

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

TB Consortium Bias Removal Server

RRID:SCR_008425

Type: Tool

Proper Citation

TB Consortium Bias Removal Server (RRID:SCR_008425)

Resource Information

URL: <http://tuna.tamu.edu>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented August 23, 2016. Map improvement server that returns a bias minimized, 6-fold averaged map generated from a model and diffraction data (with optional preceding Molecular Replacement). It does not build or repair the model for you (yet). For automated model building, you need to install a local copy of CCP4 and ARP/wARP (aka wARP&Trace), RESOLVE, MAID, or TEXTAL.

Synonyms: Bias Removal Server

Resource Type: software resource

Keywords: map, server, diffraction, data, electron, density, model, software

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: TB Consortium Bias Removal Server

Resource ID: SCR_008425

Alternate IDs: nif-0000-30172

Record Creation Time: 20220129T080247+0000

Record Last Update: 20260905T132622+0000

Ratings and Alerts

No rating or validation information has been found for TB Consortium Bias Removal Server.

No alerts have been found for TB Consortium Bias Removal Server.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Miller MA, et al. (2018) Parasite spillover: indirect effects of invasive Burmese pythons. Ecology and evolution, 8(2), 830.

Pornillos O, et al. (2009) X-ray structures of the hexameric building block of the HIV capsid. Cell, 137(7), 1282.