

Resource Summary Report

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R editing of HMMSEARCH output: heatmap with associated with gene tree and protein illustration

DOI:10.17504/protocols.io.bkqwkvxe

Type: Protocol

Proper Citation

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Protocol Information

URL: <https://dx.doi.org/10.17504/protocols.io.bkqwkvxe>

Authors: Kanae Nishii

Summary: R codes used for phyml tree and profile HMM heatmap visualization, protein domain illustration

Affiliations: Royal Botanic Garden Edinburgh; Kanagawa University

Version: 1

Publication Date: 2021

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