

Book of Abstracts



February 19-21, 2020 V GEQB ChemBio Group Meeting

Organized by:







ACETALS AS ACID-SENSITIVE CLEAVABLE LINKERS FOR THE DESIGN OF ANTIBODY-DRUG CONJUGATES

<u>E. Jiménez-Moreno,</u>¹ X. Ferhati,¹ P. Akkapeddi,² M. J. Matos,³ N. Salaverri,¹ G. Jiménez-Osés,^{1,4} G. J. L. Bernardes,^{2,3} and F. Corzana¹

¹ Departamento de Química, Universidad de La Rioja, Centro de Investigación en Síntesis Química, 26006 Logroño, Spain
² Instituto de Medicina Molecular, Faculdade de Medicina, Universidade de Lisboa, 1649-028 Lisboa, Portugal
³ Department of Chemistry, University of Cambridge, Lensfield Road, CB2 1EW, Cambridge, UK.

⁴ CIC bioGUNE, Bizkaia Technology Park, Building 800, 48170 Derio, Spain.

Email: ester.jimenez@unirioja.es

Several acid-cleavable linkers, based on an acetal group, that feature either a deactivated coumarin or a prodrug derived from the potent anti-cancer agent duocarmycin, have been designed. The linkers are stable in plasma and can rapidly break down in acidic pH to generate the free fluorophore or the toxic drug in situ. These scaffolds were conjugated to a Trastuzumab antibody that is specific for Her2, a receptor that is overexpressed in breast cancer cells. Interestingly, although the Trastuzumab-coumarin



conjugate is stable in plasma, the analogue that carries the duocarmycin derivative slowly decomposes under similar conditions. Molecular dynamics (MD) simulations performed on these antibody-drug conjugates (ADCs) suggest that a lysine residue of the antibody nearby the conjugation site can act as an acid, which promotes hydrolysis of the acetal bearing the duocarmycin derivative. These data demonstrate that both the conjugation site of the antibody and the combination linker-payload can modulate the stability of the conjugate. Finally, we show that both ADCs retain their specificity to cancer cells. Therefore, the straightforward synthesis combined with the possibility of incorporating different payloads, makes the use of acetals an innovative and competitive strategy for the design of targeted drug-delivery systems.

Acknowledgements

We thank MINECO (projects RTI2018-099592-B-C21 to F.C. and RTI2018-099592-B-C22 and RYC-2013-14706 to G.J.O)., the EU (Marie-Sklodowska Curie ITN, ProteinConjugates, grant agreement No. 675007 to G.J.L.B., F.C., and X.F.), Universidad de La Rioja (postdoctoral contract to E. J. -M.), Xunta da Galicia (Plan I2C–ED481B 2014/086-0 and ED481B 2018/007 to M.J.M.), Royal Society (URF/R/180019 to G.J.L.B.) and FCT Portugal (iFCT IF/00624/2015 to G.J.L.B.). We also thank Genentech Inc. for providing the Thiomab[®] antibody and Dr Vikki Cantrill for her help with the editing of this manuscript.