

Corrigendum

SEVA 3.0: an update of the Standard European Vector Architecture for enabling portability of genetic constructs among diverse bacterial hosts

Esteban Martínez-García¹, Angel Goñi-Moreno², Bryan Bartley³, James McLaughlin², Lucas Sánchez-Sampedro⁴, Héctor Pascual del Pozo⁴, Clara Prieto Hernández⁴, Ada Serena Marletta⁵, Davide De Lucrezia⁵, Guzmán Sánchez-Fernández⁴, Sofía Fraile¹ and Víctor de Lorenzo^{1,*}

¹Systems Biology Program, Centro Nacional de Biotecnología CSIC, Campus de la Universidad Autónoma de Madrid, 28049 Spain, ²School of Computing, Newcastle University NE4 5TG, UK, ³Raytheon BBN Technologies, Arlington, VA 22209, USA, ⁴Scienseed SL, 28020 Madrid, Spain and ⁵Doulix-Explora Srl, Via G. Peroni 386, 00131, Italy

Nucleic Acids Research, 2020, 48(D1): D1164–D1170, <https://doi.org/10.1093/nar/gkz1024>

Following an upgrade to the database, two links provided in the article are no longer working:

The link on page D1167:

https://doulx.com/public-biomodules/?name=&friendly_id=&o=name&biobrick_equivalent=&lab_tested=None&public=None&shared=None&description=&sequence=&amino_sequence=&collection=lrp

has been replaced by:

<https://doulx.com/collections/lrp/biomodules/>

The link on page D1167:

https://doulx.com/public-constructs/?name=&friendly_id=&o=-updated&lab_tested=None&public=None&shared=None&description=&sequence=&amino_sequence=&collection=lrp

has been replaced by

<https://doulx.com/collections/lrp/constructs/>

These corrections do not affect the results and conclusion of the study.

*To whom correspondence should be addressed. Email: vdlorenzo@cnb.csic.es