
- Reaction buffer: Phosphate-buffered saline (PBS) or other sulfhydryl-free buffer at pH 6.5-7.5. Include 5-10 mM EDTA to help prevent the reoxidation of disulfides by trace divalent metals.
- (Optional): Quenching buffer: concentrated (0.5-1 M) cycteine, DDT or other thiol containing reducing
 agents
- (Optional): Spin Desalting Columns

Protein Labeling

- Prepare reduced sulfhydryl-containing protein in reaction buffer containing 5-10 mM EDTA as described the Important Product Information section.
- Immediately before use, weigh a small quantity of DBCO-Maleimide and dissolve it in dimethylformamide (DMF) or dimethylsulfoxide (DMSO) at a 5-20 mM concentration.

- Add DBCO-Maleimide solution to the reduced protein(s) at ca. four-fold molar excess over available sulfhydryl groups.
- Note: The reaction solution may appear cloudy as a result of the low aqueous solubility of DBCO-Maleimide; usually, such solutions become clearer as the reaction proceeds. Many proteins will precipitate when the DMF or DMSO concentration exceeds 10-15% of the final reaction volume; if protein solubility is not an issue, there is no limit to the DMF or DMSO concentration that may be used.
- Incubate reaction mixture for 1 hour at room temperature or for 2 hours at 4°C.
- Quench reaction by adding Quenching Solution at 10-50 mM final and incubating for 15 minutes at room temperature. Alternatively (or in addition) remove the excess non-reacted reagent by desalting or dialysis.

Copper-free Click Reaction

- 1. Prepare the azide-containing sample in reaction buffer.
- 2. Add DBCO-protein conjugate to azide-containing sample.

Recommendation: Add 1 mole equivalent of limiting reagent to 2-4 molar equivalents of highest abundance reagent.

- 3. Incubate the reaction at room temperature for 2-12 hour.
- 4. The reaction is now ready for purification by size exclusion chromatography if required.

Troubleshooting

Problem	Possible Cause	Solution
	One or more sample is not	Confirm molecules were labeled or repeat
	labeled	activation process
No conjugation of		Allow product to equilibrate to room temperature before
DBCO with azide		opening
	DBCO-Maleimide decomposed	Prepare new solutions in the indicated dry solvents
		Avoid buffers that contain sulfhydryl
	Excess reagent not	Remove non-reacted reagent by dialysis or desalting
	quenched or removed	
Low conjugation of DBCO and azide	Suboptimal reaction conditions	Increase incubation time
		Optimize conjugation conditions by altering molar excess
		Perform conjugation reactions at 37°C