

# Profiles Research Networking Software Read Me First

**Documentation Version:** March 23, 2015

**Software Version:** ProfilesRNS\_2.6.0

## **Table of Contents**

|                        |   |
|------------------------|---|
| Introduction .....     | 2 |
| Where to Go Next ..... | 3 |
| More Information.....  | 4 |
| Acknowledgements.....  | 5 |

## **Introduction**

Profiles Research Networking Software is an open source tool to speed the process of finding researchers with specific areas of expertise for collaboration and professional networking. Profiles RNS imports and analyzes "white pages" information, publications, and other data sources to create and maintain a complete searchable library of web-based electronic CV's. Built-in network analysis and data visualization tools allow administrators to generate research portfolios of their institution, discover connections between parts of their organization, and understand what factors influence collaboration.

Profiles RNS is a Semantic Web application that uses the Resource Description Framework (RDF) data model. In RDF, every entity (e.g., person, publication, concept) is given a unique URI. Entities are linked together using "triples" that contain three URIs--a subject, predicate, and object. For example, the URI of a Person can be connected to the URI of a Concept through a predicate URI of hasResearchArea. An instance of Profiles RNS can have millions of URIs and triples. Semantic Web applications use an ontology, which describes the classes and properties used to define entities and link them together. Profiles RNS uses the VIVO Ontology, which was developed as part of an NIH-funded grant to be a standard for academic and research institutions. A growing number of sites around the world are adopting research networking platforms that use the VIVO Ontology. Because RDF can link different triple-stores that use the same ontology, software developers are able to create tools that span multiple institutions and data sources.

## **Where to Go Next**

This document provides a brief introduction to Profiles RNS and information on the resources that are available to help you install and use the software. The other documentation files are:

- [ProfilesRNS\\_InstallGuide.pdf](#) – Follow the instructions in this document to install a new instance of Profiles RNS or upgrade from an older version.
- [ProfilesRNS\\_APIGuide.pdf](#) – An important feature of Profiles RNS is its ability to share data. The API Guide describes the various types of ways this can be done. It also contains a brief introduction to the ontology used by Profiles RNS. The API\_Examples folder contains related files, such as XSDs and example API request messages.
- [ProfilesRNS\\_ArchitectureGuide.pdf](#) – This document describes how Profiles RNS works and how you can extend the ontology, add custom data feeds, or modify the user interface.
- [ProfilesRNS\\_ReleaseNotes.pdf](#) – This document lists new features, bug fixes, and known issues with each release of Profiles RNS.

## **More Information**

For more information about Profiles RNS, please visit

<http://profiles.catalyst.harvard.edu>

The Harvard development team can be reached at [profiles@hms.harvard.edu](mailto:profiles@hms.harvard.edu). We will try to reply promptly, though we cannot guarantee that we will be able to answer all questions.

## **Acknowledgements**

Profiles Research Networking Software was developed under the supervision of Griffin M Weber, MD, PhD, with support from Grant Number 1 UL1 RR025758-01 to Harvard Catalyst, The Harvard Clinical and Translational Science Center from the National Center for Research Resources and support from Harvard University and its affiliated academic healthcare centers.

### Harvard Development Team

The software implementation is led by the Harvard Medical School Information Technology Department. The current and past members of the development team include Nick Benik, Nick Brown, Niraj Desai, Paul Gomez, John Halamka, Ken Huling, Shashank Jain, Melissa Kenny, Kevin Laitinen, Kellie Lucy, Krishna Nellutla, James B. Norman, Rob Piscitello, George Rakauskas, Jeff Rosen, Michele Sinunu, Franco Valentino, Marlon Violette, Griffin Weber, and Steve Wimberg.

### UCSF Development Team

The UCSF Profiles team includes Mini Kahlon, Eric Meeks, Kristine Moss, Rachael Sak, and Leslie Yuan. UCSF has developed innovative promotional strategies for research networking, assisted with quality assurance, and are adding OpenSocial support to Profiles RNS. Mini Kahlon was the co-chair with Griffin Weber of the National CTSA Research Networking Group, where they led an effort to create a national pilot that demonstrated interoperability among different research networking platforms.

### Recombinant Data Corp.

We thank the team at Recombinant Data Corp for their many years as an Authorized Support Provider for Profiles RNS: Kimber Barton, Nick Brown, Peter Emerson, Dan Housman, Mike Klumpenaar, Dave Legge, Mark Mischke, Matvey Palchuk, Chris Parisi, and Nancy Pickard.

### Profiles RNS Users Group

We thank the member institutions of the Profiles RNS Users Group for their willingness to be early adopters of the software and their continued feedback. For a list of member sites, please visit the Community page on <http://profiles.catalyst.harvard.edu>.