



Correction

Correction: Braun, T., et al. Expanding the Genetic Code for Site-Directed Spin-Labeling. *Int. J. Mol. Sci.* 2019, 20, 373

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The authors wish to make the following two corrections to this paper [1]:

1. In Figures 2 and 5, a false structure of the noncanonical amino acid N^ϵ -Bicyclo[6.1.0]non-2-yn-9-ylmethoxycarbonyl-L-lysine 5 has been shown. The corrected Figure 2 should be replaced with the following figure (Figure 1), the corrected Figure 5 should be replaced with the following figure (Figure 2).

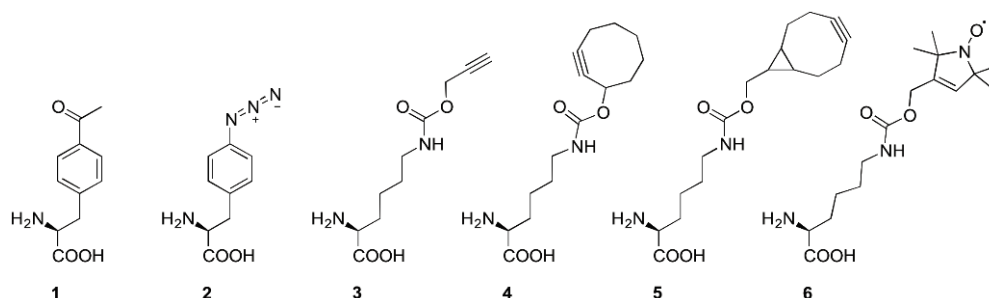


Figure 1. Overview of ncAA with reported use for SDSL. *p*-Acetyl-L-phenylalanine 1, *p*-azido-L-phenylalanine 2, N^ϵ -propargyloxycarbonyl-L-lysine 3, N^ϵ -Cyclooct-2-ynyloxycarbonyl-L-lysine 4, N^ϵ -Bicyclo[6.1.0]non-2-yn-9-ylmethoxycarbonyl-L-lysine 5, N^ϵ -1-oxy-2,2,5,5-tetramethylpyrroline-3-ylmethoxycarbonyl-L-lysine 6.

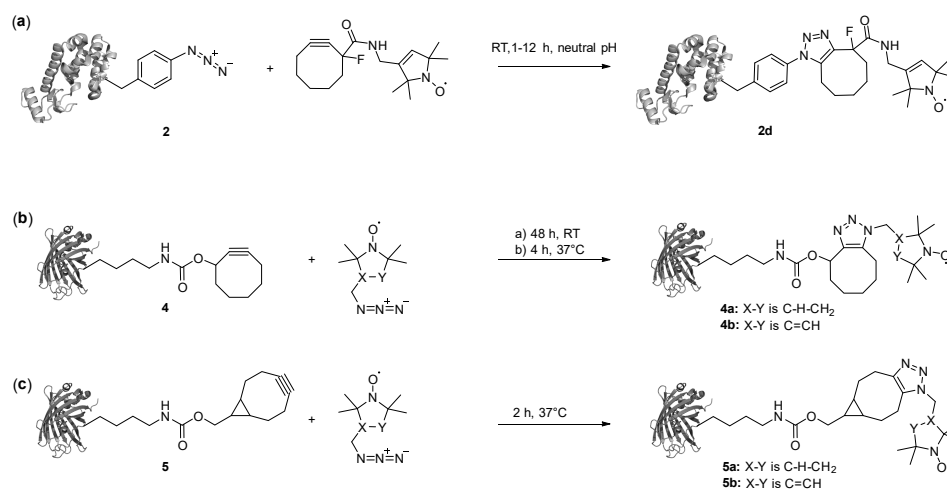


Figure 2. Conjugation of ncAA with nitroxide labels via strain-promoted azide–alkyne cycloadditions (SPAAC). The strained alkyne can either be part of the label reagent (a) or the ncAA (b) and (c).



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2. The authors wish to add the grant number “ERC-CoG EPICODE to D.S., Grant Nr. 723863” to the Acknowledgments section.

The authors would like to apologize for any inconvenience caused to the readers by these changes.

Reference

1. Braun, T.; Drescher, M.; Summerer, D. Expanding the Genetic Code for Site-Directed Spin-Labeling. *Int. J. Mol. Sci.* **2019**, *20*, 373. [\[CrossRef\]](#) [\[PubMed\]](#)