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Submission Complete

pride-support@ebi.ac.uk <pride-support@ebi.ac.uk>

Wed, Mar 27, 2019 at 2:54 AM

To: strigg@uw.eduCc: sr320@uw.edu

Dear Shelly Trigg,

We are happy to inform you that your dataset "Oyster seed time x temperature, LC-MSMS" has been successfully submitted to ProteomeXchange via the PRIDE database. The data is currently private, and can only be accessed with your account or with a single reviewer account that has been created.

Please note that it is essential that you notify us of the first (online) publication of the corresponding manuscript. Otherwise the data will remain inaccessible to readers.

Please add to your manuscript the following sentence (typically in the "Methods" section or just before/in the Acknowledgements):

"The mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium via the PRIDE [1] partner repository with the dataset identifier PXD013262".

We would recommend you to also include this information in a much abridged form into the abstract itself, e.g. "Data are available via ProteomeXchange with identifier PXD013262."

Submission details:

Project Name: Oyster seed time x temperature, LC-MSMS

Project accession: PXD013262

Project DOI: Not applicable

Reviewer account details:

Username: reviewer73630@ebi.ac.uk

Password: 0lZ9vZmB

Don't forget to include these reviewer account details in your manuscript, during the peer review process.

Thank you for choosing PRIDE and ProteomeXchange for the dissemination of your data, and look forward to receiving more of your data in the future.

Regards,

The PRIDE Team, EMBL-EBI

Website: <http://www.ebi.ac.uk/pride>

Email: pride-support@ebi.ac.uk

Twitter: [@pride_ebi](https://twitter.com/pride_ebi)

For general PRIDE reference, please use:

[1] Perez-Riverol Y, Csordas A, Bai J, Bernal-Llinares M, Hewapathirana S, Kundu DJ, Inuganti A, Griss J, Mayer G, Eisenacher M, Pérez E, Uszkoreit J, Pfeuffer J, Sachsenberg T, Yilmaz S, Tiwary S, Cox J, Audain E, Walzer M, Jarnuczak AF, Ternent T, Brazma A, Vizcaíno JA (2019). The PRIDE database and related tools and resources in 2019: improving support for quantification data. *Nucleic Acids Res* 47(D1):D442-D450 (PubMed ID: 30395289).

For general ProteomeXchange reference, please use:

[2] Deutsch EW, Csordas A, Sun Z, Jarnuczak A, Perez-Riverol Y, Ternent T, Campbell DS, Bernal-Llinares M, Okuda S, Kawano S, Moritz RL, Carver JJ, Wang M, Ishihama Y, Bandeira N, Hermjakob H, Vizcaíno JA (2017). The ProteomeXchange Consortium in 2017: supporting the cultural change in proteomics public data deposition. *Nucleic Acids Res* 54(D1):D1100-D1106 (PubMed PMID: 27924013).

For PRIDE Inspector reference, please use:

[3] Perez-Riverol Y, Xu QW, Wang R, Uszkoreit J, Griss J, Sanchez A, Reisinger F, Csordas A, Ternent T, del Toro N, Dianes JA, Eisenacher M, Hermjakob H, Vizcaíno JA (2016). PRIDE Inspector Toolsuite: moving towards a universal visualization tool for proteomics data standard formats and quality assessment of ProteomeXchange datasets. *Mol Cell Proteomics* 15(1):305-17 (PubMed ID: 26545397).

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