

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 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>

**Proper Citation:** TB Consortium Bias Removal Server (RRID:SCR\_008425)

**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.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [Kravitz](#) .

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.