

Update Notice: Efficient Generation of Genome-wide Libraries for Protein–ligand Screens Using Gibson Assembly

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After official publication in Bio-protocol (<https://bio-protocol.org/e4558>), we would like to add an updated information to the Acknowledgments section: “This protocol is derived from the original research paper by Rhiannon Heslop et al. (2023).”

The updated version of the Acknowledgments section is “This work was funded by the Canadian Institutes of Health Research (CIHR PJT-175222, to IC), the Natural Sciences and Engineering Research Council of Canada (NSERC RGPIN-2019-05271, to IC), the Canada Foundation for Innovation (JELF 258389, to IC), and by McGill University (130251, to IC). We thank Andressa Lira and Sahil Sanghi for their technical support at the early stages of this work. This research was enabled in part by computational resources provided by Calcul Quebec (<https://www.calculquebec.ca/en/>) and Compute Canada (www.computeCanada.ca). **This protocol is derived from the original research paper by Rhiannon Heslop et al. (2023).**

Accordingly, a new reference is added to the References list, “17. Heslop, R., Gao, M., Brito Lira, A., Sternlieb, T., Loock, M., Sanghi, S. R. and Cestari, I. (2023). [Genome-Wide Libraries for Protozoan Pathogen Drug Target Screening Using Yeast Surface Display](#). ACS Infect. Dis: e2c00568.”