

# **ENM DICTIONARY**

RELEASE DATE:	13 July 2016
USE:	Terms used in project documentation
VERSION:	V.1.0.
AUTHORS:	Roland Grafström, Penny Nymark, Micha Rautenberg, Nina Jeliazkova, Philip Doganis, Georgia Tsiliki
PARTNER:	All eNanoMapper partners
CONTACT DETAILS:	http://www.enanomapper.net/contacts
LICENCE:	CC-BY 4.0







Term or abbreviation	Explanation
AMBIT.Js	A Java Script Library for interaction with AMBIT instances;
AMBIT	AMBIT ( <u>http://ambit.sf.net</u> ) is an open source web application for cheminformatics data management, modelling and read across. It offers graphical user interface and REST web services, including implementation of OpenTox API and eNanoMapper API extensions. The eNanoMapper database <u>http://data.enanomapper.net</u> runs a customized version of AMBIT.
ΑΡΙ	Application Programming Interface A way computer programs talk to one another. Can be understood in terms of how a programmer sends instructions between programs. The API specifies how software components should interact. A good API makes it easier to develop a program by providing all the building blocks. A programmer then puts the blocks together.
ArrayAnalysis.org	Offers user-friendly solutions for gene expression data analysis, from raw data to biological pathways. It contains modules of three types that can be launched individually or successively as an integrated workflow; QC and pre-processing, statistical analysis and pathway analysis.
ArrayExpress	A database with data from high-throughput functional genomics experiments stored in a standardized format, providing these data for reuse to the research community. It also includes data on nanomaterials. https://www.ebi.ac.uk/arrayexpress/
Bioclipse	A Java-based, open source advanced workbench for chem- and bioinformatics. It provides 2D-editing and 3D-visualization of molecules, proteins and sequences, calculation of chemical properties, QSAR and much more; all fully integrated into a user-friendly desktop application. <u>http://www.bioclipse.net/</u>
caNanoLab	<b>ca</b> ncer <b>Nano</b> technology <b>Lab</b> oratory <i>is a data sharing portal providing</i> <i>support for the annotation of nanomaterials with characterizations resulting</i> <i>from physico-chemical,</i> in vitro <i>and</i> in vivo <i>assays and the sharing of these</i> <i>characterizations and associated nanotechnology protocols in a secure fashion.</i> <u>https://cananolab.nci.nih.gov/caNanoLab/</u>
ChEMBL	A manually curated chemical database of bioactive molecules with drug-like properties developed at the European Molecular Biology Laboratory. It also contains data on nanomaterials. <u>https://www.ebi.ac.uk/chembl/</u>
Chipster	An open source user-friendly analysis software for omics and high-throughput data. It contains over 350 analysis tools for next generation sequencing (NGS), microarray, proteomics and sequence data. Users can save and share automatic analysis workflows, and visualize data interactively.
CTD	<b>Comparative Toxicogenomics Database</b> <i>is a manually curated robust,</i> <i>publicly available database providing information about chemical–gene/protein</i> <i>interactions, chemical–disease and gene–disease relationships, integrated with</i> <i>functional and pathway data to aid in development of hypotheses about the</i> <i>mechanisms underlying environmentally influenced diseases. It also contains</i> <i>data on nanomaterials.</i> <u>http://ctdbase.org/</u>
eNM	eNanoMapper – A Database and Ontology Framework for Nanomaterials Design and Safety Assessment (EU Project)
GEO	A database with data from high-throughput functional genomics experiments





	stored in a standardized format, providing these data for reuse to the research community. It also includes data on nanomaterials. http://www.ncbi.nlm.nih.gov/geo/
ISA-TAB	<ul> <li>The Investigation / Study / Assay tab - delimited format is a general purpose framework with which to collect and communicate complex metadata (i.e. sample characteristics, technologies used, type of measurements made) from experiments employing a combination of technologies. In particular, ISA - TAB has been developed for —but not limited to — experiments using genomics, transcriptomics, proteomics or metabol/nomics techniques.</li> <li>For example, consider an investigation into the effect of a compound that induces liver damage, which looks at changes in (i) the metabolite profile of urine and (ii) gene expression in the liver (by mass spectrometry and microarray technologies, respectively). The general motivation for this work is the fulfillment of the needs</li> </ul>
ISO 13528:2015	of two group. <u>http://www.isa-tools.org/</u> Statistical methods for use in proficiency testing by interlaboratory comparison; from International Standard Organization (ISO)
IUCLID5	International Uniform Chemical Information Database version 5 plays a central role in the IT environments of all organisations that have to cope with data submission requirements of REACH and other programs (OECD HPV, EU Biocides and others). Industry stakeholders, EU Member States, the European CHemicals Agency (ECHA), and any other interested party obtain the IUCLID installation kit from the web site; once installed, the local IUCLIDs are the essential tool to capture & store, submit, and exchange data on chemical substances stored according the format of the OECD Harmonised Templates <u>http://iuclid.eu/</u>
Jaqpot Quattro	An eNanoMapper developed web application that currently supports data preprocessing, as well as statistical, data mining and machine learning algorithms and methods for defining the Domain of Applicability of a predictive nQSAR model. It is an extension of the Jaqpot web application, which was originally developed during the OpenTox project.
JSON	JavaScript Object Notation is a lightweight data-interchange format. It is easy for humans to read and write. It is easy for machines to parse and generate. JSON is a text format that is completely language independent but uses conventions that are familiar to programmers of the C-family of languages, including C, C++, C#, Java, JavaScript, Perl, Python, and many others. It is based on a subset of the JavaScript Programming Language, Standard ECMA-262 3rd Edition - December 1999. http://json.org/
MOPAC	Molecular Orbital PACkage is a semiempirical quantum chemistry program for generating QSAR descriptors.
MySQL	My-Structural Query Language is an open-source relational database management system, enabling delivery of reliable, high-performance and scalable Web-based and embedded database applications. "My" is the name of daughter to one of the founders. <u>https://www.mysql.com/</u>
Nano-lazar	<b>Nano-Lazy</b> structure- <b>a</b> ctivity <b>r</b> elationships is an eNanoMapper developed tool for read across toxicity prediction ( <u>https://nano-lazar.in-silico.ch/predict</u> ).
NanoWiki	Originally developed as an internal knowledgebase of the toxicity of, primarily, metal oxides at the Karolinska Institutet and Maastricht University. Uses <u>https://semantic-mediawiki.org/</u> for data input. The data is exported as RDF





	dump , available at <u>http://dx.doi.org/10.6084/m9.figshare.1330208</u> .
	The NanoWiki RDF dump is converted with a custom parser and imported into the
	eNanoMapper database at <u>http://data.enanomapper.net</u>
OECD HT	The <b>OECD H</b> armonized <b>T</b> emplates (OHTs) are structured (XML) data
	formats for reporting safety-related studies on chemical substances. The OHTs
	<u>http://www.oecd.org/ehs/templates/</u> ) and the supporting IT tool (IUCLID5,
	<u>http://iuclid.eu</u> ) are used in a regulatory context, for preparation of substance
	dossiers for REACH and for other regulatory frameworks operating in Europe; as
	well as by the JRC NanoHub database
Ontology	A relational controlled vocabulary; deals with questions concerning what entities
enterogy	exist or may be said to exist, and how such entities may be grouped, related
	within a hierarchy, and subdivided according to similarities and differences.
OpenTox	A predictive toxicology framework with a unified access to toxicological data,
openiex	(Q)SAR models and supporting information developed under the grant agreement
	FP7-HEALTH-2007-A- 200787
Open Source	<b>Open-source</b> software (OSS) is computer software with its source code
	made available with a license in which the copyright holder provides the rights to
	study, change, and distribute the software to anyone and for any purpose. Open-
	source software may be developed in a collaborative public manner.
PathVisio	An open-source and free pathway analysis and pathway drawing software.
Patrivisio	Allows drawing, editing and analysis of biological pathways.
	http://www.pathvisio.org/
PROAST	An R-based software package that has been developed by RIVM for the statistical
PROASI	analysis of dose-response data. It can be used for 1) dose-response modelling, 2)
	deriving a Bench Mark Dose in human risk assessment, and 3) deriving an effect
	concentration in ecotoxicological risk assessment.
PubChem	
labellem	The <b>PubChem</b> Compound Database <i>contains validated chemical depiction</i>
	information provided to describe substances in PubChem Substance. Structures
	stored within PubChem Compounds are pre-clustered and cross-referenced by
	identity and similarity groups. It also contains data on nanomaterials.
	https://pubchem.ncbi.nlm.nih.gov/
RDF	The <b>R</b> esource <b>D</b> escription <b>F</b> ramework <i>is one of the core common</i>
	standards and data exchange formats of the Semantic Web. RDF it is based upon
	the idea of making statements about resources (in particular web resources) in
	the form of subject-predicate-object expressions. These expressions are known
	as triples in RDF terminology. <u>https://www.w3.org/RDF/</u>
	Example: "The sky has the color blue" in RDF is as the triple: a subject denoting
	"the sky", a predicate denoting "has", and an object denoting "the color blue".
REST	<b>Re</b> presentational state transfer is an abstraction of the architecture of the
	World Wide Web; more precisely, REST is an architectural style consisting of a
	coordinated set of architectural constraints applied to components, connectors,
	and data elements, within a distributed hypermedia system. REST ignores the
	details of component implementation and protocol syntax in order to focus on
	the roles of components, the constraints upon their interaction with other
	components, and their interpretation of significant data elements. OpenTox web
	services are based on the REST architecture.
RO	Research Object are semantically rich aggregations of resources that bring
	together data, methods and people in scientific investigations.





RRegrs	An eNanoMapper developed tool. It is an R package based on a collection of regression tools from R that can be used to search the best regression models for any dataset. The initial use of the script is aimed at finding QSAR models for chemoinformatics / nanotoxicology.
SPARQL	Simple Protocol And RDF Query Language is a semantic query language for databases, able to retrieve and manipulate data stored in RDF format.
ToxBank	A dedicated web-based warehouse for toxicity data management and modelling, a 'gold standards' compound database and repository of selected test compounds, and a reference resource for cells, cell lines and tissues of relevance for in vitro systemic toxicity research carried out across the FP7 HEALTH.2010.4.2.9 Alternative Testing Strategies SEURAT program.

Abbreviation	Explanation
1D	1-Dimensional
3D	3-Dimensional
A&A	Authentication and Authorisation
AAI	Authentication and Authorisation Interface
ANOVA	Analysis of Variance
ARFF	Attribute-Relation File Format
BAO	BioAssay Ontology
bao (namespace)	http://www.bioassayontology.org/bao#
BFO	Basic Formal Ontology
BMD	Benchmark Dose
BP	Biological Processes
CC	Cellular Components
CC	Creative Commons
CCZero	Creative Commons Zero waiver
CDK	Chemistry Development Kit
ChEBI	Chemical Entities of Biological Interest
CHEMINF	Chemical Information Ontology
CL	Cell Ontology
clo (namespace)	http://purl.obolibrary.org/obo/
CML	Chemical Markup Language
CMS	Content Management System
CSV	Comma-separated values
CV	Cross-Validation
DB	Database
DC	Dublin Core Metadata Ontology
DFT	Density Functional Theory
DoA	Domain of Applicability
DoW	Description of Work
EC	European Commission
EHS	Environment and Health Safety
EJB	Enterprise JavaBeans
EN	Elastic Net
ENET	Elastic Net regression





ENM	Engineered Nano Materials
ENPRA	Risk Assessment Of Engineered Nanoparticles (EU Project)
ENVO	Environment Ontology
EU	European Union
FFNN	Feed Forward Neural Network
GLM	Generalized Linear Model with Stepwise Feature Selection
GNU GPL	GNU General Public License
GO	Gene Ontology
GPW	Gaussian and plane waves
GSEA	Gene Set Enrichment Analysis
GUI	Graphical User Interface
HC	Hierarchical Clustering
HF	Hartree-Fock
НОМО	Highest Occupied Molecular Orbital
HTTP	Hypertext Transfer Protocol
IAO	Information Artifact Ontology
ID3	Iterative Dichotomiser 3
IRI	International Resource Identifier
JPDI	Jaqpot Protocol of Data Interchange JQ Jaqpot Quattro
JRC	Joint Research Centre https://ec.europa.eu/jrc/
JSF	JavaServer Faces
KEGG	Kyoto Encyclopedia of Genes and Genomes
kNN	
KS-DFT	K Nearest Neighbour
	Kohn-Sham density functional theory
	Least Absolute Shrinkage and Selection Operator
LC-MS/MS	Liquid chromatography–Mass Spectrometry/ Mass Spectrometry
	Lightweight Directory Access Protocol
LM	Linear Model
L00	Leave-One-Out
	Leave one out cross validation
LUMO	Lowest Unoccupied Molecular Orbital
MARINA	Managing Risks of Nanomaterials (EU Project)
MeOx	Metal oxides
MF	Molecular Functions
MIREOT	Minimum Information to Reference an External Ontology Term
MLR	Multiple Linear Regression
MODENA	Modelling Nanomaterial Toxicity (EU COST Action)
ModNanoTox	Modelling nanoparticle toxicity: principles, methods, novel approaches (EU Project)
MOPAC	Molecular Orbital PACkage
MS	Mass Spectrometry
MSigDB	Molecular Signature Databases
	Biological Foundation for the Safety Classification of Engineered Nanomaterials
NANOSOLUTIONS	(ENM): Systems Biology Approaches to Understand Interactions of ENM with
	Living Organisms and the Environment (EU Project)
NANoPUZZLES	Nanoparticles safety collaboration platform
nanoQSAR	Nano- Quantitative Structure-Activity Relationship
NANoREG	A common European approach to the regulatory testing of Manufactured

eNanoMapper
-------------

July 2016 ENM DICTIONARY





	Nanomaterials (EU Project)
NCBI	National Center for Biotechnology Information
NM	Nanomaterial
NN	Neural Networks
NPO	NanoParticle Ontology
NPs	Nanoparticles
nQSAR	Nano- Quantitative Structure-Activity Relationship
NSC	NanoSafety Cluster
OBI	Ontology for Biomedical Investigations
OECD	Organisation for Economic Co-operation and Development
OLS	Ordinary Least Squares
013	
	OpenAM is an open source access management, entitlements and federation
OpenAM	server platform http://en.wikipedia.org/wiki/OpenAM. OpenAM is the software
OpenAM	that allows OpenTox services running at each partner facility to use one and the
	same user credentials (user name and password) this is known as Single Sign On
	capability.
OWL	Web Ontology Language, a family of knowledge representation languages for
DATO	authoring ontologies
PATO	Phenotype and Trait Ontology
PLS	Partial Least Squares
PMML	Predictive Model Markup Language
PR	Protein Ontology (alternative abbreviation)
PRO	Protein Ontology
PubMed	US National Library of Medicine National Institutes of Health
QC	Quality Control
QM	Quantum Mechanics
QSAR	Quantitative Structure-Activity Relationship
RA	Risk Assessment
RBF	Radial Basis Function
RBF-DDA	Radial Basis Dynamic Decay Adjustment
RF	Random Forest
RFE	Recursive Feature Elemination
RMSE	Root Mean Square Error
RO	Relations Ontology
RO	Research Object
RSS	Rich Site Summary
SAB	Scientific Advisory Board
SDF	Structure Data File
SME	Small and Medium Enterprise
SEO	
SMILES	Search Engine Optimization
	Simplified molecular-input line-entry system
SMW	Semantic MediaWiki
SOP	Standard Operating Procedure
SPARQL	SPARQL Protocol and RDF Query Language (recursive acronym)
SSL	Secure Sockets Layer
SSO	Single Sign-On
SUN	Sustainable Nanotechnologies (EU Project)
SUNDS	Sustainable Nanotechnologies Decision Support System





SVM	Support Vector Machines
SVN	SubVersioN
SVR	Support Vector Regression
TEM	Transmission Electron Microscopy
TLS	Transport Layer Security
UI	User Interface
UniProt	Universal Protein Database
UO	Unit Ontology
URI	Uniform Resource Identifier
URL	Uniform Resource Locator
UX Design	User eXperience Design
VIP	Very Important Person
VIP	Variable Importance to Projection
WP	Work Package
WS	Web Services
XLS	eXceL Spreadsheet
XML	eXtensible Markup Language
Xvfb	X virtual framebuffer