## **Resource Summary Report**

Generated by NIF on May 25, 2025

# toxprofileR

RRID:SCR\_017027

Type: Tool

### **Proper Citation**

toxprofileR (RRID:SCR\_017027)

#### **Resource Information**

URL: https://git.ufz.de/iTox/toxprofileR

Proper Citation: toxprofileR (RRID:SCR\_017027)

**Description:** Software R package to derive toxicogenomic fingerprints from microarray data.

**Resource Type:** software library, software application, software toolkit, data analysis software, data processing software, software resource

**Keywords:** toxicology, zebrafish, toxicogenomics, transcriptomics, microarray, regression, modelling, fingerprint, data

**Funding:** 

Availability: Free, Available for download, Freely available

Resource Name: toxprofileR

Resource ID: SCR\_017027

License: GNU General Public License

**Record Creation Time: 20220129T080333+0000** 

**Record Last Update:** 20250525T031502+0000

### Ratings and Alerts

No rating or validation information has been found for toxprofileR.

No alerts have been found for toxprofileR.

## Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

Schüttler A, et al. (2019) Map and model-moving from observation to prediction in toxicogenomics. GigaScience, 8(6).