



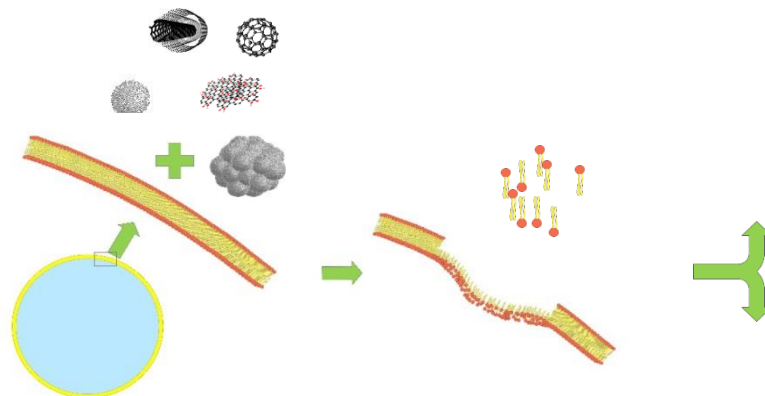
International workshop on

COMPUTATIONAL APPROACHES IN NANOSCIENCES

„COMPInNANO“

Ljubljana, 2nd and 3rd October 2015

PROCEEDINGS/ZBORNIK PREDSTAVITEV



	AOI	BYI	COI	DOI	EVI
Local Items	CHI	vanilium	SE	absorption	SE
Units	ug/ml				
Continents	0.048	-0.54873	0.93747	-0.30347	2.19615
2	0.095	3.78827	0.89346	-0.79767	3.13791
3	0.143	-1.14892	0.86192	1.4673	3.00195
4	0.19	2.41315	2.19399	1.03939	4.30036
5	0.238	-0.47774	1.48005	0.63919	4.06539
6	0.285	1.48045	1.16392	2.4172	2.76555
7	0.333	2.62165	1.01524	0.27144	1.11283
8	0.381	2.01816	0.36998	0.60213	2.11892
9	0.428	2.62389	1.1475	0.38447	0.80264
10	0.475	0.43487	1.31501	0.67781	0.65301
11	0.48	0.33899	1.39373	0.84727	1.66427
12	0.95	0.80964	0.78264	14.84744	3.10073
13	1.43	11.89043	1.7668	12.36747	2.81408
14	1.9	6.03273	1.18687	18.48617	2.4661
15	2.38	9.89952	3.5019	28.87679	5.89309
16	2.86	14.10287	18.81671	27.47116	3.87934
17	3.33	23.78387	2.266	18.842	1.77772
18	3.81	18.91839	4.8302	24.45538	4.27895
19	4.28	26.15495	0.39201	23.131	6.6224
20	4.76	32.48231	5.13691	28.07863	3.57394
21	4.8	22.24332	3.89306	24.45968	1.47634
22	5.5	31.87978	2.53889	38.09026	0.66953
23	14.3	41.68165	3.92339	49.3272	2.07146
24	19	44.29513	8.44832	42.33348	3.25218
25	23.8	43.87283	1.05972	35.16602	4.5489
26	28.4	55.25453	4.72205	48.30661	1.08666
27	33.3	57.37951	2.00002	44.95186	2.48287
28	38.1	55.09568	1.66513	52.41895	5.37739
29	42.8	55.10879	0.32342	49.7806	3.07665
30	47.6	60.00601	1.29428	47.76005	3.8844

International workshop/Mednarodni sestanek
COMPUTATIONAL APPROACHES IN NANOSCIENCES: „COMPInNANO“

Ljubljana, 2nd and 3rd October 2015

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INTERNATIONAL Workshop on Computational Approaches in Nanosciences (2015 ; Ljubljana)
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1. Jemec, Anita
283801856

Foreword

The production of nanomaterials has led to growing public and regulatory concern about their safety. The aim of a number of ongoing or past EU projects has been to address innovative, safe and sustainable nano-enabled products. The NanoValid project, as one of them, has developed a set of reliable reference methods and materials for their fabrication, physicochemical characterization, hazard identification and exposure assessment. However, since the experimental toxicological testing of nanomaterials is costly and time-consuming, it is necessary to develop a new approach based on knowledge, methods and tools to reach the goal of predictive nanotoxicology.

The workshop “Computational approaches in Nanosciences” hosted the participants from three large scale EU FP7 projects NanoValid, NanoMile, and Modern in order to disseminate the results on computational methods for NP hazard characterisation and exchange ideas with representatives from regulatory bodies and industry. Modellers, computational scientists, experimental (eco)toxicologists and experts from different fields of nanoscience discussed the future of *Computational Approaches in Nanosciences*.

The participants agreed that the most crucial points in the data sharing among academia, industry and regulatory bodies are (i) the organisation of data in large data sets and synchronised communication between different fields and sectors. Calibration and validation of computational models is impossible without utilizing high quality experimental data. Therefore, close collaboration between the computational chemists and experimentalists from different areas (i.e. toxicologists, specialists on characterization) would be crucial for the success.

The NanoValid project has significantly contributed toward defining the quality of nanotoxicity data and harmonization of test. In the future they have to be integrated into databases and shared. Namely, new approaches for the data gap filling are needed which have to be dynamic and consider scientific aspects and new developments in nano-sciences. The participants concluded that the future of nanosafety will rely on both, experimental data and computational methods but we have to adopt successful communication strategies.



Friday, 2nd October

16.15 – 16.30

Welcome speech & Introduction to the workshop
Prof. Dr. Damjana Drobne, Assist. Prof. Anita Jemec

TOPIC 1: ORGANISATION OF LARGE DATA SETS FOR MODELLING IN GENERAL

The topic will deal with the organisation of large data sets using different statistical approaches.

16.30 – 17.00

Assist. Prof. Ddr. David Bogataj, The Mediterranean Institute for Advanced Studies:
„Reproducible Research“

17.00 – 17.30

Assist. Prof. Dr. Cene Fišer, Biotechnical Faculty University of Ljubljana, Slovenia:
„Web-databases in systematics and taxonomy“

17.30 – 18.00

COFFEE BREAK

TOPIC 2: DATA SHARING-Academia, Industry and Regulatory bodies

The stakeholders will present their view on the nanoscience data sharing.

18.00 – 18.30

Mag. Vladimir Vrečko, Cinkarna Celje (TiO₂ producers), Slovenia:
„The importance of data interpretation and dissemination for the future of nanotechnology.“

18.30 – 19.00

Dr. Mojca Kos Durjava, National Laboratory of Health, Environment and Food
and **Mag. Karmen Krajnc**, Chemical Office of the Republic of Slovenia, Slovenia:
„Challenges of Regulation and Risk Assessment of Nanomaterials“

Saturday, 3rd October

TOPIC 3: QUANTITATIVE NANOSTRUCTURE-ACTIVITY (PROPERTY) RELATIONSHIPS (QNAAR, QNPR) MODELLING

9.00 – 9.30 **Prof. Dr. Marjan Vračko**, National Institute of Chemistry, Slovenia:
„Chemometric analysis and QSAR modelling in NANO-toxicology“

9.30 – 10.00 **Dr. Villem Aruoja**, National Institute of Chemical Physics and Biophysics, Estonia:
„Toxicity of metal oxide nanoparticles to algae, bacteria and protozoa:
FP7 project MODERN“

10.00 – 10.30

COFFEE BREAK and poster session

TOPIC 4: MODELLING OF INTERACTIONS BETWEEN NANOMATERIALS AND BIOLOGICAL MODELS

Presentation of different databases aimed to collect information about Nanosafety related topics.

10.30 – 11.00 **Prof. Dr. Alok Dhawan**, CSIR-Indian Institute of Toxicology Research, Lucknow, India:
„Toxicity of Nanomaterials: The need for Novel Computational Tools and Approaches in Safety Assessment“

11.00 – 11.30 **Dr. Lokesh Baweja**, Institute of Life Sciences, Ahmedabad, Gujarat, India:
„Computational approaches to understand the interaction of nanomaterials with biomolecules“

11.30 – 12.00 **Dr. Fabrice Carnal**, Institut F.A. Forel, University of Geneva, Switzerland:
„Monte Carlo Modelling of Interaction Processes between Nanoparticles and Biomacromolecules of Variable Hydrophobicity“

12.00 – 12.30

Maja Sopotnik and Prof. Dr. Kristina Sepčič, Biotechnical Faculty, University of Ljubljana, Slovenia:
„Interaction of carbon-based nanomaterials with cholinesterases and serum proteins“

12.30 – 13.30

LUNCH BREAK and poster session

Saturday, 3rd October

TOPIC 5: ORGANISING DATA FOR RISK ASSESSMENT/DATA QUALITY

- 13.30 – 14.00 **Prof. Dr. Rishi Shanker**, Institute of Life Sciences, Ahmedabad, Gujarat, India:
„Tracking Nano-footprints in microbial food chain: Observations, Data & Challenges“
- 14.00 – 14.30 **Dr. Ashutosh Kumar**, Institute of Life Sciences, Ahmedabad, Gujarat, India:
„Importance of *in silico* approaches in understanding the interactions of nanoparticles with biological membrane“
- 14.30 – 15.00 **Assist. Prof. Dr. Anita Jemec**, Biotechnical Faculty, University of Ljubljana, Slovenia:
„Quality of nanotoxicity data and the importance of harmonization“
- 15.00 – 16.00 COFFEE BREAK, poster session and open discussion

REPRODUCIBLE RESEARCH

ddr. David Bogataj

MEDIFAS

Reproducible research

Reproducible research is the idea that data analyses, and more generally, scientific claims, are published with their data and software code so that others may verify the findings and build upon them.

The goal of reproducible research is to tie specific instructions to data analysis and experimental data so that scholarship can be recreated, better understood and verified.

In the instructions the complex of conditions should be described precisely.

Replication

The ultimate standard for strengthening scientific evidence is replication of findings and conducting studies with independent:

- Investigators
- Data
- Analytical methods
- Instruments

Replication

- Replication is particularly important in studies that can impact broad policy or regulation decisions

What is Reproducible research?

- Scientific Question
- Research Protocol
- Nature (complex of conditions, entities and their relations)
- **1. Measured data (dataset regarding realization of an experiment – raw data)**
 - Data processing code
- **2. Analytic data (cleansed data)**
 - Analytical code
- **Computational results**
 - Presentation code (figures, tables, Numerical Summaries)
- Published article (text)

Up to now in published articles only short overview of data and code (1 and 2) is presented which is not enough.

What problem does Reproducibility solve

- Black Box Problem
- Transparency
- Data Availability
- Software/Methods Availability
- Improved Transfer of Knowledge

Reproducibility

- The premise of reproducible research is that with data/code available, researchers can check each other and the whole system is self-correcting
- Addresses the most “downstream” aspect of the research process – post-publication
- Assumes everyone plays by the same rules and wants to achieve the same goals (i.e. scientific discovery)

Who Reproduces Research?

For reproducibility to be effective as means to check validity, someone needs to do something

- Re-run analysis; check if results match
- Check the code for bugs/errors
- Try alternate approaches/check sensitivity

Validation of the data analysis

Availability:

- Description of complex of conditions in which the experiment run
- Data
- Algorithms

Reasons:

- Other people can run the same algorithms on the sama data and can come to the same conclusions as the researcher

What you do not have

- Independent data
- Independent method

Impact of new technologies

- Allow us to collect data at much higher throughput
- We can get very complex high dimensional datasets almost instantaneously
- Computing power that allows us to merge databases in even bigger „megadatabases“

Minimum standards

- Experiment(observation) should be described in exactly known complex of conditions.
- If the complex of conditions is changed (environment), realization of the experiment can give different results.
- It could lead to different conclusions.

Problems

- Authors must undertake considerable effort to put data/results on the web
- Readers must download data/results individually and piece together which data go with which code section, etc
- Authors/readers must manually interact with websites
- There is no single document to integrate data analysis with textual representations; data, code, and text are not linked

Simplify the process

Put the data and the code together in the same document

- People can execute code in the right order
- Data are read at the right times

Document integrates data analysis with all the textual representations (descriptions) that everything is linked together

How do I Make My Work Reproducible

- Decide to do it (ideally from the start)
- Keep track of things, perhaps with versioning control system to track snapshots/changes
- Use statistical software whose operation can be coded (R)
- Don't save output (store raw dataset)
- Save raw data and the process that get you there
- Save data in non-proprietary formats (ASCII)

Single research report document

- To document the analysis and
- to have the code of the analysis in the same document

Documentation of total observation and research process

- Documentation preparation system – description of research process including metadata (LaTeX)
- Description of complex of conditions in which the experiment run
- Raw data - dataset
- Code – Programming Language (R) for data analysis, statistics, forecasting, optimization, sensitivity analysis and simulations.

Literate Statistical Programming with knitr (R)

- Text and code all in one place, logical order
- Data where results of observations are automatically updated that reflect external changes
- Code is live – you need to run code. When error appears it needs to be resolved.

CONCLUSION

1.- Reproducibility brings transparency (wrt code+data) and increases transfer of knowledge. Therefore complex of conditions which describe the environment for observation procedures have to be very clearly described.

2.- Important currant discussion is about how to convince researchers to share data. The owners of data should have incentives to publish the dataset, metadata and code

-by the system similar as IF evaluation (citation index) of their publications and

- market system like available by ScienceDirect, Springer and others where the databases are able to buy or sell if there is not obligatory open system.

- The founder of a research should clearly determine which data should be publicly available and which are available on the data market - for sale.

Web-databases in systematics and taxonomy

Cene Fišer

SubBio Lab

Odd. za Biologijo

Biotehniška Fakulteta

Univerza v Ljubljani

Univerza v Ljubljani



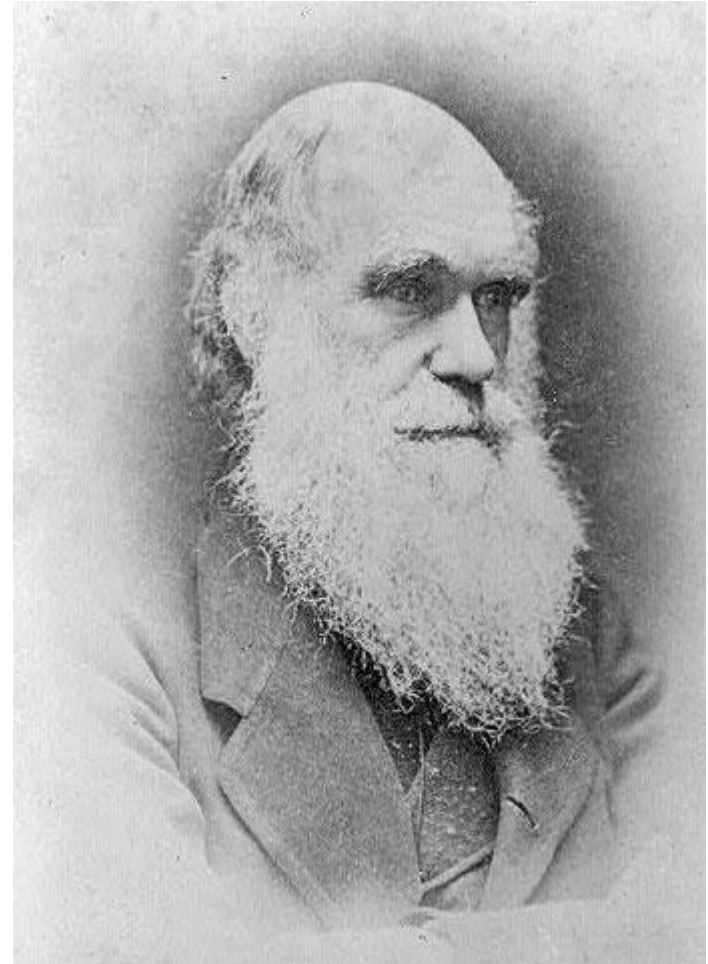
Ljubljana, 2.10.2015

taxonomy-systematics?

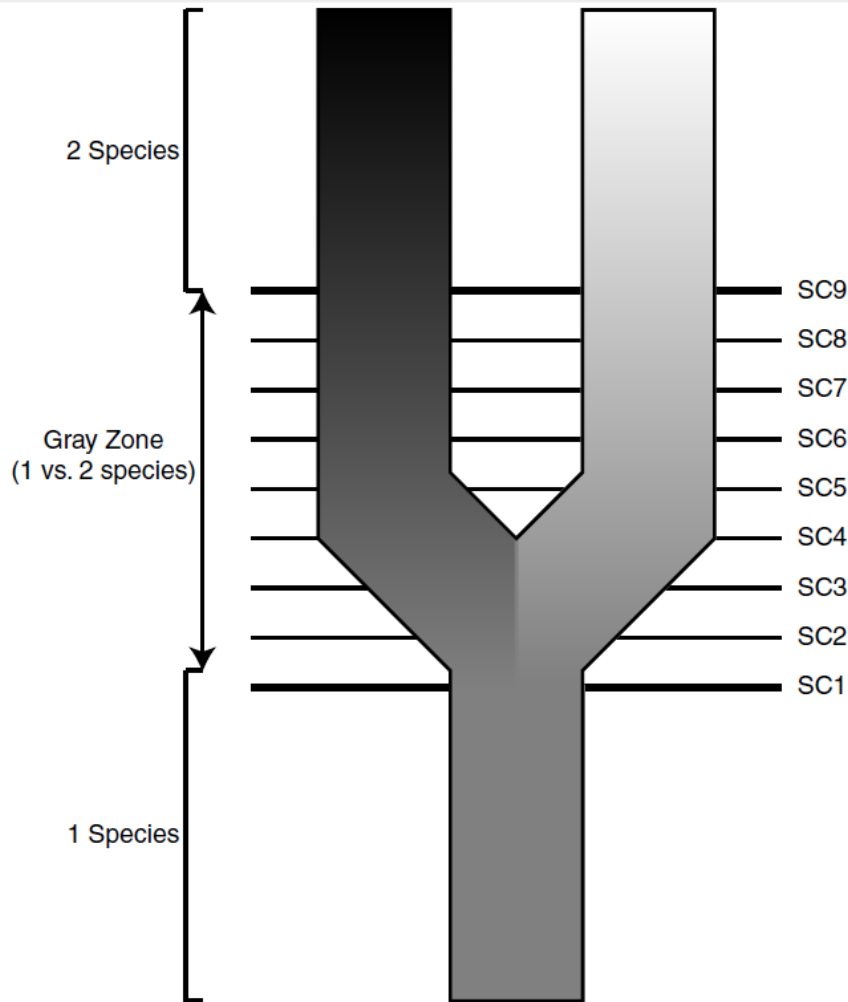
1. Taxonomy in a narrow sense: describing and naming species.
2. Taxonomy in a broad sense: species identification, research of species biology, distribution and conservation.
3. Systematics: inference of species relatedness and hierarchical categorization of species in higher taxonomic categories.



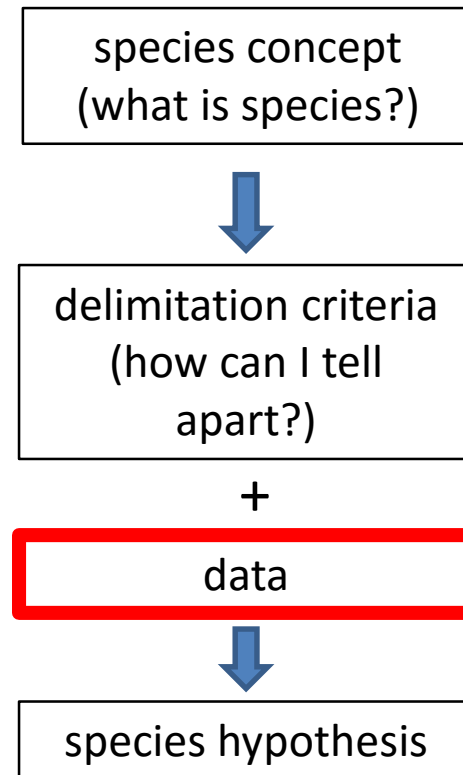
an ancient
science?



Modern synthesis: grounded in evolutionary theory....

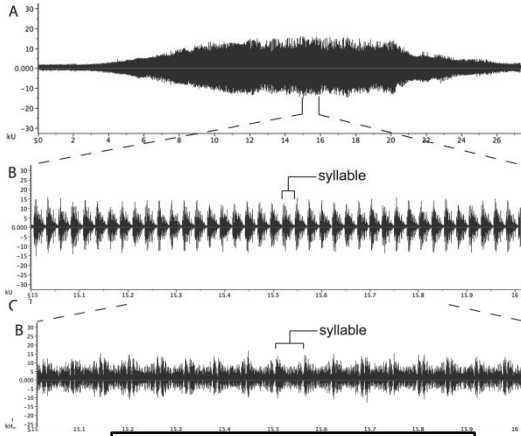


Speciation: a process of divergence



De Queiroz, Kevin (2007) 'Species Concepts and Species Delimitation', *Systematic Biology*, 56:6, 879 - 886

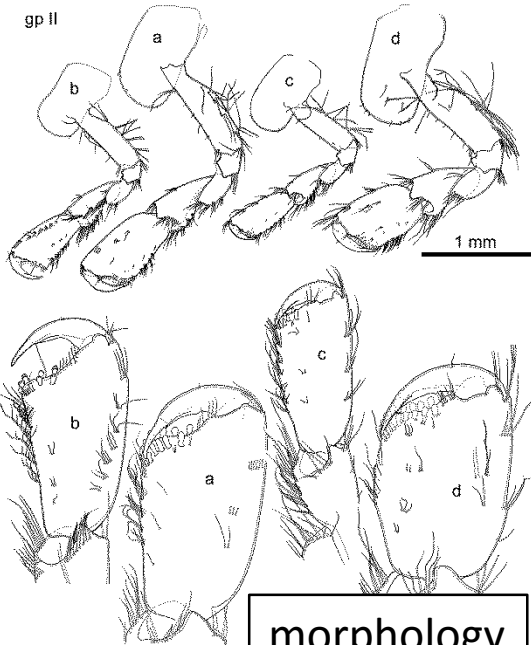
....interdisciplinary science



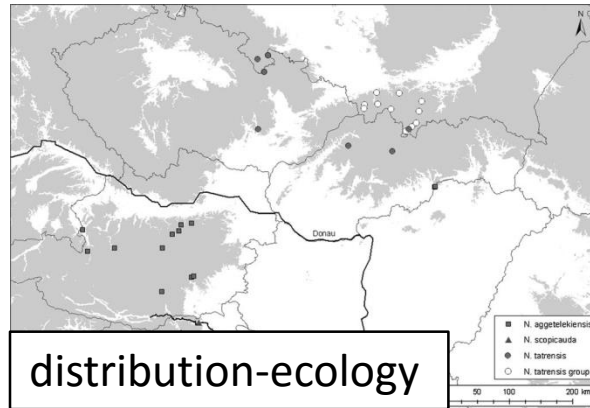
communication

TYPE: NO LIMITS

LIMITS IN DATA
DEPOSITION AND
RETRIVAL



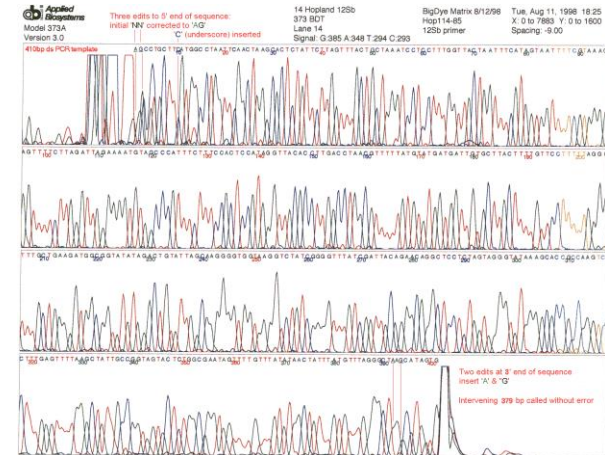
morphology



distribution-ecology



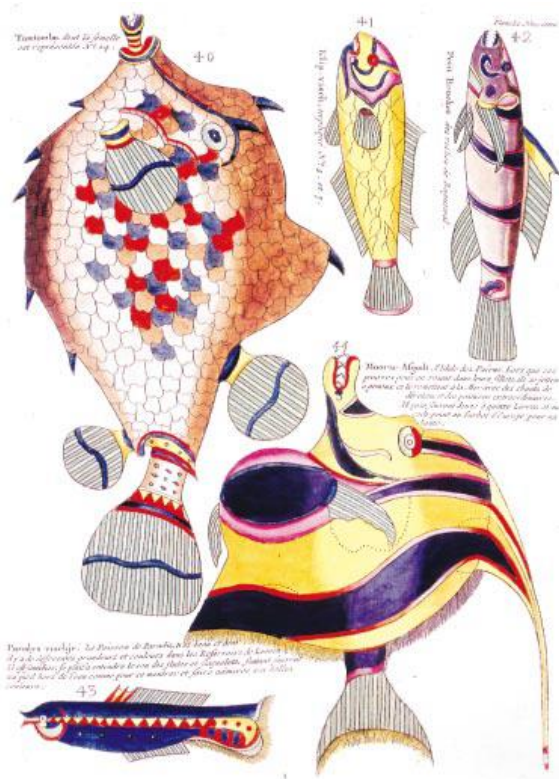
mating behavior



molecular data

Challenges for taxonomy

The discipline will have to reinvent itself if it is to survive and flourish.



This discipline is made for the web: it is information-rich and often requires copious illustrations.

Godfray, H.C.J. 2002 **Challenges for taxonomy.** *Nature* 417, 17-19 (Commentary; reply to correspondence arising: “Towards taxonomy’s ‘glorious revolution’”

Taxon specialized websites: not necessarily databases

Make your own website:
Scratchpads:

<http://scratchpads.eu/>

The screenshot shows the Scratchpads website interface. The header includes the logo and navigation links: Home, News, Explore, Support, Develop, and About us. A sidebar on the left lists various categories like 'Why choose Scratchpads?', 'Concept', 'Policies', 'Partner projects', 'Sponsors', 'Resources', 'Team', and 'How to cite'. The main content area is titled 'Publish your data online' and features a 'PUBLISH' button on a laptop icon. Below this, a paragraph explains that Scratchpads are used for publishing data on the web, constructing databases, or assembling manuscripts. A table titled 'Overview of features:' lists several benefits with checkmarks in the right column:

Feature	Description
Share your data across communities	Scratchpad lets you connect with your community. ✓
Search engine friendly	All Scratchpads sites are search engine friendly. Scratchpads sites usually rank on top in Google search results when relevant taxa and keywords are used. ✓
Publish your content online	Scratchpads lets you create your content online. ✓
Publish in peer reviewed Journals	Submit original data present in your scratchpad directly to peer-reviewed journals for publication. ✓

This screenshot shows a different view of the Scratchpads website. The header is consistent with the previous image. The main content area is titled 'Why choose Scratchpads?' and features four columns of text with corresponding images: 'Create your own site', 'Publish your data online', 'Contribute to biodiversity knowledge', and 'Collaborate with your peers'. Below this, there is a 'Join the community!' section with links for 'Create a Scratchpad', 'Join a Scratchpad', 'Register for training', 'Try our sandbox', 'Become a Scratchpads ambassador', 'Develop Scratchpads source code', and 'Read about our policies'. To the right, there is a 'Case studies' section featuring a 'DEST' case study with a small image. Below that is a 'Feeds' section with buttons for 'News' and 'Twitter'. At the bottom, there is a 'Scratchpads around the world' section with a globe icon and text indicating the number of active users and pages. The footer contains logos for the Natural Environment Research Council, eMonocot, Vibrant, and the Natural History Museum, along with a detailed navigation menu and footer links for 'Website feedback', 'Terms & Conditions', 'Privacy Policy', and 'Cookies'.

...moving to level of an individual : increase in strength of analyses

**Analysis I :
moleclar/morphological
differentiation**

downolad sequences,
quantitative or
continous data

+

newly measured
data



analysis

**Analysis II :
spatial circumstances of
speciation**

downolad spatial
data

+

geomorphological
characteristcs



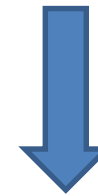
analysis gene-flow
barries

**Analysis III :
ecological niche
reconstruction**

downolad spatial
data

+

Climate data
(Hijmans et al. 2005,
J.climat)



analysis of ecological
similarity

Taxon specific interactive database

SubBio Database: interactive database of subterranean fauna

Three main sources of data:

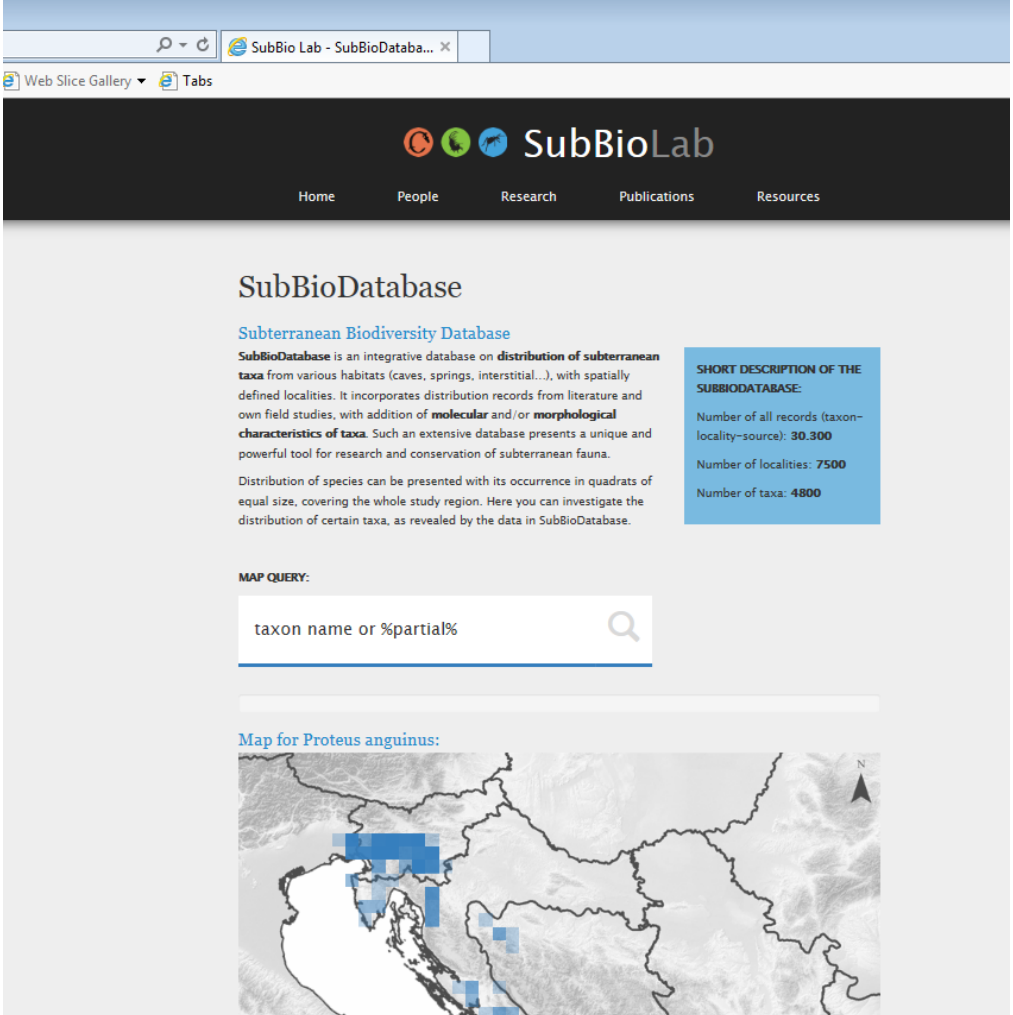
- species lists
- distributional data
- DNA sequences

Sources of data:

- published sources
- own data

Quantity:

- number of records:
30.300
- number of localities :
7.500
- number of taxa: 4.800



The screenshot shows the SubBioLab website interface. The browser address bar displays "SubBio Lab - SubBioDataba...". The website header includes the SubBioLab logo and navigation links: Home, People, Research, Publications, and Resources. The main content area is titled "SubBioDatabase" and describes it as a "Subterranean Biodiversity Database". It provides a short description of the database, stating it is an integrative database on the distribution of subterranean taxa from various habitats (caves, springs, interstitial...), with spatially defined localities. It incorporates distribution records from literature and own field studies, with addition of molecular and/or morphological characteristics of taxa. Such an extensive database presents a unique and powerful tool for research and conservation of subterranean fauna. The distribution of species can be presented with its occurrence in quadrats of equal size, covering the whole study region. Here you can investigate the distribution of certain taxa, as revealed by the data in SubBioDatabase.

MAP QUERY:

taxon name or %partial%

Map for *Proteus anguinus*:

The map shows the distribution of *Proteus anguinus* in the Balkans region, with blue squares indicating the presence of the species in various quadrats. A north arrow is visible in the top right corner of the map.

SHORT DESCRIPTION OF THE SUBBIODATABASE:

- Number of all records (taxon-locality-source): 30.300
- Number of localities: 7500
- Number of taxa: 4800

Taxon specific interactive database

SubBio Database: interactive database of subterranean fauna

The screenshot displays the SubBioDatabase access web application in a browser window. The browser's address bar shows the URL <http://subbio.net/dbaccess/dbquery/>. The application header includes the logo and text "SubBioDatabase access".

The main interface is organized into several sections:

- Search:** A search bar with a dropdown menu for "Locality" and a "Search" button.
- Advanced DB Query:** A section with a "DB-QUERY" button.
- Add:** Buttons for "+ New Locality", "+ New Literature", and "+ Multiple DNA entries".
- View:** A "Multi-Survey" button.
- Log in / log out** and **Edit profile** links.

The central content area is titled "Tabele / Polja" and contains six panels, each with a list of fields:

- Literatura:** ID literature, Kratka oznaka literature, Podrobna oznaka literature, Avtorji, Leto objave, Naslov, Revija / Knjiga, Vol / Issue, Strani, Opombe, Oznaka - fascikli.
- Lokaliteta:** ID lokalitete, Ime lokalitete, Kraj ob, Vešji kraj, Drugo ime / Sinonimi, Država, Geog. dolžina (WGS_X), Geog. širina (WGS_Y), Koordinate po, Nadmorska višina, UTM kvadrant, Izvor koordinat, Natančnost 1, Natančnost 2, Katastrska številka jame.
- Popis:** ID popisa (intern), Št. popisnega lista, Popisovalci, Datum popisa, Opombe (opažanja ob obisku lokalitete), Trajanje - OD, Trajanje - DO, Vir (če iz tuje zbirke), Namen, Metode, Vzorec za prebiranje, Opis vzorca, Pasti (vzorčenje s pastmi).
- Sistem:** Takson ID, Celotno ime taksona, Sinonimi, Podvrsta, Vrsta, Podrod, Rod, Podružina, Družina, Red, Razred, Avtor - ime, Avtor - leto.
- Voucherji / DNA:** Voucher ID, Voucher - stara koda, Št. vrsta voucherskega osebk, Opombe k voucherskemu osebk, DNA ID, Datum izolacije, Št. vrsta DNA, Gen, Fwd, Rev, PCR, Sekvenca, Accession num., Dnevnik.
- Opaženi taksoni / Zbirka:** PopisDetalji ID, Izvorna določitev (Takson_original), Locus typicus, Določevalci, Št. opaženih osebkov, Št. vzetih osebkov, V vodi / Na kopnem, Mikrobioti, Metoda vzorčenja, Opombe (za posamezen opažen takson).

At the bottom of the main content area, there are buttons for "Dodaj izbrana polja +", "Odstrani -", "Izvrši poizvedbo", and "Izvrši poizvedbo".

The Windows taskbar at the bottom shows the system clock as 20:53 on 1.10.2015, along with various application icons and system tray icons.

Taxon specific interactive database

SubBio Database: interactive database of subterranean fauna

Niphargus_DNABaza_23maj14 - Microsoft Excel

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R
	Celotno_ime	DNA_id	Datum_izolacije	created	voucher_id	shramba	gen	fwd	rev	PCR	sekvenca	acc_num	dnevnik	Lokalit_ime	Drzava	WGS_Xdd	WGS_Ydd	
1	Microniphargus leruthi	1532	15/05/2011		0 NA741	stiroporna škat							Laboratorijski	Velika Britanija	UK			
2	Microniphargus sp.	1531	15/05/2011		0 NA740	stiroporna škat							Laboratorijski	Jama Sweetwater	Great Britain	-3,535975282	50,62151447	
3	Microniphargus sp.	1524	03/12/2010		0 NA649	stiroporna škat				neuspešno			Laboratorijski	Borehole at Wind	Great Britain	-2,376686748	50,72602791	
4	Microniphargus sp.	1528	22/02/2011		0 NA713	stiroporna škat							Laboratorijski	Borehole at Wind	Great Britain	-2,376686748	50,72602791	
5	Microniphargus sp.	1670	26/07/2012		0 NB090	stiroporna škat							Laboratorijski	Grotte de Combla	Belgium	5,57481	50,4754052	
6	Microniphargus sp.	1525	22/02/2011		0 NA710	stiroporna škat							Laboratorijski	Abime de Combla	Belgium	5,57481	50,4754052	
7	Microniphargus sp.	1526	22/02/2011		0 NA711	stiroporna škat							Laboratorijski	Abime de Combla	Belgium	5,57481	50,4754052	
8	Microniphargus sp.	1527	22/02/2011		0 NA712	stiroporna škat							Laboratorijski	Abime de Combla	Belgium	5,57481	50,4754052	
9	Niphargus aberrans	1229		1309432543	NA025		28S	28Slev3	28Sdes5	OK	AAGGCTTCAGTA			Sneberje, freatske	SLO	14,57311944	46,08042778	
10	Niphargus aberrans	1230		1309432543	NA025		H3	H3aF2	H3aR2	PONOVNO V				Sneberje, freatske	SLO	14,57311944	46,08042778	
11	Niphargus aberrans	1228		1309432543	NA025		28S	28Slev2	28Sdes2	OK	GAAAAGCACTCT EF617260			Sneberje, freatske	SLO	14,57311944	46,08042778	
12	Niphargus aggtelekiensis	1298		1309432543	NA501		28S	28Slev3	28Sdes5	OK	AAGGCTTCAGTA			Gesause ET2B	Austria	14,587258	47,565345	
13	Niphargus aggtelekiensis	1297		1309432543	NA501		28S	28Slev2	28Sdes2	OK	GAAAAGCACTCT KJ566697			Gesause ET2B	Austria	14,587258	47,565345	
14	Niphargus aggtelekiensis	1299		1309432543	NA501		H3	H3aF2	H3aR2	OK	TGCTCGGAAATC KJ566722			Gesause ET2B	Austria	14,587258	47,565345	
15	Niphargus aggtelekiensis	2119	12.08.2013	1376490795	NB541	skrinja na hodr							dnevnik Doroth	Rakoczi 1 cave	Hungary	20,74886667	48,52081667	
16	Niphargus aggtelekiensis	2120	12.08.2013	1376491015	NB542	skrinja na hodr							dnevnik Doroth	Baradla-also Cav	Hungary	20,54401667	48,4831	
17	Niphargus alpinus	1003		1309432543	NA019		28S	28Slev2	28Sdes2	OK	GAAAAGCACTCT			Ramsauer Tal, No NEM		12,900112	47,616669	
18	Niphargus alpinus	1002		1309432543	NA109		28S	28Slev3	28Sdes5	OK	AAGGCTTCAGTA			Schapbach spring	NEM	12,95833333	47,58194444	
19	Niphargus alpinus	1001		1309432543	NA109		28S	28Slev2	28Sdes2	OK	GAAAAGCACTCT EF617254			Schapbach spring	NEM	12,95833333	47,58194444	
20	Niphargus ambulator	1290		1309432543	NA504		28S	28Slev2	28Sdes2	OK	GAAAAGCACTCT KJ566699			Buco del Piombo	ITA	9,198473	45,825819	
21	Niphargus ambulator	1291		1309432543	NA504		H3	H3aF2	H3aR2	OK	CGCTCGTAAGTC KJ566723			Buco del Piombo	ITA	9,198473	45,825819	
22	Niphargus angelieri	1218		1309432543	NA200		28S	28Slev2	28Sdes2	NE DELA				Grotte des Fees	FRA	3,028236	42,910091	
23	Niphargus angelieri	1220		1309432543	NA200		H3	H3aF2	H3aR2	OK	CGCTCGCAAGTC			Grotte des Fees	FRA	3,028236	42,910091	
24	Niphargus angelieri	1219		1309432543	NA200		28S	28Slev3	28Sdes5	PONOVNO V				Grotte des Fees	FRA	3,028236	42,910091	
25	Niphargus aquilex	1214		1309432543	NA020		28S	28Slev2	28Sdes2	OK	CTCTGAAGAGAC EF617255			Wavreille, well 4	BEL	5,248482	50,121031	
26	Niphargus aquilex	1215		1309432543	NA020		28S	28Slev3	28Sdes5	OK	AAGGCTTCAGCA			Wavreille, well 4	BEL	5,248482	50,121031	
27	Niphargus aquilex	1216		1309432543	NA020		H3	H3aF2	H3aR2	OK	CGCTCGTAAGTC			Wavreille, well 4	BEL	5,248482	50,121031	
28	Niphargus aquilex	1217		1309432543	NA020		COI	LCO	HCO	OK	GGAGCTTGAGCT			Wavreille, well 4	BEL	5,248482	50,121031	
29	Niphargus aquilex	1213		1309432543	NA029		H3	H3aF2	H3aR2	OK	CGCTCGCAAGTC			Marden območje	GB	-0,855519	50,929969	
30	Niphargus aquilex	1212		1309432543	NA029		28S	28Slev2	28Sdes2	OK	CTCTGAAGAGAC			Marden območje	GB	-0,855519	50,929969	
31	Niphargus aquilex	1511		1309432543	NA194		28S	28Slev2	28Sdes2	NE DELA				Littlebourne	UK	1,168921	51,275484	
32	Niphargus aquilex	1512		1309432543	NA205		28S	28Slev2	28Sdes2	NE DELA				Littlebourne	UK	1,168921	51,275484	
33	Niphargus arbiter	672	14/08/2002	1309432543	NA052		28S	28Slev2	28Sdes2	OK	CTCTGAAGAGAC EF617287			Tounjčica Špilja	Croatia	15,32633388	45,24872352	
34	Niphargus arbiter	674		1309432543	NA052		H3	H3aF2	H3aR2	OK	TGCTCGCAAGTC			Tounjčica Špilja	Croatia	15,32633388	45,24872352	
35	Niphargus arbiter	676		1309432543	NA052		12S	?	?	OK	GCCTTTATATAT			Tounjčica Špilja	Croatia	15,32633388	45,24872352	

Web-database as a collaborative tool?

Morphology: deposition of figures – deposition of measurements

Quantitative data: MYSQL

Form - taxa and characters

[Niphargus home](#) [Submit form!](#) [Clear all!](#)

Select taxa (Check/Uncheck All)

- 1 Niphargus aquilex
- 2 Niphargus arbiter
- 3 Niphargus balcanicus
- 4 Niphargus bilecanus
- 5 Niphargus brachytelson
- 6 Niphargus camiolius
- 7 Niphargus costozzae
- 8 Niphargus crosticus
- 9 Niphargus dabarensis
- 10 Niphargus dalmatinus
- 11 Niphargus dimorphopus
- 12 Niphargus danconai
- 13 Niphargus dolichopus
- 14 Niphargus elegans

Select characters (You can select up to 100 characters.)

- 1 body length up to [mm]
- 2 rostrum
- 3 head length [of body length]
- 4 pereonites I-VI with up to [setae]
- 5 pereonite VII with
- 6 pereonite VII with [postero-ventral setae]
- 7 pleonites I-III with up to [setae]
- 8 pleonites I-III with
- 9 epimera II and III, posterior margins with [setae]
- 10 epimeral plate II postero-ventral corner
- 11 epimeral plate II, posterior margin
- 12 epimeral plate II, ventral margin
- 13 epimeral plate II, postero-ventral corner
- 14 epimeral plate II with [strong spine-like setae along ventral margin]

[Niphargus home](#) [Back](#)

	body length up to [mm]	rostrum
Niphargus arbiter	31	absent
Niphargus bilecanus	29	absent

Web-database as a collaborative tool?

Morphology: deposition of figures – deposition of measurements

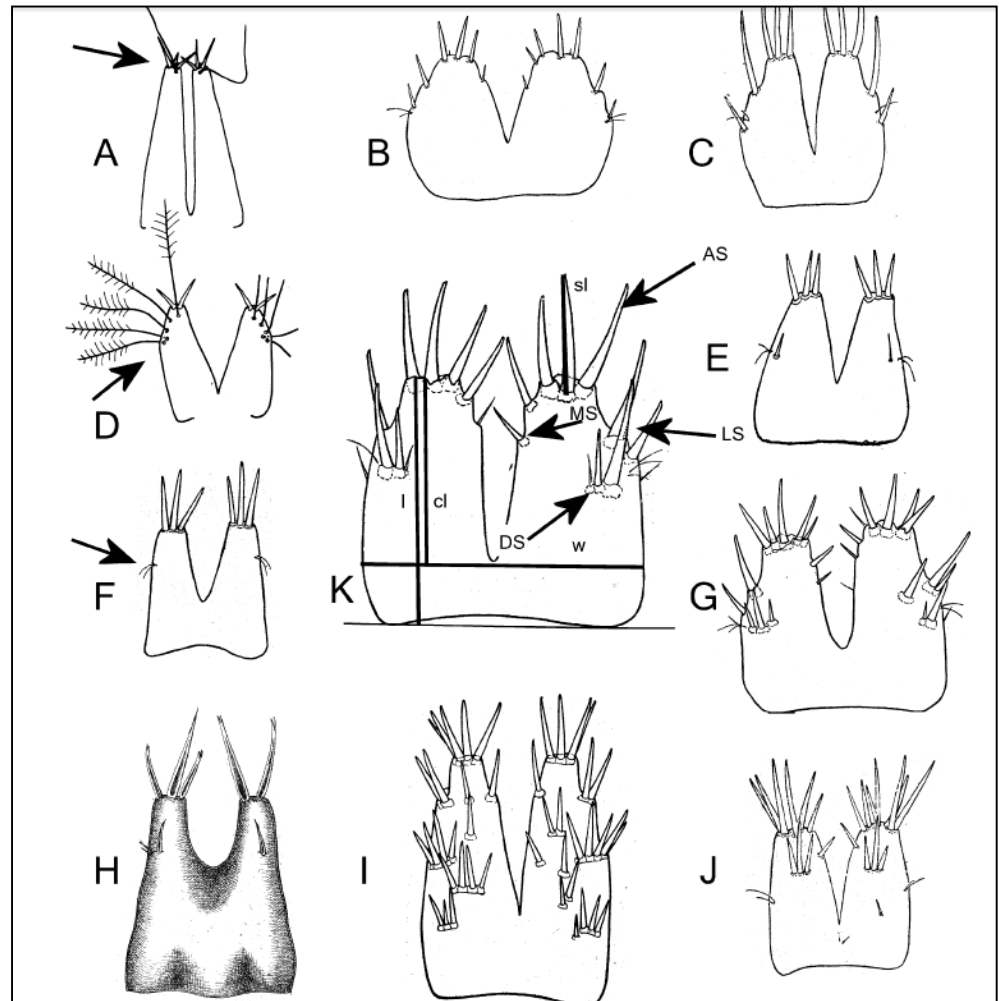
Quantitative data: MYSQL

SEVERAL IMPEDIMENT:

- taxon specific
- requires pre-defined protocols
- individual level
- unintentionally limits exploration

A CHALLENGE:

- making links with other databases (GeneBank, spatial data)



Taxon specific interactive database

Strengths:

- precise, a possibility to correct errors
- standardized protocols
- different properties linked with voucher specimens
- eases team work

Weakness:

- nature of characters may limit extent of the database (few taxa)

Global databases – character specific

Species names on a web: a chaos

The screenshot shows the Fauna Europaea website. The browser address bar displays 'http://www.faunaeur.org/'. The page features a navigation menu on the left with options like 'Name Search', 'Advanced Search', 'Taxon Tree', 'Distribution', 'Statistics', 'Experts', 'References', 'Taxonomic Resources', 'Other on-line databases', 'Acknowledgments', and 'LinkedIn'. The main content area has a 'Name search' section with dropdown menus for '(Sub) genus' and '(Sub) species', both set to 'is'. Below the search fields are 'Search' and 'Reset' buttons. A text box explains that the database contains scientific names of all European land and freshwater animals. At the bottom, there is a Creative Commons license logo (CC BY SA 4.0).

Zoobank : Niphargus 8 taxa

Fauna Europaea was supported by the European Commission under the Fifth Framework Programme and contributed to the Support for Research Infrastructures work programme with Thematic Priority Biodiversity.

Fauna Europaea is powered by MN

Niphargus – 295 taxa

The screenshot shows the WoRMS website. The browser address bar displays 'http://www.marinespecies.org/index.php'. The page features a navigation menu on the left with options like 'Home', 'About', 'Search taxa', 'Taxon tree', 'Literature', 'Distribution', 'Specimens', 'Match taxa', 'Editors', 'Statistics', 'Users', 'Webservice', 'Photogallery', 'Info downloads', 'Sponsors', and 'Activities'. The main content area has a 'Search WoRMS' section with dropdown menus for 'Common name' (set to 'contains') and 'Scientific name' (set to 'begins with'). Below the search fields are 'Search' and 'Advanced search' buttons. A text box explains that the database contains scientific names of all European land and freshwater animals. At the bottom, there is a Creative Commons license logo (CC BY SA 4.0).

Wikipedia (SLO): Niphargus - 206 taxa

EOL: Niphargus – 204-309 taxa

Niphargus – 261 taxa

Global databases – character specific

GeneBank: respository of sequences

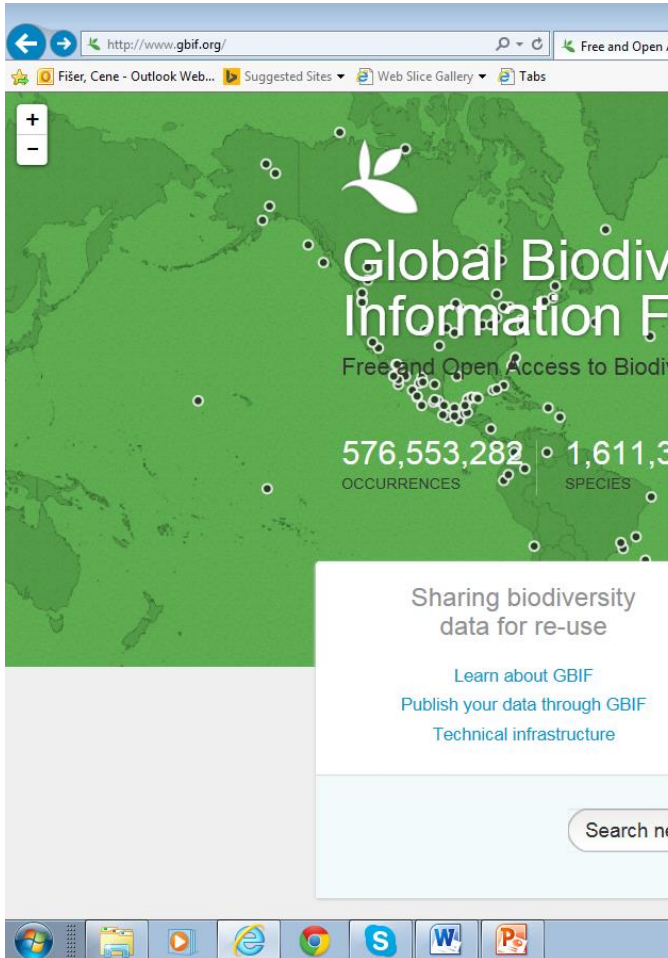
<http://www.ncbi.nlm.nih.gov/genbank/>

Main impediment: species id

The screenshot displays the GenBank website's overview page. At the top, there is a search bar with a dropdown menu set to 'Nucleotide' and a 'Search' button. Below the search bar is a navigation menu with various database categories: GenBank, Submit, Genomes, WGS, HTGs, EST/GSS, Metagenomes, TPA, TSA, and INSDC. The main content area is titled 'GenBank Overview' and includes sections for 'What is GenBank?', 'Access to GenBank', and 'GenBank Data Usage'. The 'What is GenBank?' section explains that GenBank is a NIH genetic sequence database, an annotated collection of all publicly available DNA sequences, and is part of the International Nucleotide Sequence Database Collaboration (INSDC). The 'Access to GenBank' section lists several ways to search and retrieve data, including using Entrez Nucleotide, BLAST, and the NCBI e-utils. The 'GenBank Data Usage' section states that the database is designed to provide and encourage access within the scientific community. The 'Confidentiality' section discusses the appearance of data in GenBank prior to publication. The 'Privacy' section mentions that human sequences should not include personal identity data. At the bottom of the page, there is a footer with navigation links, contact information for the National Center for Biotechnology Information, and logos for the Department of Health and Human Services and the USA.gov website. The footer also includes the text 'Last updated: 2015-05-07T14:49:35-04:00'.

Global databases – character specific

GBIF's vision: "A world in which biodiversity information is freely and universally available for science, society and a sustainable future.,,"
<http://www.gbif.org/>



The screenshot shows the GBIF website interface. At the top, there's a navigation bar with the GBIF logo and the text "Global Biodiversity Information Framework". Below this, a large green map of the world is displayed with numerous white circular data points. To the right of the map, the text "Global Biodiversity Information Framework" is visible, along with the tagline "Free and Open Access to Biodiversity Information". Below the map, two statistics are shown: "576,553,282 OCCURRENCES" and "1,611,300 SPECIES". A search bar is located at the bottom right of the page. The Windows taskbar is visible at the bottom of the screenshot, showing various application icons and the system clock displaying "15:53 30.9.2015".

- It provides a single point of access (through this portal and its web services) to more than **570,000,000 records**, shared freely by hundreds of institutions worldwide, making it the biggest biodiversity database on the Internet.
- The data accessible through GBIF relate to evidence about more than **1.6 million species**, collected over three centuries of natural history exploration and including current observations from citizen scientists, researchers and automated monitoring programmes.
- More than **1,400 peer-reviewed research publications have cited GBIF** as a source of data, in studies spanning the impacts of climate change, the spread of pests and diseases, priority areas for conservation and food security. **About one such paper is published each day.**
- Many GBIF Participant **countries have set up national portals using tools, codes and data freely available through GBIF** to better inform their citizens and policy makers about their own biodiversity.

Global databases – character specific

GBIF Global Biodiversity Information Facility

3.427 Occurrences | 255 Species

View occurrences

Niphargus Schiödte, 1849

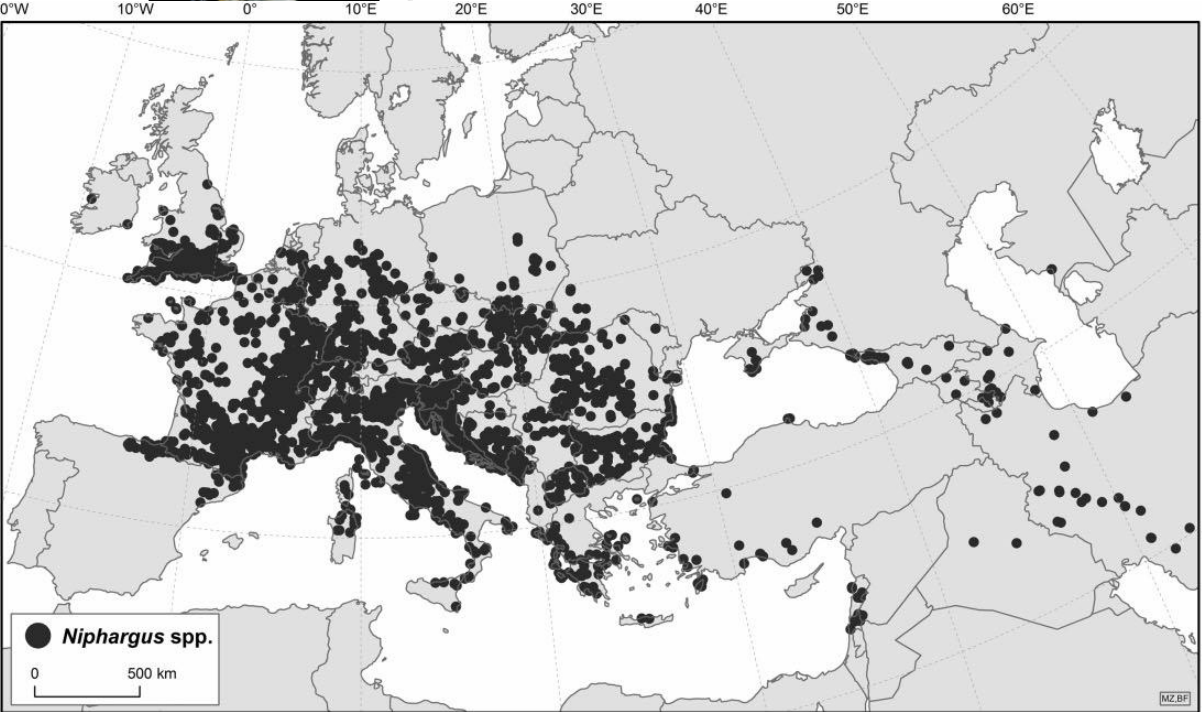

Genus in GBIF Backbone Taxonomy

Animalia · Arthropoda · Malacostraca · Amphipoda · Niphargidae

Information

Overview

FULL NAME Niphargus Schiödte, 1849	TAXONOMIC STATUS Accepted genus
COMMON NAMES · Höhlenflohkrebse deu more >	ACCORDING TO World Register of Marine S
	PUBLISHED IN Danske Selske. Skr. . (5) 2.
	HABITAT Not terrestrial, Not marine



Pre 1900 1900s 1910s 1920s 1930s 1940s 1950s 1960s 1970s 1980s

Main impediments:

- species id
- completeness of data for selected taxon
- quality of coordinates

Global databases – character specific

Strengths:

- different taxa in one place, a promising source of information for ecology and evolutionary biology

Weakness:

- difficult or impossible to revise errors
- poor connection between databases dealing with different types of characters

Global issues

1. Quality of database-quality of data.
2. Linking the taxon oriented databases with character oriented databases: increase of accuracy and number of taxa.
3. Promotion database publishing:
 - Citation problem
 - Rewarding in funding agencies
 - Role of journals
4. Collaborative network – stimulating collaborators

Thanks...

Maja Zagmajster

Martin Turjak

Roman Luštrik

The rest of SubBio lab team


Taxon specialized websites: not necessarily databases

What is the name of this creature?

Salticid spiders

<http://salticidae.org/salticid/main.htm>

**Monograph of
*Salticidae (Araneae)***
By Jerzy Prószyński
2007
Version revised in part on February 2008

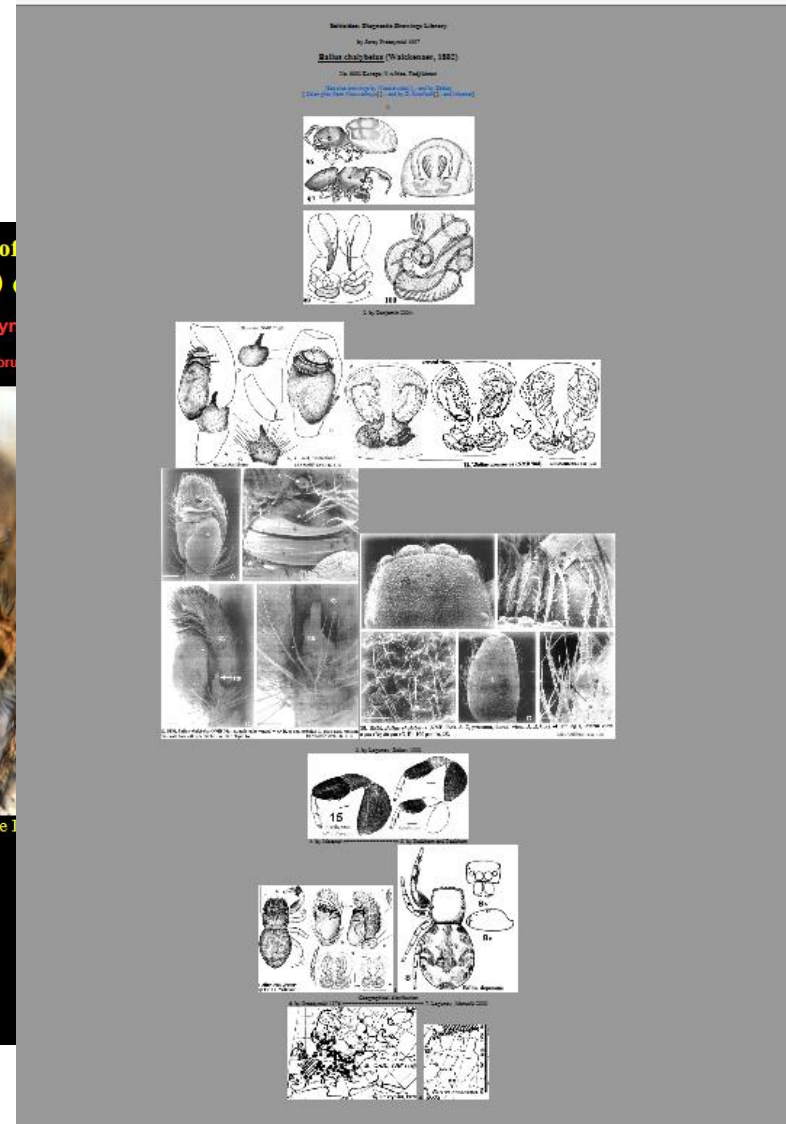


Click the picture to see list of genera of the 1

PART I. DATABASE
PART II. DIAGNOSTIC DRAWINGS
PART III. CATALOGUE OF THE *Salticidae* OF THE WORLD


Special features: [Pictorial Indexes, Keys and Taxonomic Revisions](#)
See also: [OTHER WWW SALTICIDAE PAGES](#)

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Global databases – character specific

Morphology: deposition of figures – deposition of measurements

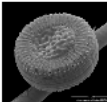

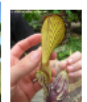
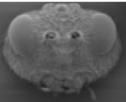

 **Welcome to Morphbank on Sky**
User: Guest [\[click to login\]](#)

[About](#) [Browse](#) [Tools](#) [Help](#)

Featured from 381962 images


MorphBank
<http://www.morphbank.net/>

click images below to [browse new uploads and collections](#)


				
Anglerfish	Chromista	<i>M. blackburni</i>	Aristolochia	Cynipidae

News and Updates


Morphbank move to new servers
The Morphbank system at FSU has moved to the Sky server farm at the FSU Research Computing System and the Nolestor storage system. These new systems will keep Morphbank running into the future. ...

WHO SAYS THE SKY IS THE LIMIT?

(Posted: 07-27-15)


Filtered Push plugin in process
The Filtered Push group have created a Morphbank plugin that extracts annotations from the Morphbank database and pushes them into the FP annotation services. ...


(Posted: 11-07-14)

Morphbank in DINA Project
The DINA project develops an open-source Web-based information management system for natural history data. At the core of the system is support for assembling, managing and sharing data associated with natural history collections and their curation ("collection management"). Target collections include zoological, botanical, geological and paleontological collections, living collections, biodiversity inventories, ...


(Posted: 11-07-14)

Troy University Herbarium Online
The Troy University Herbarium images from the Deep South Plant Specimen Imaging Project are now available. Search for "Michael Woods" to see the images and specimens ...


(Posted: 11-06-14)

[\(see all past news\)](#)

Global databases – character specific

Morphology: deposition of figures – deposition of measurements

Image 833234
User: Guest (click to login) | About | Browse | Tools | Help

Image Record: [833234] *Alloxysta victrix*

Contributor: Jordi Parceta-Martinez
Submitter: Jordi Parceta-Martinez
Group: Universitat de Barcelona
Date Submitted: 2013-10-09
Last Modified: 2013-10-09
Publish Date: 2011-07-01
Description: New image from upload

Magnification: 3000
Dimension (px): 750x576
Resolution (PPI):
Submitted as: tiff
Original File Name: ALLOX19.TIF
Photographer: Jordi Parceta-Martinez

View id: 833278
Specimen part: Metatarsal claw
Angle: Lateral
Technique: SEM
Preparation: Air dried, gold coated

Download: original (tiff) (432.11 KB)
full sized jpeg (71.72 KB) medium sized jpeg (22.9 KB)

Copyright: public domain
License: This work is free of known copyright restrictions.

Specimen
Specimen id: 833070
Basis of record: [S] - Specimen
Sex: Female
Form: Indeterminate
Stage: Adult
Catalog number:
Collector:
Date collected:

Locality [edit this locality](#)
Locality id: 101950
Continent: UNKNOWN
Water Body:
Country: UNKNOWN
State/Province:
County:
Locality: Unknown
Latitude:
Longitude:
Elevation (m):
Depth (m):

Determination
Kingdom: Animalia
Phylum: Arthropoda
Class: Insecta
Order: Hymenoptera
Family: Pimplidae
Genus: *Alloxysta*
Species: *Alloxysta victrix*

External links/identifiers
External Unique Reference:
Specimen 833070: gdforma:identifier: urn:uuid:23148099-6e03-455b-9e5a-257af6c75862
gdforma:identifier: urn:uuid:284808e8-b149-434fae32-8b6d872497cc

Determination annotations [Add Annotation...](#) | **Other Annotations** [Add Annotation...](#)

Related Annotations

Taxonomic Name	Taxon Author	Prefix	Suffix		
<i>Alloxysta victrix</i>		none	none	1	0

Image 196798
User: Guest (click to login) | About | Browse | Tools | Help

Image Record: [196798] *Dulichella spinosa*

Contributor: Dean Peake
Submitter: Dean Peake
Group: SCANIT
Date Submitted: 2007-12-15
Last Modified: 2011-09-01
Publish Date: 2007-12-15
Description: Image added to Database

Magnification: 25
Dimension (px): 1094x1176
Resolution (PPI):
Submitted as: jpg
Original File Name: Dulichella spinosa.jpg
Photographer:

View id: 196772
Specimen part: Whole body
Angle: Lateral left
Technique: Digital camera
Preparation: Wet Mount

Download: original (jpeg) (404.49 KB)
full sized jpeg (404.49 KB) medium sized jpeg (80.59 KB)

Copyright: dean@peake.com
License:

Specimen
Specimen id: 196798
Basis of record: [S] - Specimen
Sex: Male
Form: Not Applicable
Stage: Adult
Catalog number:
Collector: Grango County Sanitation District
Date collected: 2006-07-11

Locality [edit this locality](#)
Locality id: 196794
Continent: NORTH AMERICA
Water Body:
Country: UNITED STATES
State/Province:
County:
Locality: Southern California Bight
Latitude:
Longitude:
Elevation (m):
Depth (m):

Determination
Kingdom: Animalia
Phylum: Arthropoda
Class: Malacostraca
Order: Amphipoda
Family: Dulichellidae
Genus: *Dulichella*
Species: *Dulichella spinosa*

External links/identifiers
External Unique Reference:
Specimen 196798: gdforma:identifier: urn:uuid:832673ef-dea7-42ef-8326-926c79290a62
gdforma:identifier: urn:uuid:f7260c49-9185-4f6b-92d0-7040ed4100c

Determination annotations [Add Annotation...](#) | **Other Annotations** [Add Annotation...](#)

Related Annotations

Taxonomic Name	Taxon Author	Prefix	Suffix		
<i>Dulichella spinosa</i>	Stout, 1912	none	none	1	0

Beginnings of the databases...

Special software packages: DELTA, Lucid

Description Language for
Taxonomy (DELTA)

<http://delta-intkey.com/>

A TOOL FOR:

- descriptions
- identification keys (different types)
- files for phylogenetic/phenetic analysis
- comparison of taxa

MAIN IMPEDIMENT:

limited analytic frame, species – population level

The screenshot displays the INTKEY software interface for 'Beetle Larvae of the World'. It features a 'Differences' window comparing two taxa: Dorcatominae ANOBIIDAE and Ceryloninae (major part) CERYLONIDAE. The comparison is organized into several categories: Length, Body, Body (as viewed from above), Vestiture, and Dorsal surfaces. Each category lists characteristics for both taxa, with differences highlighted in bold. For example, under 'Body', Dorcatominae is 'strongly curved ventrally (c-shaped)' while Ceryloninae is 'relatively straight or only slightly curved ventrally'. Under 'Dorsal surfaces', Dorcatominae is 'very lightly pigmented or sclerotized' while Ceryloninae is 'more or less heavily pigmented or sclerotized'. A 'Remaining Taxa' list on the right shows 390 taxa, with a yellow callout 'Differences between taxa' pointing to the current comparison. Below the text, two illustrations are shown: a lateral view of a Dorcatoma sp. larva and a dorsal view of a Mychocerus sp. larva. The interface includes a menu bar (File, Window, Help), a toolbar, and a status bar at the bottom indicating '7 differences'.

Beginnings of the databases...

Special software packages: DELTA, Lucid

Description Language for
Taxonomy (DELTA)

<http://delta-intkey.com/>

SOURCE OF INFORMATION THAT
CAN BE INCLUDED:

- discrete and continuous morphological characters
- illustrations and photos
- sonograms
- videoclips

INTKEY : Beetle Larvae of the World

File Window Help

Available Characters [9]

- length
- body (whether disc-like)
- body (whether C-shaped)
- body (cross-section)
- body, as viewed from above (shape)

Remaining Taxa [390]

- ADERIDAE
- AGYRTIDAE
- ALEXIIDAE
- AMPHIZOIDAE
- ANOBIIDAE (major part)
- Dorcatominae ANOBIIDAE**
- Cryptorama ANOBIIDAE
- Ptininae ANOBIIDAE
- Anthicinae ANTHICIDAE
- Cotes ANTHICIDAE

Differences between taxa

Dorcatominae ANOBIIDAE: Dorcatoma

Control Window

Notes

Dorcatoma sp.

Ceryloninae (major part) CERYLONIDAE: Mychocerus

Control Window

Notes

Mychocerus sp.

Differences

Length

- Dorcatominae ANOBIIDAE
less than 3 mm; or 3 to 15 mm
- Ceryloninae (major part) CERYLONIDAE
less than 3 mm

Body

- Dorcatominae ANOBIIDAE
strongly curved ventrally (c-shaped)
- Ceryloninae (major part) CERYLONIDAE
relatively straight or only slightly curved ventrally

Body, as viewed from above

- Dorcatominae ANOBIIDAE
circular in cross-section, or slightly flattened
- Ceryloninae (major part) CERYLONIDAE
slightly flattened

Body, as viewed from above

- Dorcatominae ANOBIIDAE
oblong to ovate, not parallel-sided
- Ceryloninae (major part) CERYLONIDAE
elongate and more or less parallel-sided; or oblong to ovate,
not parallel-sided

Vestiture

- Dorcatominae ANOBIIDAE
consisting of fine hairs or setae only
- Ceryloninae (major part) CERYLONIDAE
including bristles, scales, expanded or complex hairs

Dorsal surfaces

- Dorcatominae ANOBIIDAE
very lightly pigmented or sclerotized
- Ceryloninae (major part) CERYLONIDAE
more or less heavily pigmented or sclerotized

Dorsal surfaces

- Dorcatominae ANOBIIDAE
generally smooth
- Ceryloninae (major part) CERYLONIDAE
generally granulate or tuberculate

7 differences.



CINKARNA

The importance of data interpretation and dissemination for the future of nanotechnology

Mag. Vladimír Vrečko

Future of nanotechnology





**Hope you enjoyed and
thank you for your attention!**

Public point of view

- Yet another potentially dangerous technology.
- Industry pursues its own interests and can not be trusted.
- Research which is funded by private funds can not be trusted.
- Even scientists don't agree with each other, so there is obviously something wrong.
- Public media informs us of the research results that prove nanotechnology is dangerous.
- We don't want the technology until we are sure it is safe.

Scientists point of view – Material scientists

- We have discovered numerous new materials and developed their applications.
- Everyone would benefit from their use.
- We don't see, why industry does not recognise the potential and finance us extensively.
- If we could just get money, we would have immediate success on market (everyone would buy our products).
- The potential risks, costs, availability of the materials and public opinion are not our concern.
- Yet another proof that we are not understood and appropriately valued.

Scientists point of view – Risk Assessment groups

- We got financed to reveal the safety risks.
- Public expects us to prove we are independent.
- To prove that nano materials are dangerous is therefore a success and worth publishing.
- Such findings are great for PR (ensures headlines and recognition of the research group).
- If the material or technology does not manifest risky properties, it is a disappointment and not worth mentioning.
- Only if we reveal new safety risks will we have a chance for future funding of our research.

Industry point of view

- We are interested in new technologies, but we take decisions based on economy.
- We can manage the costs, however market is still underdeveloped because the risks and legal frameworks are not yet sufficiently defined.
- Public opinion is very sceptical and media are not in favour, publishing frightening stories.
- Customers and investors don't want to take premature financial risks.
- Can it happen to nanotechnology that it will follow the footsteps of GMOs?

How about that:

Scientific research title:

Titanium dioxide nanoparticles induce DNA damage and genetic instability in vivo in mice.

Title in newspaper:

Scientists say nanoparticles cause DNA damage.

Implications for the customers:

Be aware of the products that contain nanoparticles.

Demand that all products containing nanoparticles shall be labeled (as dangerous?).

Don't buy such products as they can damage your DNA (scientist have said so).

What is needed?

- Industry needs stable and known market conditions and legal boundaries.
- It is willing to take reasonable precautionary measures, but doesn't want unpredictable environment for investments.
- Scientists should be aware of their responsibility at shaping public opinion.
- They should publish also the results in which they confirm that some nanomaterials do not pose safety risks.
- Market will grow only if all the stakeholders will feel secure.

What is needed?

- Scientist shall pursue balanced and objective approach and help regulatory bodies at designing legal frameworks.
- Scientists shall not forget that they are after all financed from the money which is mainly created by the industry.
- Scientists and industry shall join forces and through public media present nanotechnology as mature technology, which has many advantages. And, normally, some potential risks, which we can however control through joint activities of researchers, legal bodies and manufacturers.

What is needed?

We shall persuade all stakeholders that they have nothing to be afraid of.

How shall we act?



So?

Or so?



A large, dark tree stands on the left side of a lush green field. The sun is setting in the background, creating a warm, golden glow over the rolling hills. The sky is a mix of orange and yellow, and the overall scene is peaceful and hopeful.

**Hope to see you
in bright future!**



**NACIONALNI LABORATORIJ ZA
ZDRAVJE, OKOLJE IN HRANO**

Prvomajska ulica 1, 2000 Maribor

Challenges of Regulation and Risk Assessment of Nanomaterials Part 2: ECHA - NMWG

Dr. Mojca Kos Durjava

COMPInNANO, Ljubljana, 2.-3.10.2015

Risk Assessment of Nanomaterials

EC and OECD - Risk assessment of nanomaterials can be managed through existing regulatory frameworks, adapted to take into account the specific properties of manufactured nanomaterials.

REACH, CLP - EC is modifying some of the technical provisions in the REACH Annexes - amendments in 2016

Biocides – Biocidal Products Regulation (BPR), 2013: contains a definition of nanomaterial.

Risk Assessment of Nanomaterials

EC launched a comprehensive REACH Implementation Project on Nanomaterials (RIPoN) in 2009:

- RIPON1 : Substance Identity
- RIPON2 : Information Requirements
- RIPON3 : Chemical Safety Assessment

CARACAL is an expert group which advises the European Commission and ECHA on questions related to REACH and CLP.

CASG nano - Competent Authority Subgroup on Nanomaterials

- Nanomaterials in REACH
- Classification, Labelling and Packaging of Nanomaterials in REACH and CLP.

OECD - WPMN

Working Party on Manufactured Nanomaterials - WPMN

- is linked to the implementation of REACH - TG and guidance.

OECD and WPMN SG on Risk Assessment and Regulatory Mitigation SG-AP

- SG on Exposure and Exposure Mitigation Exp. Mitigation
- SG on the Environmentally Sustainable use of Nanomaterials LCA
- **SG on Testing Assessment of Manufactured Nanomaterials SG-TA:**
 - TG update
 - Assessment of data
 - In vitro work
 - Alternative methods (e.g. read-across)

ECHA - NMWG

NMWG – Nanomaterials Working Group

The aim – to discuss topical scientific and technical issues relevant to the implementation of REACH, CLP and Biocidal Product Regulation in relation to nanomaterials.

From 2013, 7th meeting in November 2015, chair Frank Le Curieux from ECHA

50 or more participants at meetings, twice a year:

- MSCA – Member States Competent Authorities
- European Commission (DG ENT and DG ENV)
- DG Joint Research Centre
- ECHA-NMWG Accredited Stakeholders Observers (ASO)
- ECHA representatives
- Invited speakers from industry, science

ECHA - NMWG

NMWG – Nanomaterials Working Group

- ECHA presentations on their work on NM
- Industry presentations of dossiers of registered NM
(10 NM – cerium oxide, calcium carbonate, zinc oxide, multi-wall nanotubes, titanium dioxide,...)
- Scientists presentations on their research in the field of NM (NanoREG, NanoValid, Marina,...)
- Other presentations

Discussion, working in groups.

ECHA - NMWG

ECHA guidance on RA of NM

Already developed by ECHA: a technical manual on how to include information on nanomaterial in a IUCLID dossier which is an integral part of every REACH registration.

REACH amendments of Annexes (VII-X) for nanomaterials



ECHA guidance on RA of NM before 2018 registration deadline

- ECHA guidance (updating)
- Practical guides/examples

Advice and expertise from the NMWG.

ECHA - NMWG

IUCLID 6

IUCLID is a software application to capture, store, maintain and exchange data on intrinsic and hazard properties of chemical substances. Report generator for Chemical Safety Report - CSR.

„Assessment entity“ concept – a feature for IUCLID 6

- Enable transparent reporting of hazard, use and exposure information for NM.

Multiple composition or multiple forms of the same substance.

Many constituents within the substance.

Variable composition of the substance.

Forming of transformation products on use.

ECHA - NMWG

Read across and grouping of NM

Colaboration of ECHA, JRC, NanoREG and RIVM.

Many nanoforms on the market:

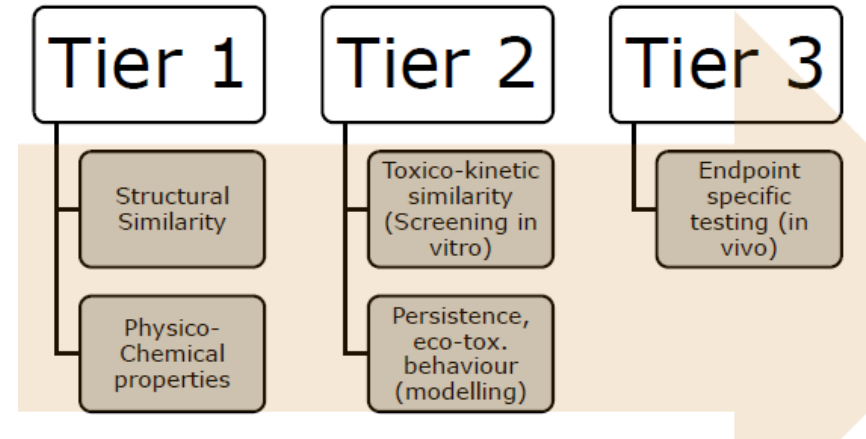
- Are studies from one form applicable to other nanoforms?
- How and when can data be used on nanoforms or beetwen non-nano and nanoforms?

Case studies presented;

Develop a decision framework;

Link to the OECD discussion;

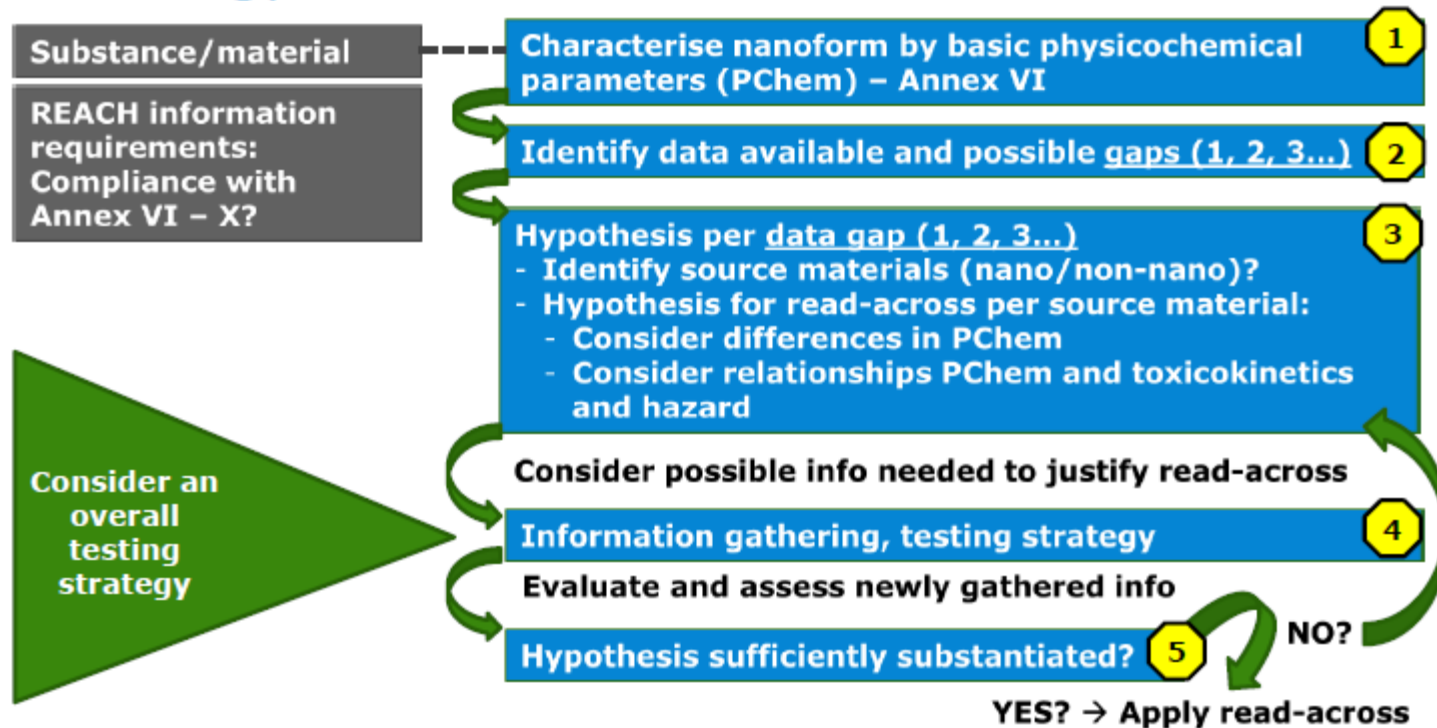
Template for reporting on read – across.



ECHA - NMWG

Read across and grouping of NM

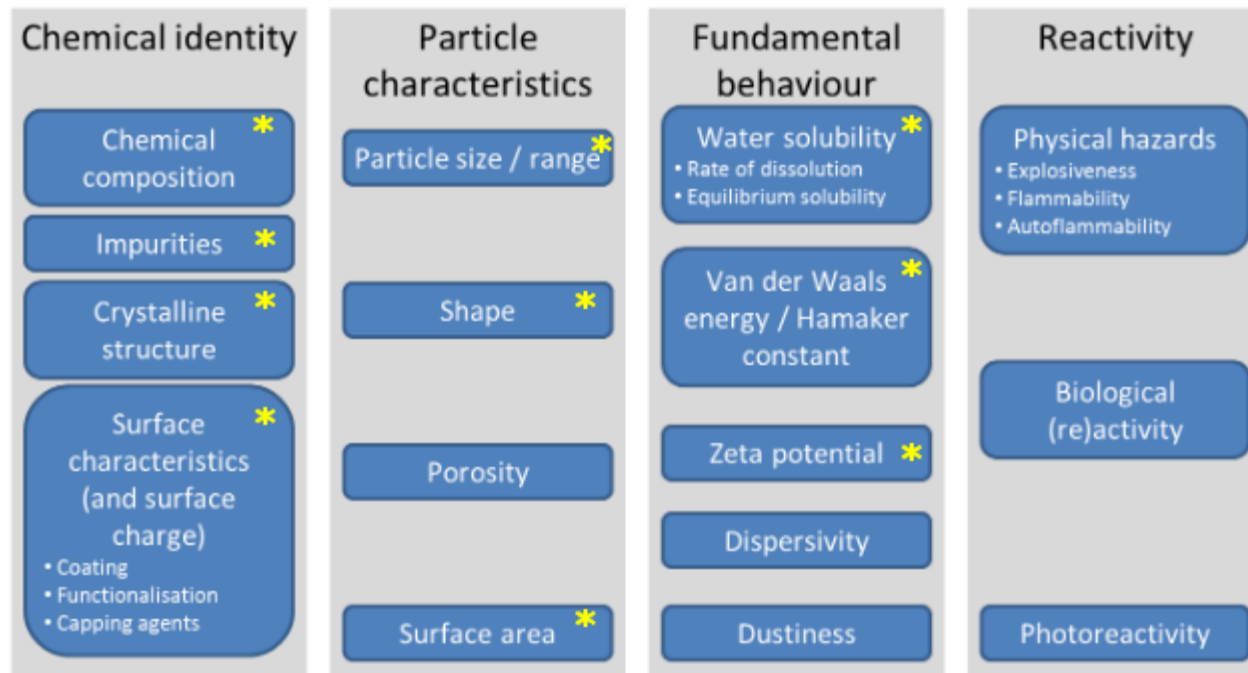
Strategy for read-across of nanomaterials



ECHA - NMWG

Read across and grouping of NM

Parameters influencing nanomaterial behaviour:



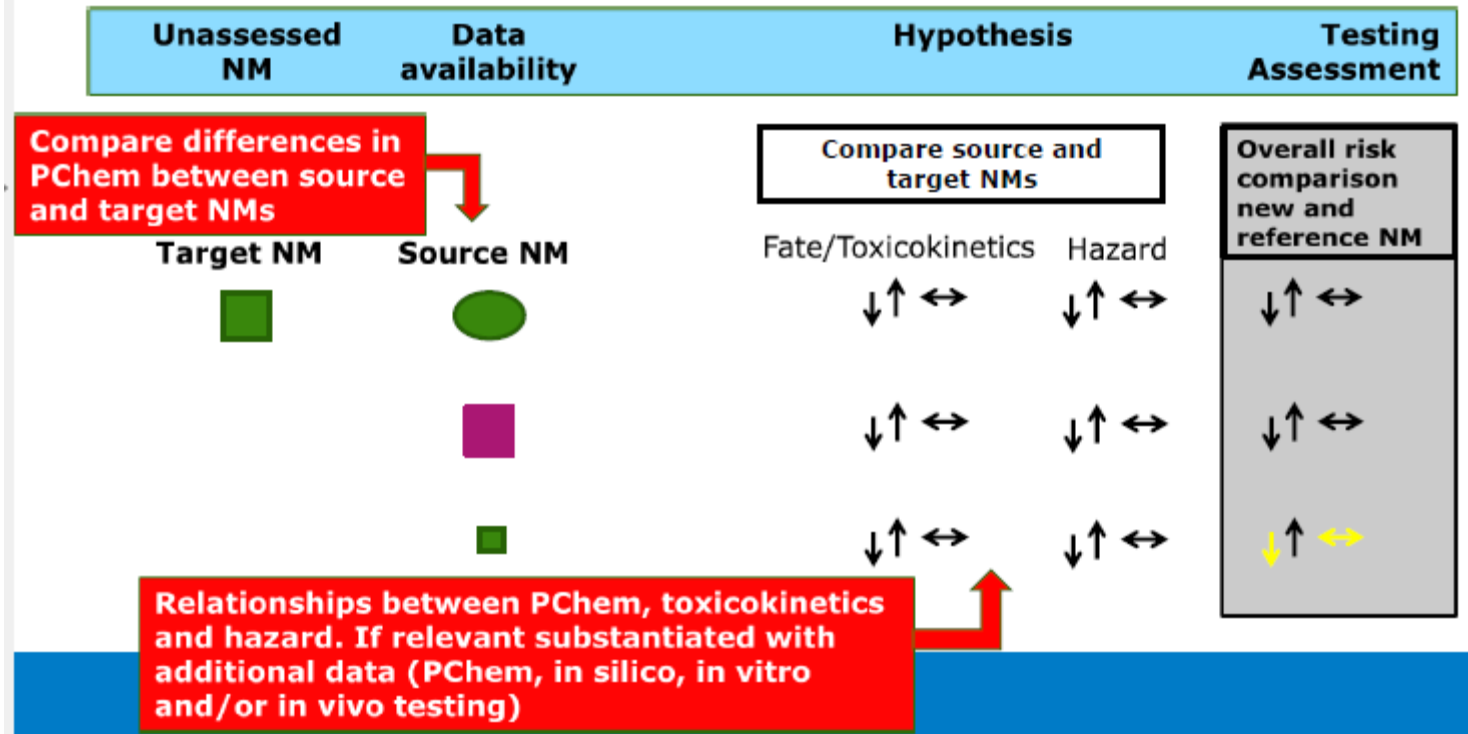
* Essential information to characterise nanomaterials

ECHA - NMWG

Read across and grouping of NM

Hypothesis on read across per data point

- Find source materials with information for same exposure route
- Assess:



ECHA - NMWG

Read across and grouping of NM

- The focus of the framework is on Pchem data as well as their relation to the different steps in the life cycle and biological pathway (exposure, kinetics, hazard).
- Identify Pchem that matter and provide strategy for systematic assessment.
- Needs to be validated in practice.
- Future implementation of growing experience and understanding on behaviour, fate, toxicokinetics and toxicity.

ECHA guidance will be developed on read across.

ECHA - NMWG

NMWG group

- until guidances for REACH and CLP are developed.

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Challenges of Regulation and Risk Assessment of Nanomaterials

part.1

Karmen Krajnc, M.Sc.
Chemicals Office of the Republic of Slovenia

EU – strategic documents

- **Towards a European Strategy for Nanotechnology 2004**
 - **European Nanotechnology Action Plan 2005-2009**
 - Implementation Reports 2007, 2009
- **Code of conduct for nanotechnology research 2008**
- Communication from the Commission to the European Parliament, THE Council and the European Economic and Social Committee
2008 (first regulatory review)
2012 (second regulatory review)

http://ec.europa.eu/research/industrial_technologies/pdf/policy/communication-from-the-commission-second-regulatory-review-on-nanomaterials_en.pdf

Definition

(Commission Recommendation 2011/696)

- 'Nanomaterial' means a natural, incidental or manufactured material containing particles, in an unbound state or as an aggregate or as an agglomerate and where, for **50 %** or more of the particles in the number size distribution, one or more external dimensions is in the size range **1 nm-100 nm**.
- In specific cases and where warranted by concerns for the environment, health, safety or competitiveness the number size distribution threshold of 50 % may be replaced by a threshold between 1 and 50 %.
- By derogation, fullerenes, graphene flakes and single wall carbon nanotubes with one or more external dimensions below 1 nm should be considered as nanomaterials.

<http://eur-lex.europa.eu/LexUriServ/LexUriServ.do?uri=OJ:L:2011:275:0038:0040:en:PDF>

- Consideration of possible changes to this definition

<https://ec.europa.eu/jrc/en/publication/eur-scientific-and-technical-research-reports/towards-review-ec-recommendation-definition-term-nanomaterial-part-3-scientific-technical>

Cosmetic regulation(1223/2009)

- definition: 'nanomaterial' means an insoluble or biopersistent and intentionally manufactured material with one or more external dimensions, or an internal structure, on the scale from 1 to 100 nm.
- colorants, preservatives and UV-filters must be authorized (positive lists: Annexes IV, V and VI)
 - titanium dioxide (nano), zinc oxide (nano), tris-biphenyl triazine (nano) and carbon black (nano) shall be added shortly
- notifications of **safety information** for nano substances(art.16)
- **labeling: „nano“**

Biocidal regulation (528/2012)

- specific approval of nano form of biocidal substance (art. 4(4))
- labeling obligation for treated articles, art. 58(3):
the name followed by the word 'nano' in brackets
- nano-definition

REACH-nanomaterials registered

- Carbon black
- Cerium dioxide
- Calcium carbonate
- Zinc oxide
- Silver
- MWNT (multi-wall nanotubes)
- MWNT as a form of graphite
- Titanium dioxide
- Silicate(2-), hexafluoro-, disodium, reaction products with lithium magnesium sodium silicate

Substance evaluation

- **Substances selected for CoRAP** (Community rolling action plan) based on initial grounds of concern: evaluated by member states, coordinated by ECHA
- **Silicon dioxide** (synthetic amorphous silica - SAS) – the Netherlands, 2012
<http://echa.europa.eu/documents/10162/a94c8df7-81c5-4946-80ae-dfa9275897e1>
- **Silver** – the Netherlands, 2012 (ongoing)
- **Titanium dioxide** – France, 2015 (not started yet)

OECD Working Party on Nanomaterials

- TESTING PROGRAMME ON MANUFACTURED NANOMATERIAL

<http://www.oecd.org/chemicalsafety/nanosafety/testing-programme-manufactured-nanomaterials.htm>

- Cerium oxide
- Multi-walled carbon nanotubes (MWCNTs)
- Single-walled carbon nanotubes (SWCNTs)
- Dendrimers
- Nanoclays
- Fullerenes (C60)
- Silicon dioxide
- Gold nanoparticles
- Silver nanoparticles
- Titanium dioxide, Zinc oxide (not publicly available yet)

REACH NANO legislative challenges

- Definition of „nanomaterial“
- REACH Annexes (registration purposes)
- Discussion on possible EU NANO DATABASE

Chemicals Office of the RS-nano activities

- Nanoportal
- Identified 30 companies (producers, users of nanosubstances)
 - Ag, Zn, Al, TiO₂, soot, ZnO, SiO₂, Fe.H₂SO₄, Si, Al₂O₃, graphite-C, SiC,Cu, CdS, FeCl₃, B, Fe, Cr, CuO, MoSi, MoS₂, Wox
 - Coatings/Surface modification,
 - Pigments,
 - Polymers/Composits,
 - Cosmetic,
 - Medical devices & Medicinal products,
 - Textiles,
 - Electronics.
- Awareness raising

Nanoportal

<http://www.uk.gov.si/>

www.uk.gov.si

AKTUALNO

Nanoportal



Z namenom okrepitve povezav med slovensko industrijo in številnimi priznanimi slovenskimi znanstveniki, ki se pri nas ukvarjajo s posameznimi vidiki nanomaterialov in nanovarnosti, smo v sodelovanju z slovenskimi raziskovalci, pripravili NANOPORTAL.

Portal je sestavljen iz dveh delov:

- [Splošna predstavitev področja, skupaj s koristnimi povezavami.](#)
- [Seznam več kot 100 slovenskih raziskovalcev, vključno z navedbo področja njihovega delovanja.](#)

Načrtujemo, da se bo portal sproti dopolnjeval, vse morebitne predloge za spremembe lahko posredujete na URSK, mag. Karmen Krajnc (tel: 478 6054, mail: [karmen.krajnc\(at\)qov.si](mailto:karmen.krajnc(at)qov.si))

DRUGE VSEBINE

[Registracija, evalvacija,](#)

[Razvrščanje, pakiranje in](#)

[Biocidni proizvodi](#)

Elektronsko sporočanje podatkov o kemikalijah



Zavezancem za sporočanje podatkov o kemikalijah na podlagi 35. člena Zakona o kemikalijah in Pravidnika o sporočanju podatkov o kemikalijah je na voljo spletna aplikacija ISK za elektronsko sporočanje podatkov. Aplikacija ISK omogoča pregled, vnos, spreminjanje in ukinjanje sporočil o kemikalijah ter pregled in vnos letnih količin o kemikalijah. Več o elektronskem sporočanju podatkov, pogojih za uporabo aplikacije, načinu registracije uporabnikov in uporabi aplikacije najdete na zavihku [e-sporočanje](#).

NOVICE

22. 9. 2015
[Objava prostega delovnega mesta projektneга svetovalca Twinning projekta v Beogradu](#)

3. 9. 2015
[Javni posvet glede identifikacije snovi, ki vzbujajo veliko zaskrbljenost \(SVHC\)](#)

2. 9. 2015
[Registracija za uporabnike anhidrida očetne kisline](#)

[Več ▶](#)

DOGODKI

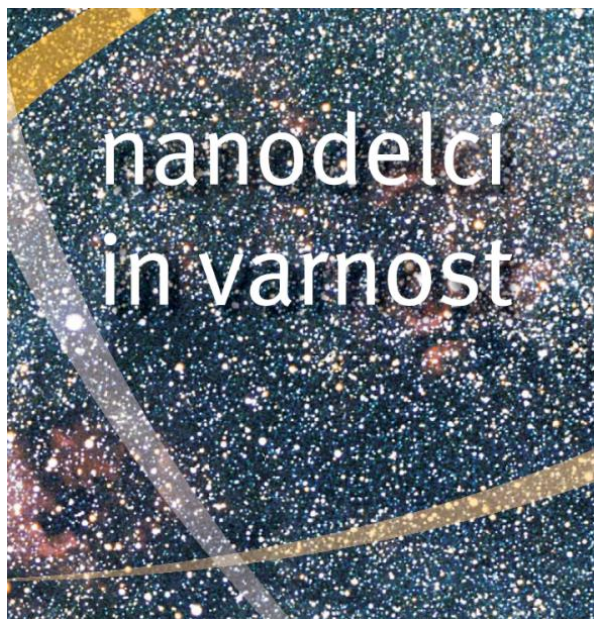
27. 5. 2015
[Predstavitve s seminarja "Strokovno srečanje svetovalcev za kemikalije", 20.05.2015](#)

13. 4. 2015
[Strokovno srečanje svetovalcev za kemikalije, 20.05.2015](#)

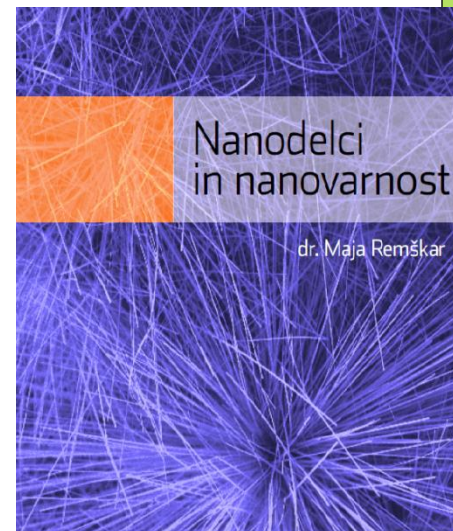
2. 4. 2015
[Biocidni simpozij 2015](#)

A	B	C	D
Raziskovalna organizacija	Kontaktna oseba	E-mail	Nanomateriali
Cinkarna Celje	Dejan Verhovšek	dejan.verhovsek(at)cinkarna.si	TiO2
Cinkarna Celje	Nika Veronovski	nika.veronovski(at)cinkarna.si	TiO2
Helios	Znoj Bogdan	bogdan.znoj(at)helios.si	nanoclay
Industrijski razvojni center slovenske predilne industrije	Žlabravec Verica	vera.zlabravec(at)litija.com	nanodelci na tekstilu
Institut "Jožef Stefan"	Bernik Slavko	slavko.bernik(at)ijs.si	ZnO
Institut "Jožef Stefan"	Gyergyek Sašo	saso.gyergyek(at)ijs.si	magnetni nanodelci
Institut "Jožef Stefan"	Jesih Adolf	adolf.jesih(at)ijs.si	Molx
Institut "Jožef Stefan"	Kobe Spomenka	spomenka.kobe(at)ijs.si	SiC
Institut "Jožef Stefan"	Kralj Slavko	slavko.kralj(at)ijs.si	magnetni nanodelci
Institut "Jožef Stefan"	Kutnjak Zdravko	zdravko.kutnjak(at)ijs.si	MoS2
Institut "Jožef Stefan"	Lisjak Darja	darja.lisjak(at)ijs.si	feriti
Institut "Jožef Stefan"	Makovec Darko	darko.makovec(at)ijs.si	magnetni nanodelci
Institut "Jožef Stefan"	Milošev Ingrid	ingrid.milosev(at)ijs.si	nanodelci kovinskih implantov
Institut "Jožef Stefan"	Mozetič Miran	miran.mozetic(at)ijs.si	FeOx, NbOx
Institut "Jožef Stefan"	Mrzel Aleš	ales.mrzel(at)ijs.si	MoSi
Institut "Jožef Stefan"	Muševič Igor	igor.musevic(at)ijs.si	SiO2
Institut "Jožef Stefan"	Novak Saša	sasa.novak(at)ijs.si	SiC
Institut "Jožef Stefan"	Panjan Peter	peter.panjan(at)ijs.si	TiAlN/CrN
Institut "Jožef Stefan"	Pribošič Irena	irena.pribosic(at)ijs.si	KNbO3
Institut "Jožef Stefan"	Remškar Maja	maja.remskar(at)ijs.si	MoS2
Institut "Jožef Stefan"	Suvorov Danilo	danilo.suvorov(at)ijs.si	funkcionalni in bio nanodelci - ele
Institut "Jožef Stefan"	Škapin D. Srečo	sreco.skapin(at)ijs.si	poly(d,1-lactide-co-glycolide)/hy
Institut "Jožef Stefan"	Štrancar Janez	janez.strancar(at)ijs.si	TiO2
Institut "Jožef Stefan"	Vaupotič Janja	janja.vaupotic(at)ijs.si	detekcija aerosolov (5-1100 nm)
Institut "Jožef Stefan"	Vesel Alenka	alenka.vesel(at)ijs.si	Fe2O3
Institut "Jožef Stefan"	Vukomanović Marija	marija.vukomanovic(at)ijs.si	Au
Inštitut za kovinske materiale in tehnologije	Jenko Monika	monika.jenko(at)imt.si	FeOx
Kemijski inštitut	Bele Marjan	marjan.bele(at)ki.si	Carbon NP , TiO2, SiO2, LiMPO4 (M
Kemijski inštitut	Crnjak Orel Zorica	zorica.crnjak.orel(at)ki.si	Cu/ZnO, CuO/ZnO, Pd/CuO/ZnO, f
Kemijski inštitut	Dominko Robert	robert.dominko(at)ki.si	LiMPO4 (M=Mn,Fe) Li2FeSiO4, Car
Kemijski inštitut	Dražič Goran	goran.drazic(at)ki.si	TiO2
Kemijski inštitut	Gaberšček Miran	miran.gaberscek(at)ki.si	Carbon NP , TiO2, SiO2, LiMPO4 (M

Awareness raising



OGNJEMETI in druga zabavna
PIROTEHNIKA
ZASTRUPAJAJO OZRAČJE



- http://www.kemijskovaren.si/files/nano_knjiga.pdf

Additional links:

- http://ec.europa.eu/nanotechnology/index_en.html,
- http://ec.europa.eu/environment/chemicals/nanotech/index_en.htm
- <http://echa.europa.eu/regulations/nanomaterials>
- <https://ec.europa.eu/jrc/en/scientific-tool/jrc-web-platform-nanomaterials>
- <https://ec.europa.eu/jrc/en/scientific-tool/nanohub>
- http://ec.europa.eu/health/nanotechnology/policy/index_en.htm

Thank you for
your attention

<http://www.uk.gov.si/>

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<http://www.ki.si>

Chemometric analysis and QSAR modelling in NANO-toxicology

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Chemometric analysis and QSAR modelling in NANO-toxicology

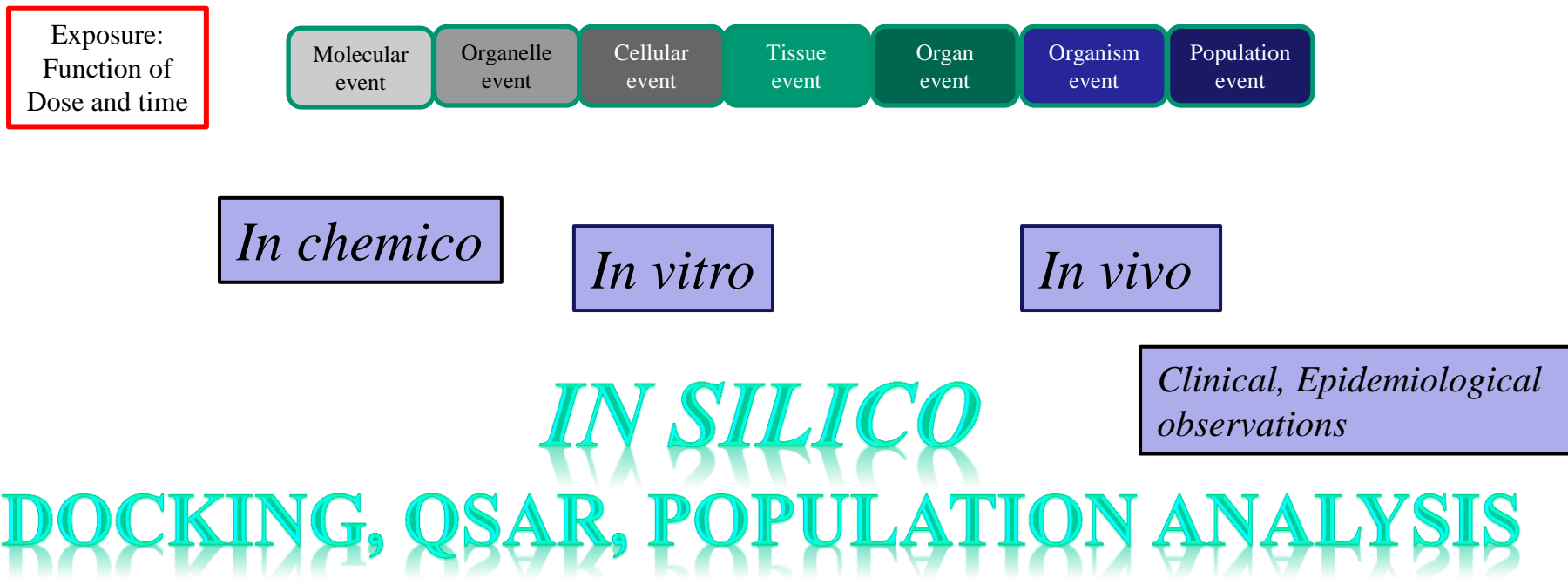
Outline

1. Introduction
2. New toxicological paradigm
3. QSAR (**Q**uantitative **S**tructure-**A**ctivity **R**elationship) working scheme and strategy
4. NANO-descriptors
5. Chemometrical analysis of proteomic data

New paradigm in toxicology...

Adverse Outcome Pathways

Is a sequential chain of causally related events leading to the toxic effect...



OECD document...

Adverse Outcome Pathway (AOP)

OECD.org Data Publications More sites News Job vacancies

OECD BETTER POLICIES FOR BETTER LIVES

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OECD Home > Chemical safety and biosafety > Testing of chemicals > Adverse Outcome Pathways, Molecular Screening and Toxicogenomics

- > Testing of chemicals
- > Assessment of chemicals
- > Risk management of chemicals
- > Chemical accident prevention, preparedness and response
- > Pollutant release and transfer register
- > Safety of manufactured nanomaterials
- > Agricultural pesticides and biocides
- > Biosafety - BioTrack

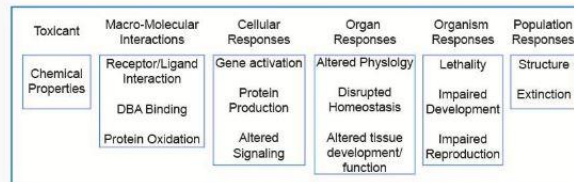
Adverse Outcome Pathways, Molecular Screening and Toxicogenomics

- [Documents](#)
- [How to make a project proposal?](#)
- [Process for the development of AOP at OECD](#)
- [List of projects on the AOP development programme workplan](#)
- [AOP wiki](#)
- [Other activities on Molecular Screening and Toxicogenomics](#)

What is an Adverse Outcome Pathway (AOP)

In 2012, the OECD launched a new programme on the development of Adverse Outcome Pathways. An Adverse Outcome Pathway (AOP) is an analytical construct that describes a sequential chain of causally linked events at different levels of biological organisation that lead to an adverse health or ecotoxicological effect (see figure). AOPs are the central element of a toxicological knowledge framework being built to support chemical risk assessment based on mechanistic reasoning.

Figure: schematic representation of the Adverse Outcome Pathway (AOP) illustrated with reference to a number of pathways.



The AOP development programme addresses the needs of:

- the OECD [Test Guidelines Programme](#) for the identification of new *in vitro* test methods that are candidates to become OECD Test Guidelines;
- The OECD [QSAR Project](#) for the identification of new methods/profilers for grouping chemicals, and;
- the OECD [Hazard Assessment activities](#) for the development of Integrated Approaches to Testing and Assessment (IATA), also known as Integrated Testing Strategies, for defined hazard endpoints.

The OECD co-ordinates its activities with the [WHO/IPC work on Mode of Action](#), as the AOP concept and the Mode of Action are closely related.

Documents

[Guidance, template, format available](#)

[AOP project proposal form](#)

Workshop on COMPUTATIONAL APPROACHES IN NANOSCIENCES
„COMPInNANO“, Ljubljana, 2. - 3. October 2015

Example: skin sensitization (chemicals)...

ENV/JM/MONO(2012)10/PART1

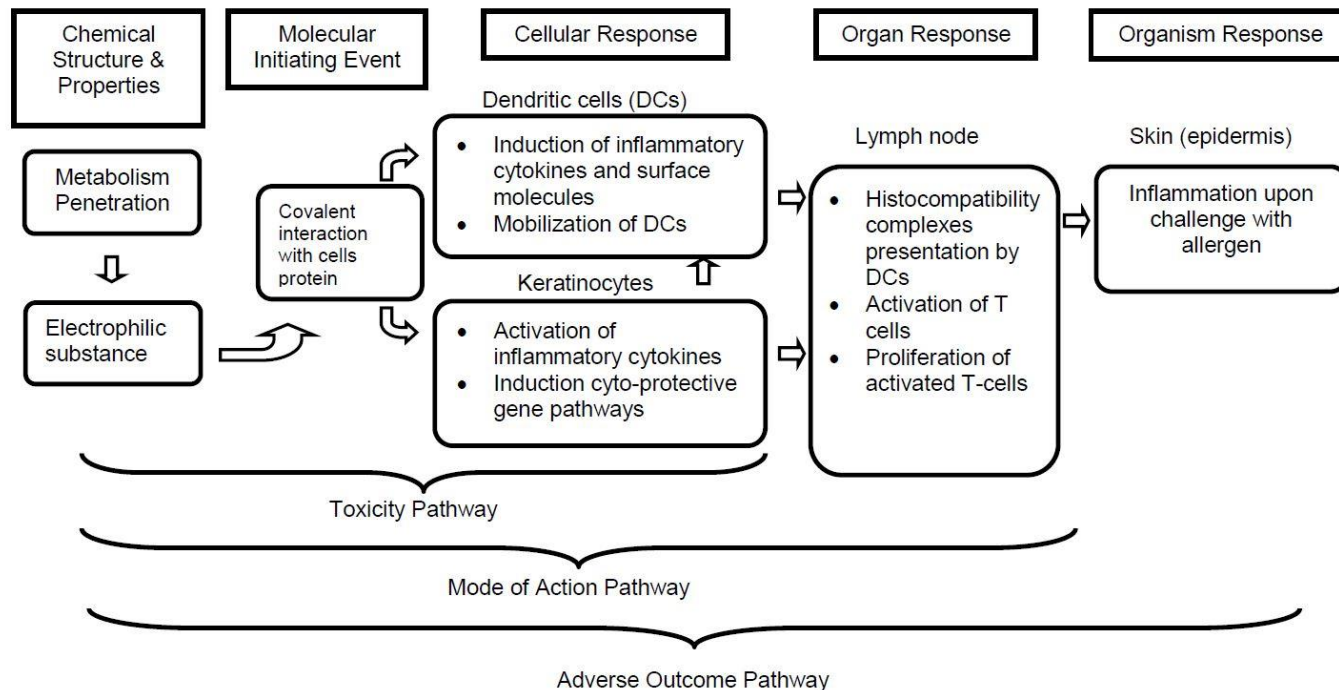
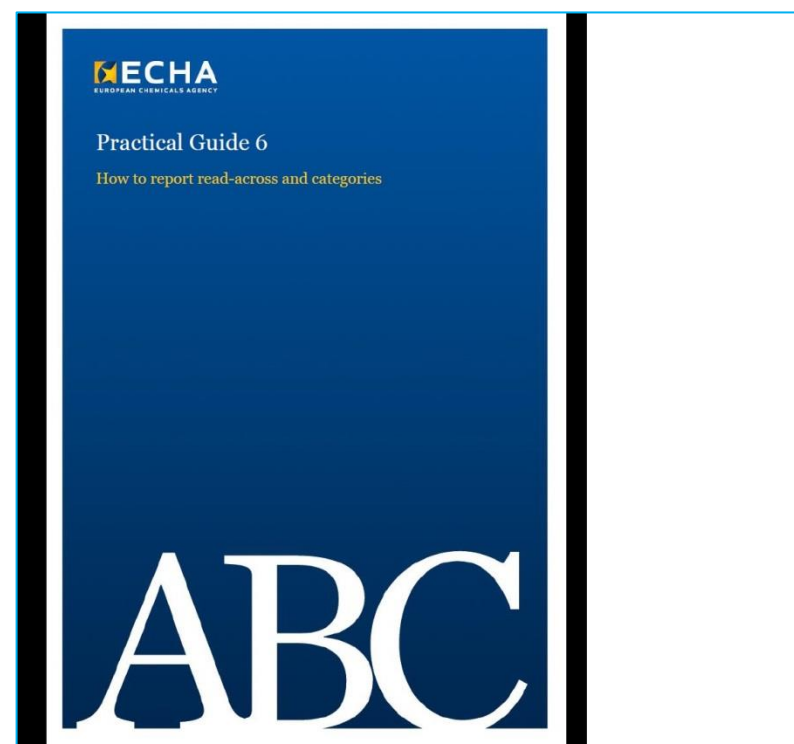
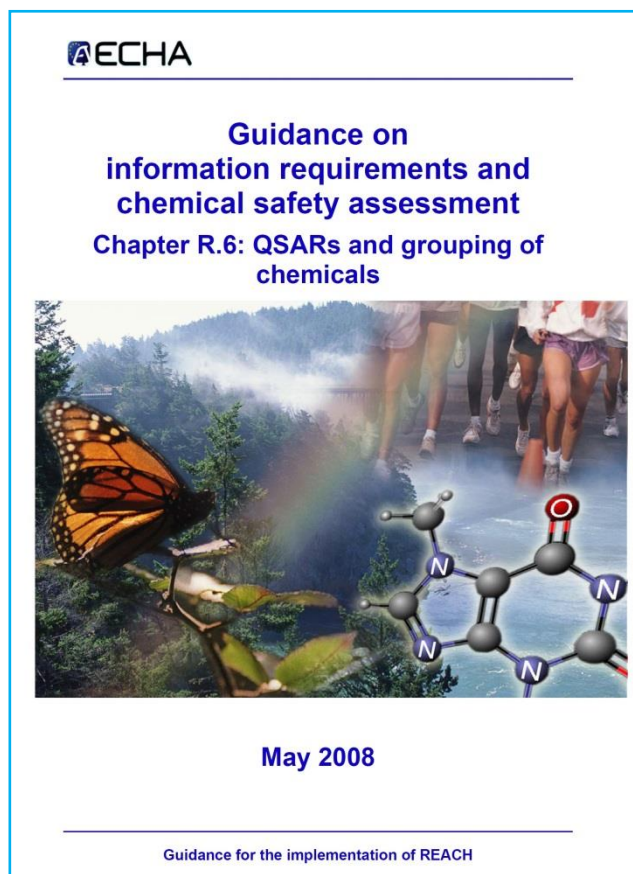


Figure 3. Flow diagram of the pathways associated with skin sensitisation.

In silico methods...

- *In silico* methods are important to acquire the data for AOP scheme...
- Under term *In silico* we understand different computational modelling approaches; molecular mechanics, dynamics, **QSAR**, modeling of ADME properties, and also **chemometrical analysis of (proteomics, genomics, metabonomics) data,...**
- **MOLECULES - NANO particles?**

QSAR, Read across (grouping) in REACH and ECHA



NANO and EU



Brussels, 3.10.2012
SWD(2012) 288 final

COMMISSION STAFF WORKING PAPER

Types and uses of nanomaterials, including safety aspects

Accompanying the

**Communication from the Commission to the European Parliament, the Council and the
European Economic and Social Committee**

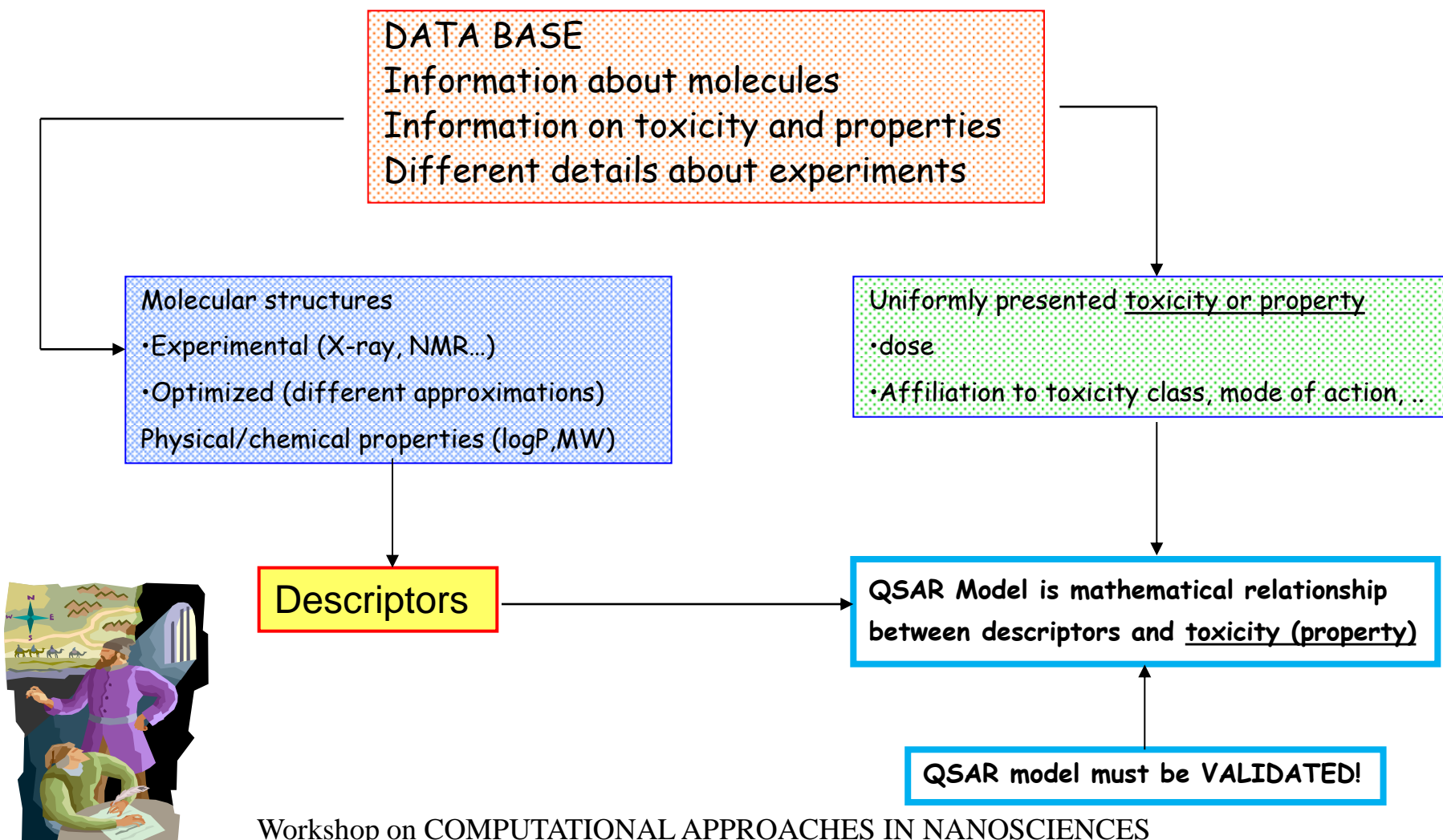
on the Second Regulatory Review on Nanomaterials

{COM(2012) 572 final}

Workshop on COMPUTATIONAL APPROACHES IN NANOSCIENCES
„COMPInNANO“, Ljubljana, 2. - 3. October 2015

Scheme of QSAR - strategy

How to extract the hidden knowledge from DATA SET



Scheme of QSAR - strategy

How to extract the hidden knowledge from DATA SET?

QSAR hypothesis: The property is a function of chemical structure!

The validated QSAR model can be used to predict the property for a '**new - hypothetical**' compound.

ONLY FOR THE PROPERTY FOR WHICH IT WAS TRAINED.



Two main application areas of QSAR models

Active substances (drug) research:

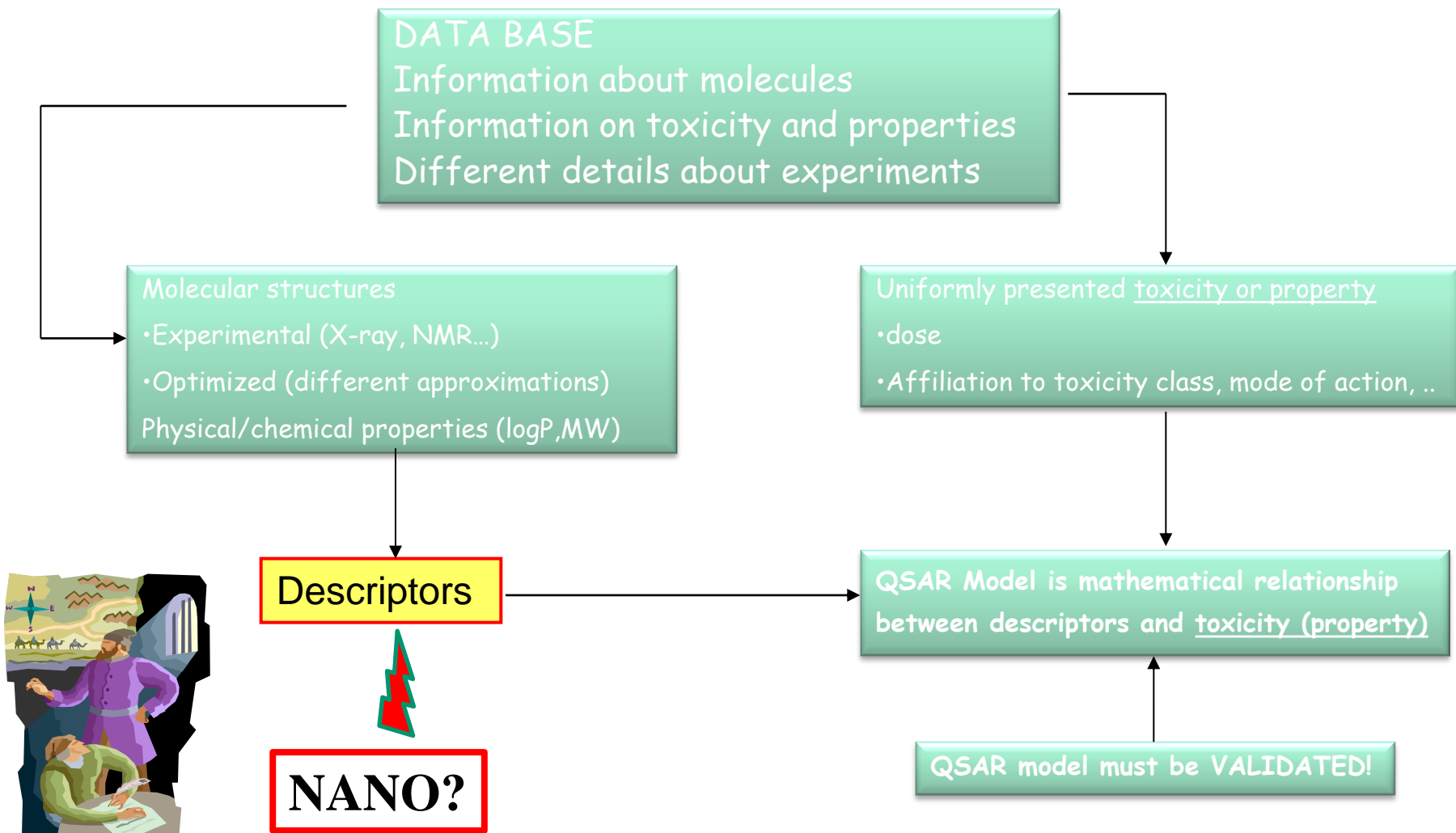
1. Searching for new lead compounds
2. Searching for better analogues

Regulatory assessment:

1. Predicting of 'missing' values for risk assessment
2. Categorisation of compounds for labeling
3. Priority setting

NANO: Scheme of QSAR - strategy

How to extract the hidden knowledge from DATA SET



Molecules – NANO particles ??

The properties of NANO particles (strongly) depend on their the size and the shape.

An accent is placed is on *descriptors*.

Gold nanoparticles are red to black.

Red glass (Wikipedia):

First produced in late Roman Empire.

The knowledge was lost and rediscovered in the 17th century by either Johann Kunckel in Potsdam or by the Florentine glassmaker Antonio Neri in Italy.

Chemist and winner of the 1925 Nobel Prize in Chemistry Richard Adolf Zsigmondy was able to understand and explain that small colloids of gold were responsible for the red colour.

Representation of a molecule - *How to apply for NANO?*

Different levels of the representation of molecules:

- 1D - Information on constituents (which atoms, or which groups of atoms)
- 2D - Structural formula gives information on atoms and bonds between atoms, but no information on metrical parameters (distances between atoms, angles between bonds)
- 3D - Coordinates of all atoms (information on all metrical properties)
- Quantum chemical descriptors are calculated from QC results - they describe the electronic properties
- Structure can be described by fragments
- Etc.



1D descriptors

Molecules (constitutional):

Information on constituents:

- Number of atoms
- Number of particular atom groups (fragments)
- ...

NANO particles:

- Size
- Shape
- Chemical constitution
- ...



2D descriptors - Topological indices

Topological indices are numbers deduced from structural formula of a molecule (2D representation).

NANO particles:

- Shape indices
- Analysis from electron microscopy pictures.

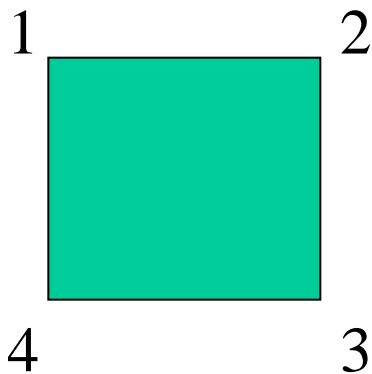
Ghorbani, Modjtaba. Computing Wiener index of C-24 fulleren.

Journal Of Computational And Theoretical Nanoscience, 2015,12, 1847-1851.

Example:

Wiener index (name proposed by H. Hosoya) is an integer number deduced from structural formula (graph).

Wiener index for cyclobutane:



<u>Molecule</u>	<u>W</u>
Methane	0
Propan	4
Cyclopropane	3
N-butane	10
Isobutane	9
Cyclobutane	8

$$\begin{aligned} W &= d_{12} + d_{13} + d_{14} + d_{23} + d_{24} + d_{34} \\ &= 1 + 2 + 1 + 1 + 2 + 1 = 8 \end{aligned}$$

Wiener index has a high degeneracy: different non-isomorphic graphs have the same W .

Attempts to lower the degree of degeneration:

- Hyper-Wiener index
- Szeged index
- Three-dimensional Wiener index
- Hosoya's index Z

Second generation topology indices

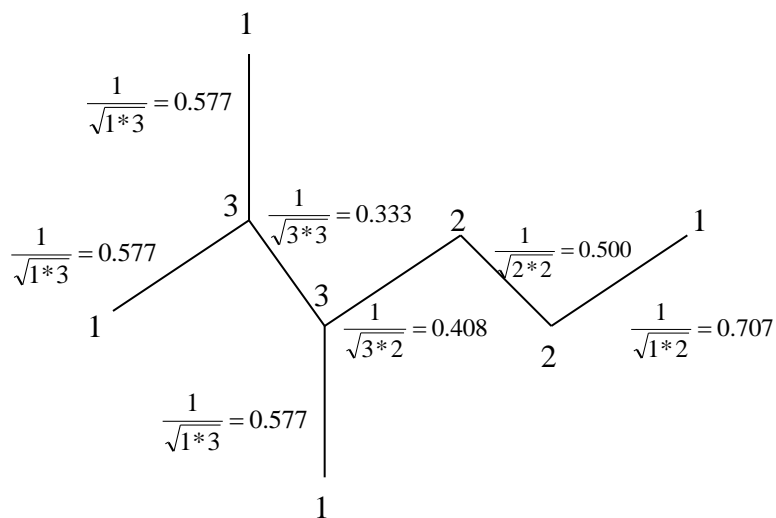
They are real numbers deduced from graphs. One of the most successful is molecular connectivity index c proposed by Randić:

$$\chi = \sum_{ij} (v_i v_j)^{-1/2}$$

Sum runs over edges (bonds), v is the vertex degree on the endpoints.

Calculation of connectivity index for 3-methylheptane

$$\chi = \sum_{ij} (v_i v_j)^{-1/2}$$



$$\chi = 3 \cdot 0.577 + 0.333 + 0.408 + 0.500 + 0.707 = 3.679$$

3D descriptors

Three-dimensional structure of molecules is not unanimously defined.

Rigid molecules are rare, most of the molecules are flexible.

A molecule can have a different 3D structure *in vacuo*, in crystalline form, in water environment, or in protein environment.

Experimental determinations:

- X ray diffraction measurements
- 2D-NMR measurements - method enables determination of ligand-receptor geometry

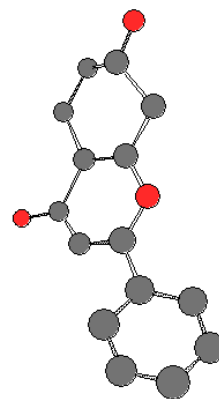
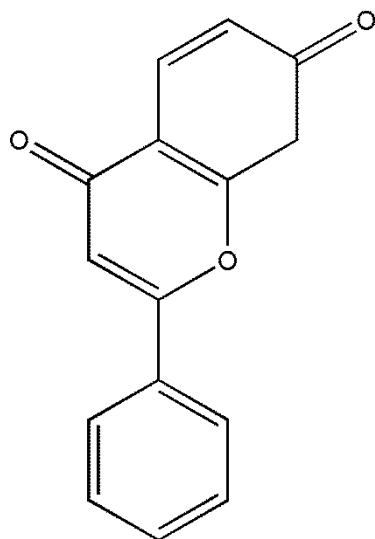
Theoretical determination:

- Quantum chemical optimization
- Molecular dynamics

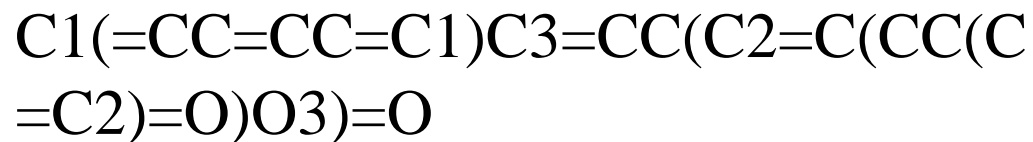
Result (3D structure) depends on the selected method.



A Flavonoid derivate: 3D structure determined with rule based generator

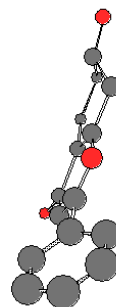


Tautomer of 7-hydroxy-2-phenyl-4-benzopyrone

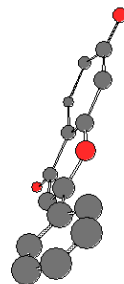


A Flavonoid derivate: 3D structure determinated with rule based generator or
with Molecular Mechanics Optimization

Rule based system



MM optimization



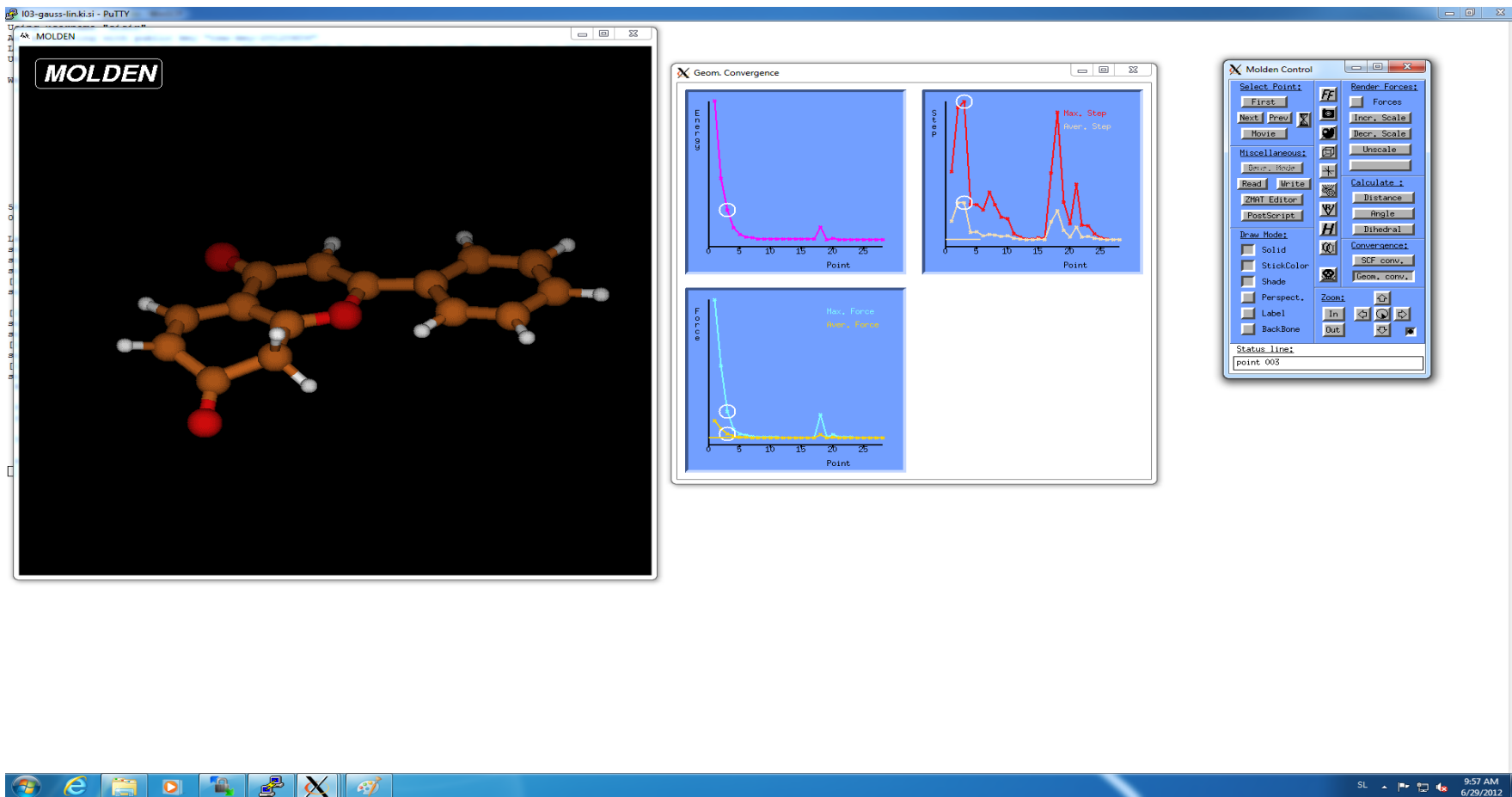
A Flavonoid derivate: 3D structure determined with rule based generator or
with QC program GAUSSIAN (HF-6-31g approximation)
Input

```
# HF/6-31G(d) opt
Test1
0 1
C 0 -0.809889 -2.481124 0.027570
C 0 -0.809889 -3.818124 0.027570
C 0 0.347987 -4.486624 0.027570
C 0 1.505863 -3.818124 0.027570
C 0 1.505863 -2.481124 0.027570
C 0 0.347987 -1.812624 0.027570
C 0 0.347987 -0.475624 0.027570
O 0 -0.853279 0.151261 0.027570
C 0 -0.722044 1.499890 0.027570
C 0 0.450229 2.142809 0.027570
C 0 1.574749 1.394039 0.027570
C 0 1.371375 0.058435 0.027570
C 0 -2.020291 2.245149 0.039936
C 0 -1.793136 3.659064 -0.435801
C 0 -0.663327 4.173114 0.051266
C 0 0.479580 3.479435 0.039418
O 0 2.679662 1.882333 0.027570
O 0 -2.545154 4.254249 -1.170300
H 0 -1.762517 -1.931124 0.027570
H 0 -1.762517 -4.368124 0.027570
H 0 0.347987 -5.586624 0.027570
H 0 2.458491 -4.368124 0.027570
H 0 2.458491 -1.931124 0.027570
H 0 2.266345 -0.581120 0.027570
H 0 -2.427220 2.263510 1.075717
H 0 -2.745599 1.738385 -0.635258
H 0 -0.671772 5.189531 0.471773
H 0 1.444041 4.008410 0.039418
```

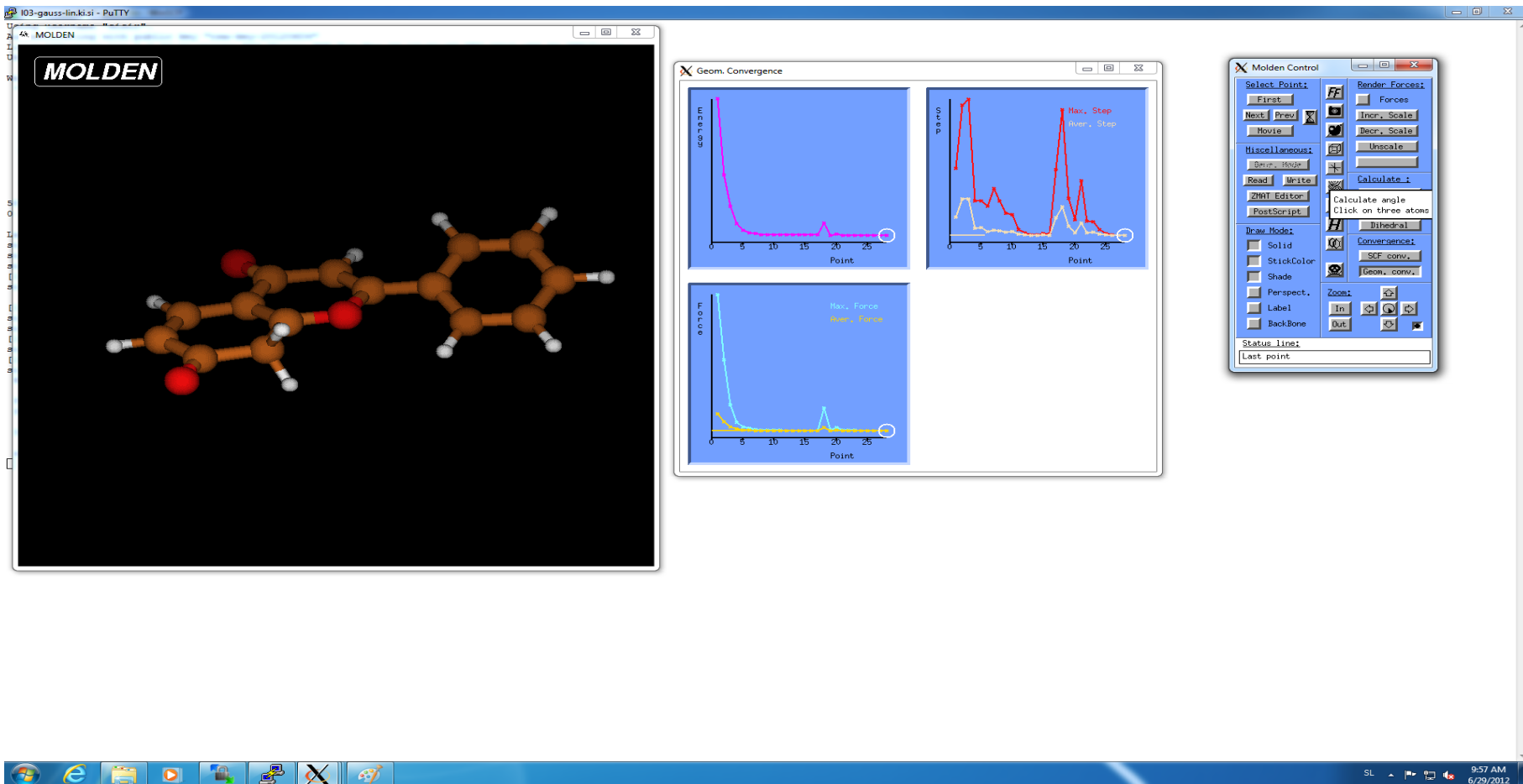
A Flavonoid derivate: 3D structure determined with rule based generator or with QC program GAUSSIAN (HF-6-31g approximation)
Original

The screenshot displays a software interface for molecular visualization and analysis. On the left, a window titled 'MOLDEN' shows a 3D ball-and-stick model of a flavonoid derivative. The central window, 'Geom. Convergence', contains three plots: 'Energy' vs 'Point', 'Force' vs 'Point', and 'Max. Step' vs 'Point'. The 'Energy' plot shows a sharp initial drop followed by a plateau. The 'Force' plot shows a similar trend with a small peak at the end. The 'Max. Step' plot shows a series of peaks and troughs, indicating convergence steps. On the right, the 'Molden Control' panel offers various options for rendering and analysis, including 'Select Point', 'Render Forces', 'Miscellaneous', 'New Model', and 'Zoom'. The Windows taskbar at the bottom shows the system clock as 9:55 AM on 6/29/2012.

A Flavonoid derivate: 3D structure determined with rule based generator or with QC program GAUSSIAN (HF-6-31g approximation)
Midle point



A Flavonoid derivate: 3D structure determined with rule based generator or with QC program GAUSSIAN (HF-6-31g approximation) Final



Quantum chemical descriptors

HF calculations are time consuming. For our molecule from Gaussian output:

•
•

ON A TOMBSTONE, "HERE LIES LESTER MOORE,
FOUR SLUGS FROM A 44, NO LES, NO MORE".

Job cpu time: 0 days 1 hours 26 minutes 23.4 seconds.

File lengths (MBytes): RWF= 58 Int= 0 D2E= 0 Chk= 6 Scr= 1

Normal termination of Gaussian 09 at Thu Jun 28 17:48:11 2012.

Approximations to HF (semiempirical methods) : AM1, CNDO, MINDO, etc.

Quantum chemical descriptors

They are calculated from eigenvalues and eigenvectors of Hartree-Fock equation, or alternatively, from an approximation to it (AM1, CNDO, MINDO, etc.)

Eigenvalues are molecular orbital energies. According Koopmans' theorem:

$$\text{Ionisation potential} = -E_{\text{HOMO}}$$

$$\text{Electron affinity} = -E_{\text{LUMO}}$$

$$\text{Gap} = E_{\text{LUMO}} - E_{\text{HOMO}}$$

Eigenvalues are used to calculate the charge distribution and descriptors related to it (dipole moment and higher moments).

Example: CODESSA calculates about 300 quantum chemical descriptors.

Quantum chemical descriptors

Gaussian output

Our molecule:

1. Optimized geometry
2. $E_{\text{HOMO}} = -0.33343$ Hartree, $E_{\text{LUMO}} = 0.06510$ Hartree
3. Multipole moments:

Electronic spatial extent (au): $\langle R^2 \rangle = 5332.5111$

Charge= 0.0000 electrons

Dipole moment (field-independent basis, Debye):

X= 5.2339 Y= -1.4411 Z= 0.0844 Tot= 5.4293

Quadrupole moment (field-independent basis, Debye-Ang):

XX= -101.3395 YY= -116.5999 ZZ= -103.6957
 XY= -7.8858 XZ= 1.5863 YZ= 4.5278

Traceless Quadrupole moment (field-independent basis, Debye-Ang):

XX= 5.8722 YY= -9.3882 ZZ= 3.5160
 XY= -7.8858 XZ= 1.5863 YZ= 4.5278

Octapole moment (field-independent basis, Debye-Ang²):

XXX= 127.9448 YYY= -41.1843 ZZZ= 1.2466 XYY= 66.4476
 XXY= 59.6734 XXZ= 2.9705 XZZ= -14.6642 YZZ= -3.5759
 YYZ= -1.1079 XYZ= 6.4639

Hexadecapole moment (field-independent basis, Debye-Ang³):

XXXX= -5387.5410 YYYY= -1605.9836 ZZZZ= -157.2575 XXXY= -384.5756
 XXXZ= 75.5886 YYYYX= -15.4537 YYYZ= 32.9132 ZZZX= -3.2400
 ZZZY= -4.7132 XXYY= -1293.9863 XXZZ= -1012.2994 YYZZ= -250.1014
 XXYZ= 57.5382 YYXZ= -5.9352 ZZXY= 21.7033

Etc.

Quantum chemical descriptors in NANO

A promised area in NANO QSAR, possible descriptors:

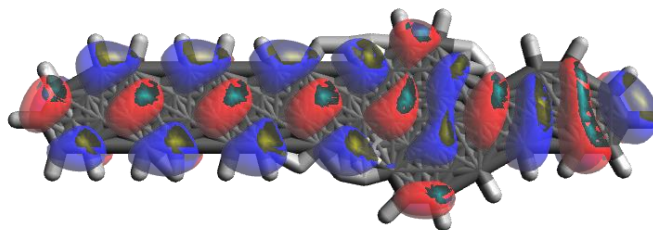
- gaps,
- proton, electron affinity,
- surface charges,
- electronic properties related to reactivity (formation of metal cations).

Problems:

- NANO particles are large, poorly defined systems. *Ab initio* calculations require large scale computer abilities.

I. Lynch et al. A strategy for grouping of nanomaterials based on key physico-chemical descriptors as a basis for safer-by-design NMs. *Nano Today*, 2014, 9, 266-270.

Electron distribution of graphene



Descriptors - résumé

- Descriptors are parameters, which represent the molecular structure in QSAR model
- Different chemical and physical parameters can be used as descriptors (logP, solubility, etc.)
- Thousands of descriptors can be calculated from chemical structures
- Dozens of programs are available (commercial and free) to calculate the descriptors (DRAGON, CODESSA, POLLY, MDL, PETRA.....)
- **We are far from clear NANO-QSAR concept...**
- **Far from NANO particle design (hypothetical NP ?)**

E. Burello, A. Worth. Predicting toxicity of nanoparticles. Nature Nanotechnology, 2011, 6, 138-139.

NANO data bases relevant for toxicity assessment

N. Jeliaskova et al. The eNanoMapper database for nanomaterial safety information. *Beilstein J. Nanotechnology*, 2015, 6, 1609-1634.

Data bases relevant for toxicity assessment:

- <http://www.nanomaterialregistry.org>
- <http://www.nanoparticlelibrary.net>
- <http://nbi.oregonstate.org>
- <http://cananolab.nci.nih.gov/caNanoLab/>
- <http://www.internano.org>
- <http://icon.rice.edu/report.cfm>
- <http://ncl.cancer.gov>
- <http://www.napira.eu>
- <http://nanopartikel.info>
- <http://nanowerk.com>
- <http://www.nanosafetycluster.eu/>

Chemometrical analysis of -omic data..

In vitro study:

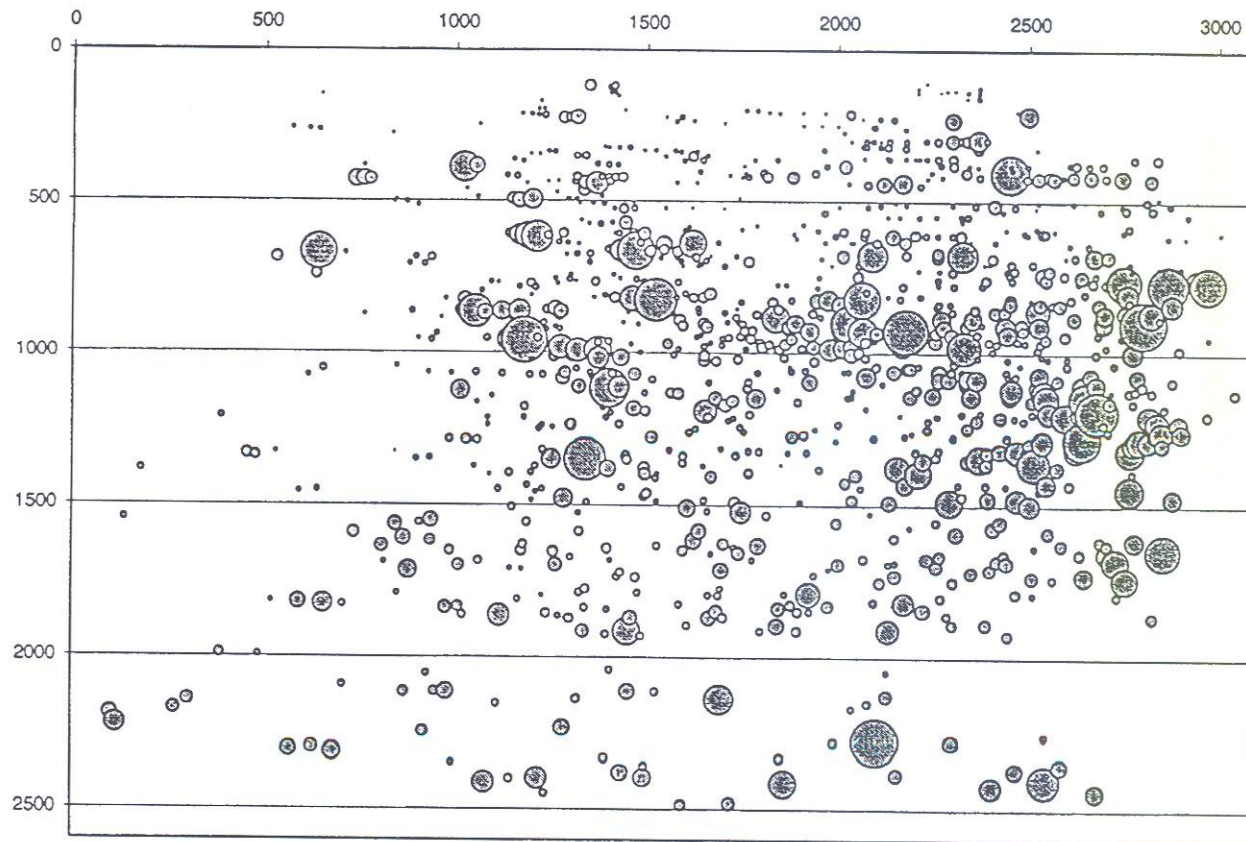
1. Cells are treated with different NANO particles...
2. The -omic status is measured (ca 3000 proteins pro measurements)
3. Statistical relevant -omics are selected?



Pilot study 1: Biodescriptors-2D proteomic maps

Study :

M. Vracko, S. C. Basak. Similarity study of proteomic maps. *Chemometrics and Intelligent Laboratory Systems* 70 (2004) 33–38



Data from:

N.L. Anderson, R. Esquer-Blasco, F. Richardson, P. Foxworthy, P. Eacho, The effects of peroxisome proliferators on protein abundances in mouse liver, *Toxicol. Appl. Pharmacol.* 135 (1996) 75–89.

Similarity indices between control map and treated maps are reported:

The similarity index:
$$s^{a,b} = \frac{1}{N} \sum_{i=1}^N \frac{z_i^a z_i^b}{\max(z_i^a, z_i^b)^2}$$

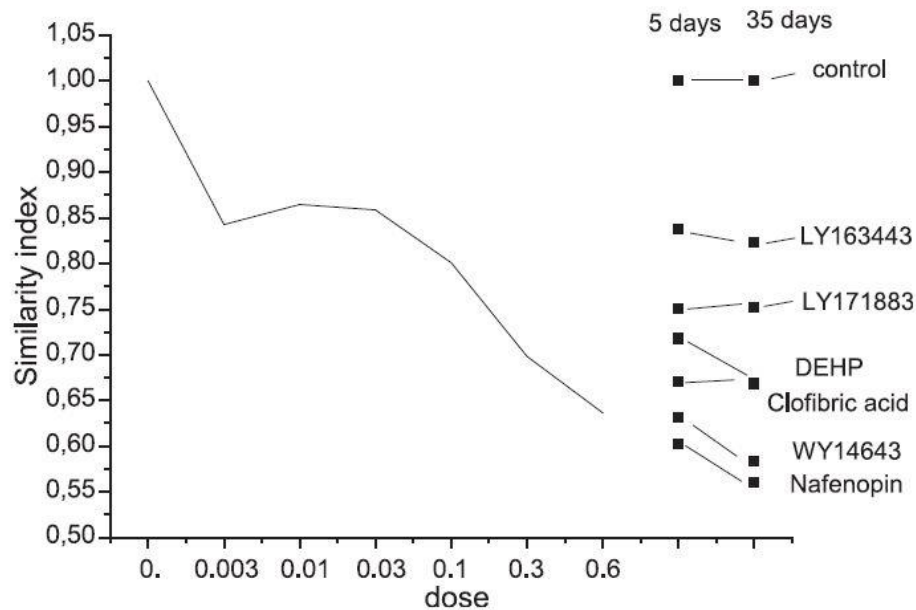


Fig. 2. Similarity index dependent on administrated dose of LY171883. On spots the right side show similarity index for 5 and 35 days exposures.

Study 2: Information on biological status of cells biodescriptors, proteomics

M. Vracko, S. C. Basak, K. Geiss, F. Witzmann. Proteomic Maps-Toxicity Relationship of Halocarbons Studied with Similarity Index and Genetic Algorithm. *J. Chem. Inf. Model.* 2006, 46, 130-136.

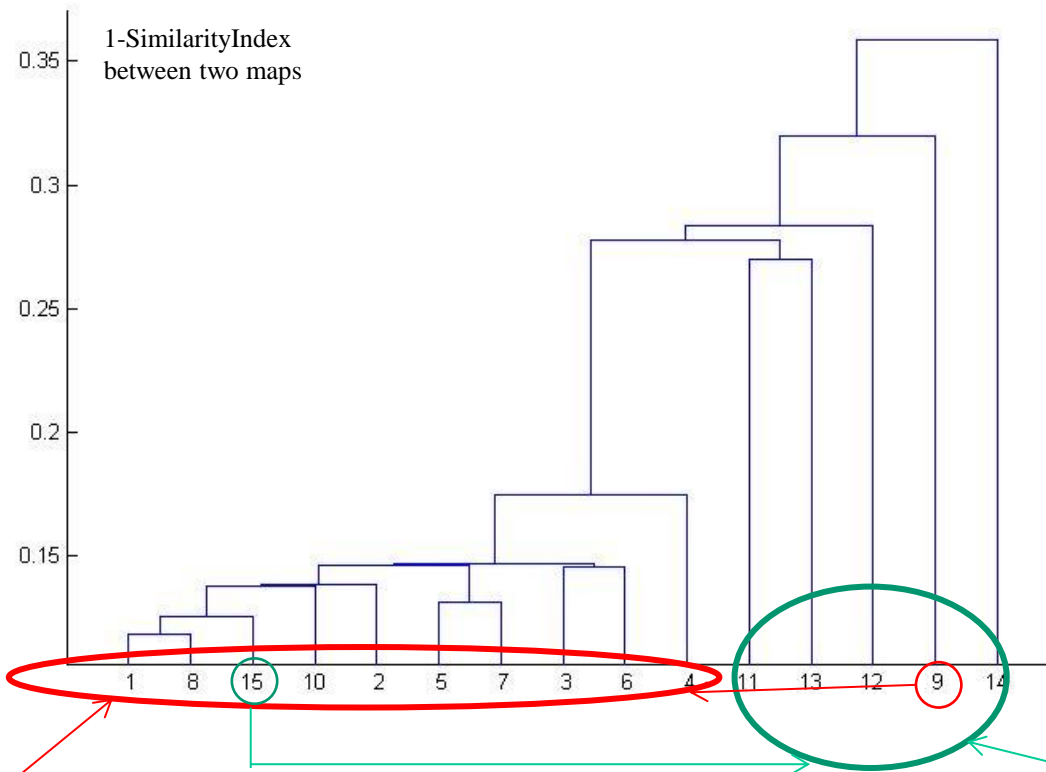
- The comparison of proteomic maps obtained from hepatocytes, which were treated 14 halocarbons.
- Six biological endpoints were determined *in vitro*: $EC50_{MIT}$, LEC_{ROS} , $EC20_{SH}$, $EC50_{LDH}$, LEC_{LP} , LEC_{CAT}
- From each map 263 spots were taken to study the similarity among maps.
- The similarity between two maps was expressed with similarity index.
- The clustering structure is graphically presented with hierarchical clustering method
- With genetic algorithm we selected proteins related to endpoints.

Information on biological status of cells biodescriptors, proteomics

The similarity index:
$$s^{a,b} = \frac{1}{N} \sum_{i=1}^N \frac{z_i^a z_i^b}{\max(z_i^a, z_i^b)^2}$$

	control	111- C2Cl3H3	112- C2Br3H3	112- C2Cl3H3	12-C2Br2H4	12- C2BrClH4	12-C2Cl2H4	C2Cl3H	C2Cl4	C2Cl4H2	CBr2H2	CBrClH2	CCl2H2	CCl3H	CHBr3
control	1.0000	0.8614	0.8529	0.7890	0.8323	0.8380	0.8540	0.8816	0.6426	0.8438	0.6850	0.6972	0.7220	0.5177	0.8748
111- C2Cl3H3	0.8614	1.0000	0.8270	0.7762	0.8128	0.8419	0.8426	0.8576	0.6685	0.8577	0.6715	0.6736	0.6914	0.5205	0.8498
112- C2Br3H3	0.8529	0.8270	1.0000	0.7812	0.7699	0.8542	0.8209	0.8413	0.6458	0.8484	0.6553	0.6751	0.6890	0.5081	0.8451
112- C2Cl3H3	0.7890	0.7762	0.7812	1.0000	0.8091	0.7777	0.8251	0.7521	0.6801	0.7607	0.6718	0.6585	0.6685	0.5404	0.7637
12- C2Br2H4	0.8323	0.8128	0.7699	0.8091	1.0000	0.7866	0.8688	0.7926	0.6483	0.7842	0.6822	0.6725	0.6737	0.5431	0.7865
12- C2BrClH4	0.8380	0.8419	0.8542	0.7777	0.7866	1.0000	0.8279	0.8409	0.6287	0.8400	0.6383	0.6489	0.6489	0.5007	0.8300
12- C2Cl2H4	0.8540	0.8426	0.8209	0.8251	0.8688	0.8279	1.0000	0.8081	0.6620	0.8208	0.6839	0.6851	0.6847	0.5269	0.8195
C2Cl3H	0.8816	0.8576	0.8413	0.7521	0.7926	0.8409	0.8081	1.0000	0.6281	0.8332	0.6618	0.6900	0.7025	0.5050	0.8746
C2Cl4	0.6426	0.6685	0.6458	0.6801	0.6483	0.6287	0.6620	0.6281	1.0000	0.6473	0.6390	0.6243	0.6187	0.5812	0.6519
C2Cl4H2	0.8438	0.8577	0.8484	0.7607	0.7842	0.8400	0.8208	0.8332	0.6473	1.0000	0.6328	0.6694	0.6837	0.5008	0.8619
CBr2H2	0.6850	0.6715	0.6553	0.6718	0.6822	0.6383	0.6839	0.6618	0.6390	0.6328	1.0000	0.6712	0.7298	0.6413	0.6657
CBrClH2	0.6972	0.6736	0.6751	0.6585	0.6725	0.6489	0.6851	0.6900	0.6243	0.6694	0.6712	1.0000	0.7162	0.5275	0.6750
CCl2H2	0.7220	0.6914	0.6890	0.6685	0.6737	0.6489	0.6847	0.7025	0.6187	0.6837	0.7298	0.7162	1.0000	0.5355	0.7084
CCl3H	0.5177	0.5205	0.5081	0.5404	0.5431	0.5007	0.5269	0.5050	0.5812	0.5008	0.6413	0.5275	0.5355	1.0000	0.5048
CHBr3	0.8748	0.8498	0.8451	0.7637	0.7865	0.8300	0.8195	0.8746	0.6519	0.8619	0.6657	0.6750	0.7084	0.5048	1.0000

Hierarchical clustering of 14 halocarbons + control



Two carbon atoms

One carbon atom



M. Vracko, S. C. Basak, K. Geiss, F. Witzmann. Proteomic Maps-Toxicity Relationship of Halocarbons Studied with Similarity Index and Genetic Algorithm. *J. Chem. Inf. Model.* 2006, 46, 130-136.

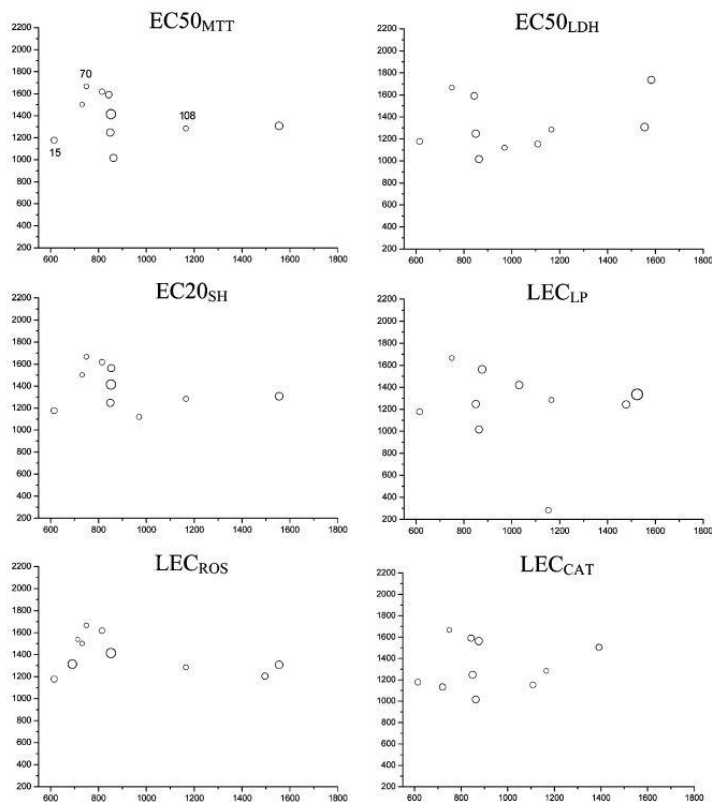


Figure 6. Ten mostly occurring spots for six different biological parameters as they appear in the 2D gel. The size of the bubbles is proportional to the logarithm of intensity. (The integrated intensity of protein spots was calculated by the image analysis software, PDQuest.) In the first picture ($EC50_{MTT}$) the spots 15, 70, and 108 are labeled.



Laboratory for chemometrics

Thank you for your attention!

Researchers

- Marjana Novič
- Marjan Tušar
- Natalja Fjodorova
- Marjan Vračko
- Špela Župerl
- Viki Drgan
- Nikola Minovski
- Katja Venko

Young researchers

- Alja Plošnik
- Jure Borišek
- Lidija Avsenik

Emeritus

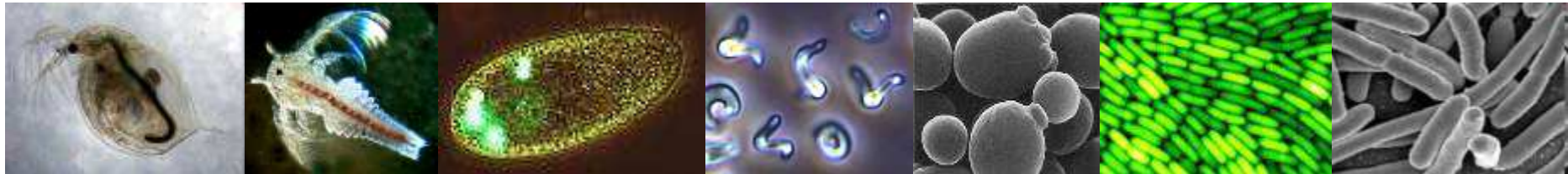
- Prof. Jure Zupan
- Prof. Milan Randić



Toxicity of metal oxide nanoparticles: FP7 project MODERN

Villem Aruoja, PhD
Laboratory of Environmental Toxicology,
National Institute of Chemical Physics and Biophysics,
Tallinn, Estonia

Our zoo



Crustacea

Daphnia magna

Themnocephalus platyurus

Protozoa

Tetrahymena thermophila

Algae

Pseudo-kirchneriella subcapitata

Yeast

Saccharomyces cerevisiae

Bacteria

Vibrio fischeri

Escherichia coli

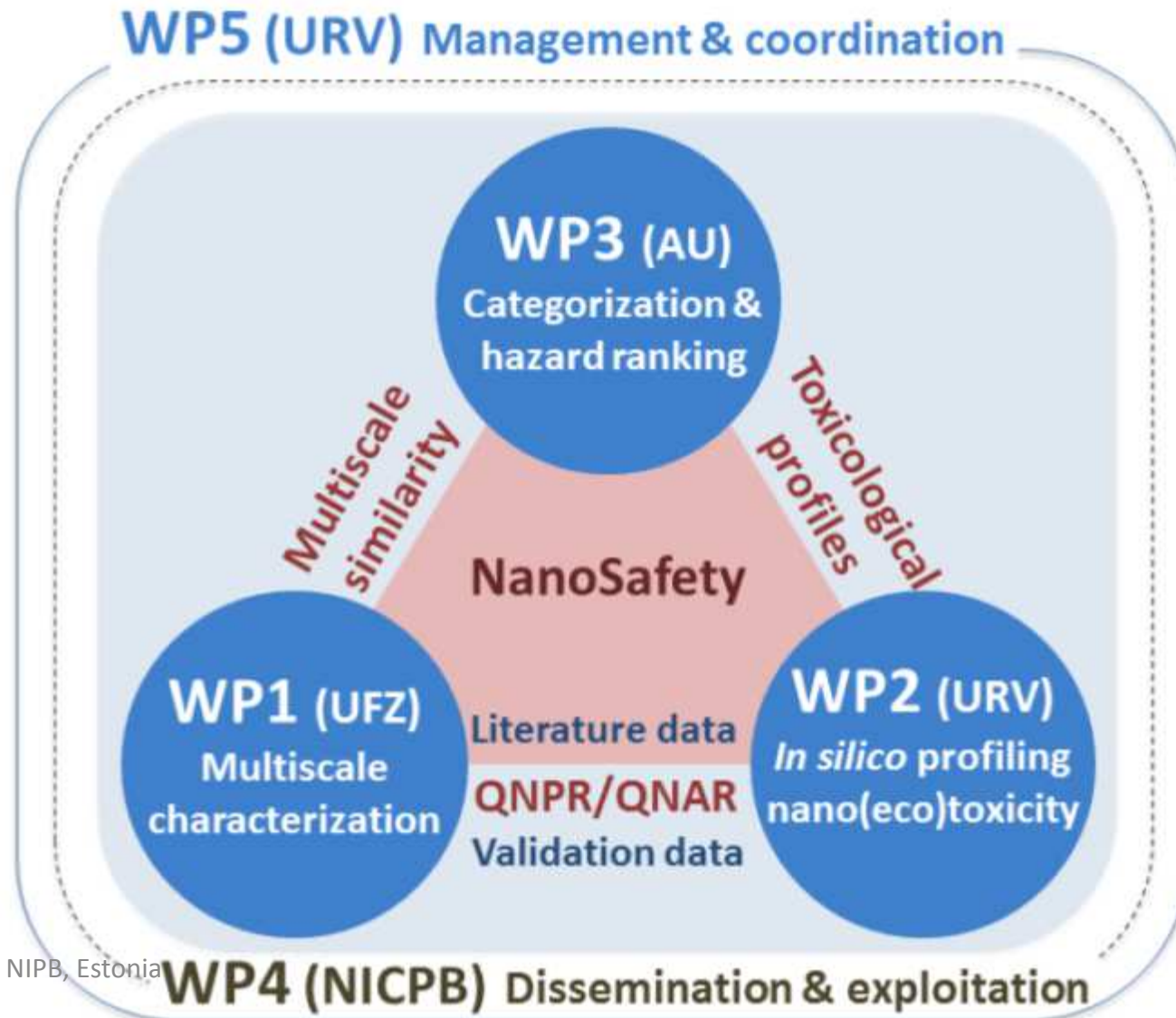
Consumers

Primary producers

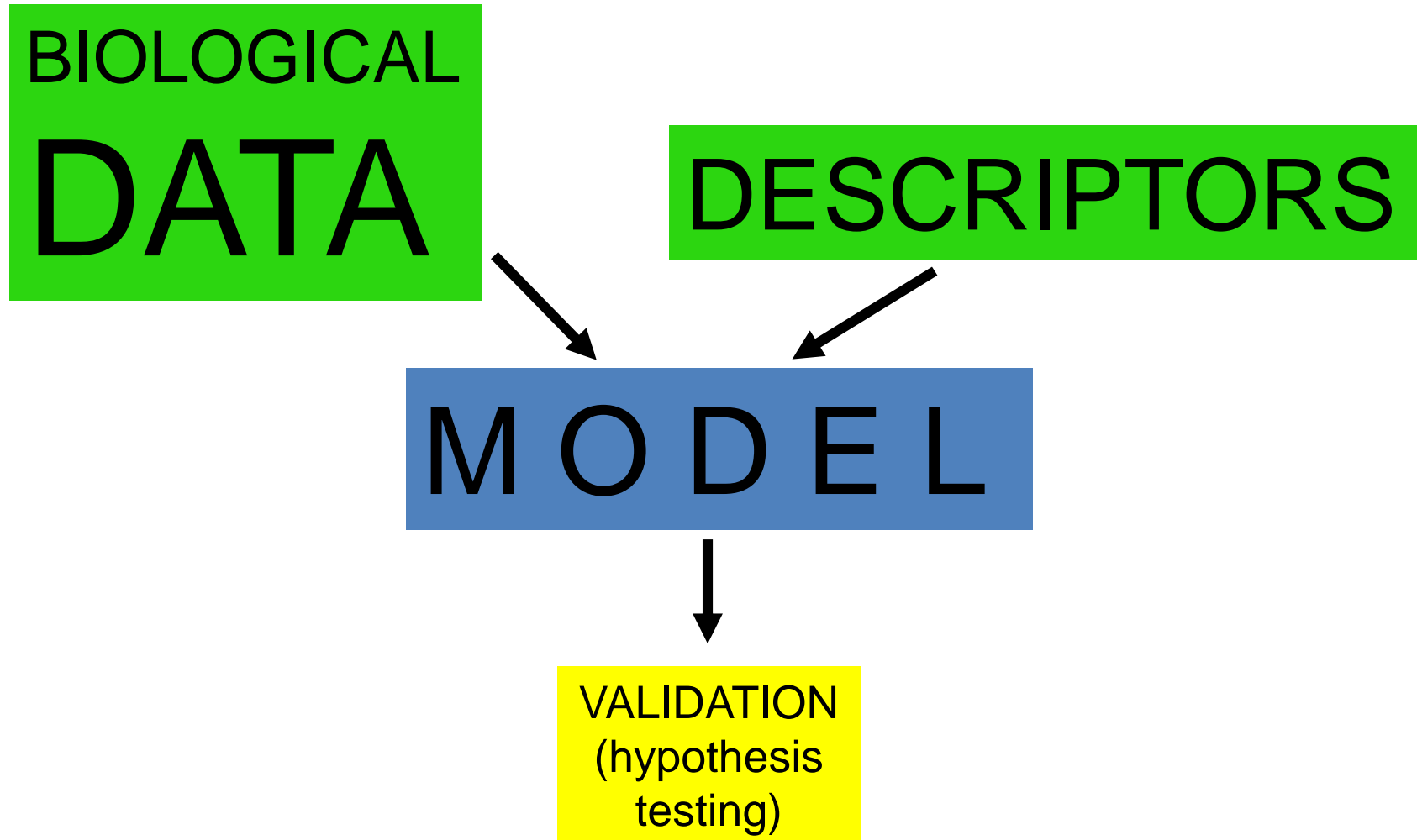
Decomposers

human and animal cell lines
 (human alveolar epithelial cells **A549**, human epithelial colorectal cells **Caco2**, murine fibroblast cell line **Balbc/3T3**)

MODeling the EnviRonmental and human health effects of Nanomaterials



QNAR



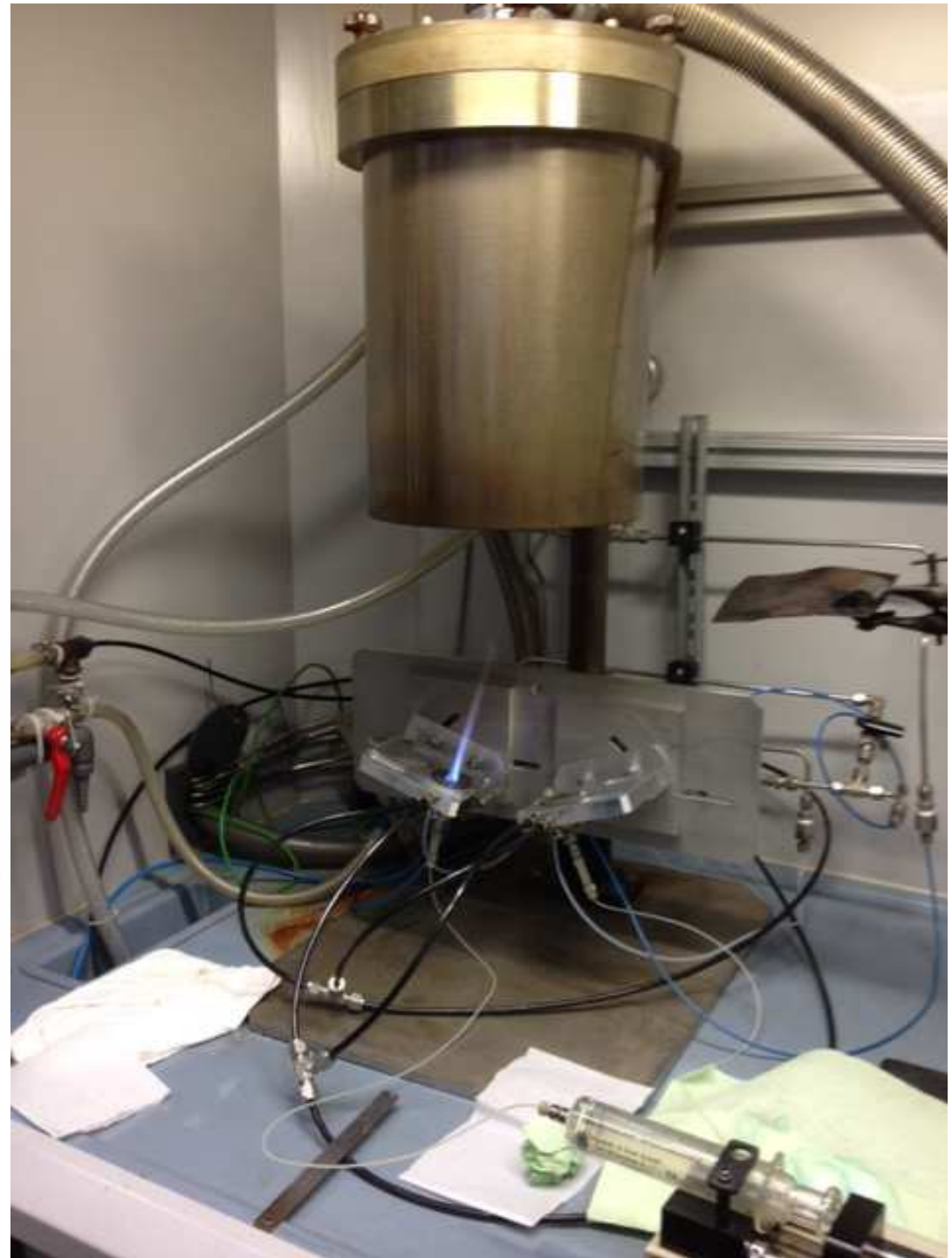
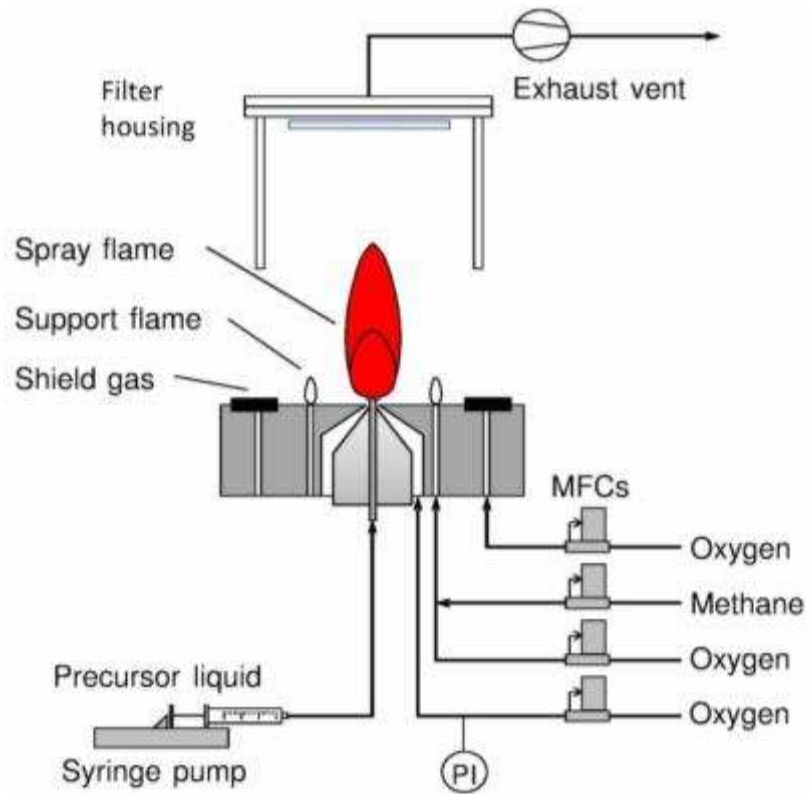
Data?

- Literature?
- Industry?

Andre Nel*: “You have 2,5 years left, start generating data now!”

- * Director of University of California's Center for Environmental Implications of Nanotechnology

Flame Spray Pyrolysis



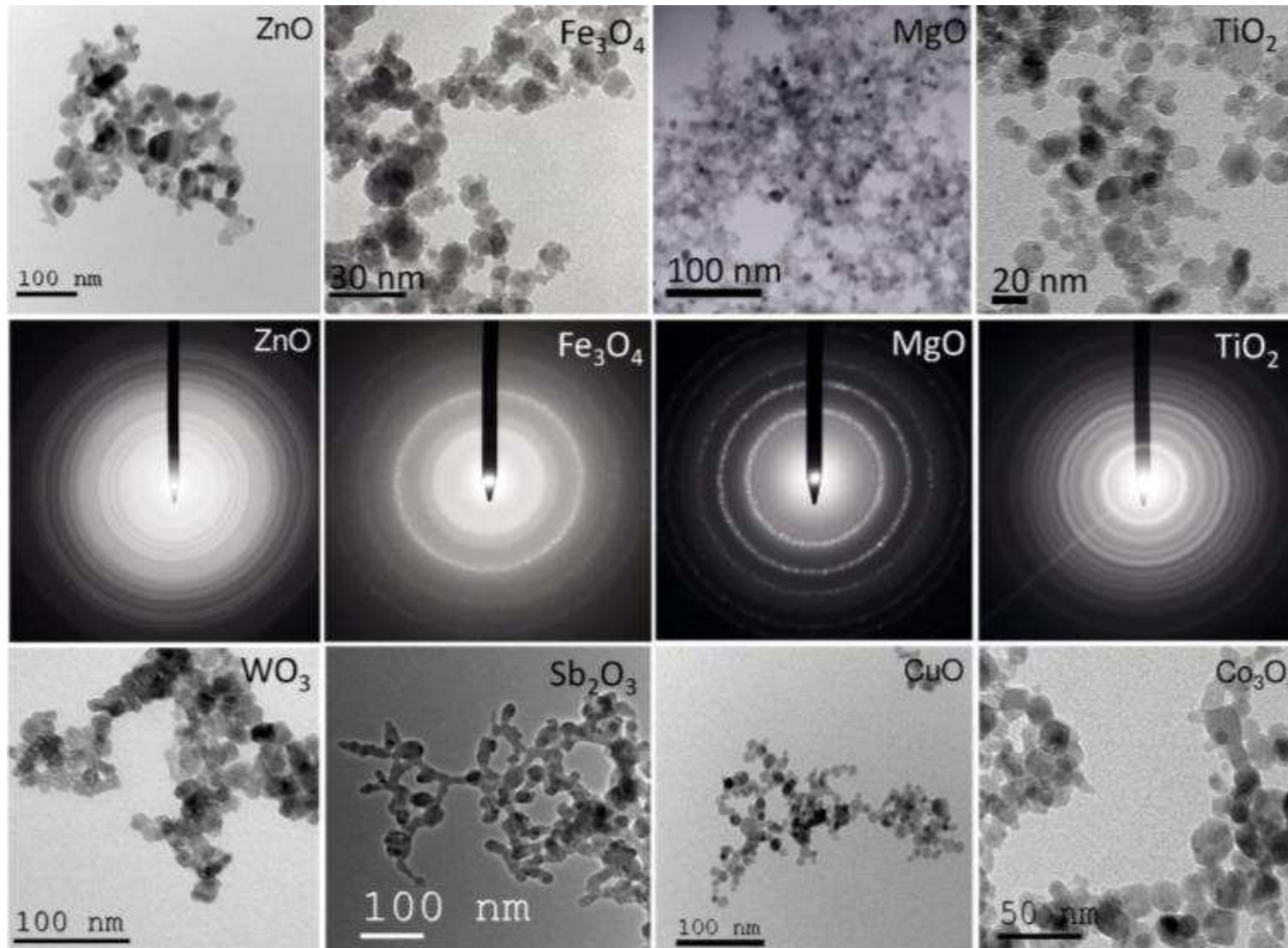
Selection of nanoparticles

- CuO, Co_3O_4 , Sb_2O_3 , TiO_2 , WO_3 , ZnO, Mn_3O_4 , Fe_3O_4 , MgO, Al_2O_3 , SiO_2 , Pd
(Initial subset in project description: 23 metal oxides: ZnO, CuO, Co_3O_4 , Fe_3O_4 , Sb_2O_3 , TiO_2 , WO_3 , Al_2O_3 , CeO_2 , Y_2O_3 , CoO, Ni_2O_3 , Cr_2O_3 , Fe_2O_3 , HfO_2 , In_2O_3 , SnO_2 , ZrO_2 , Gd_2O_3 , La_2O_3 , Mn_2O_3 , NiO, Yb_2O_3 and Ag, Au, Pt)

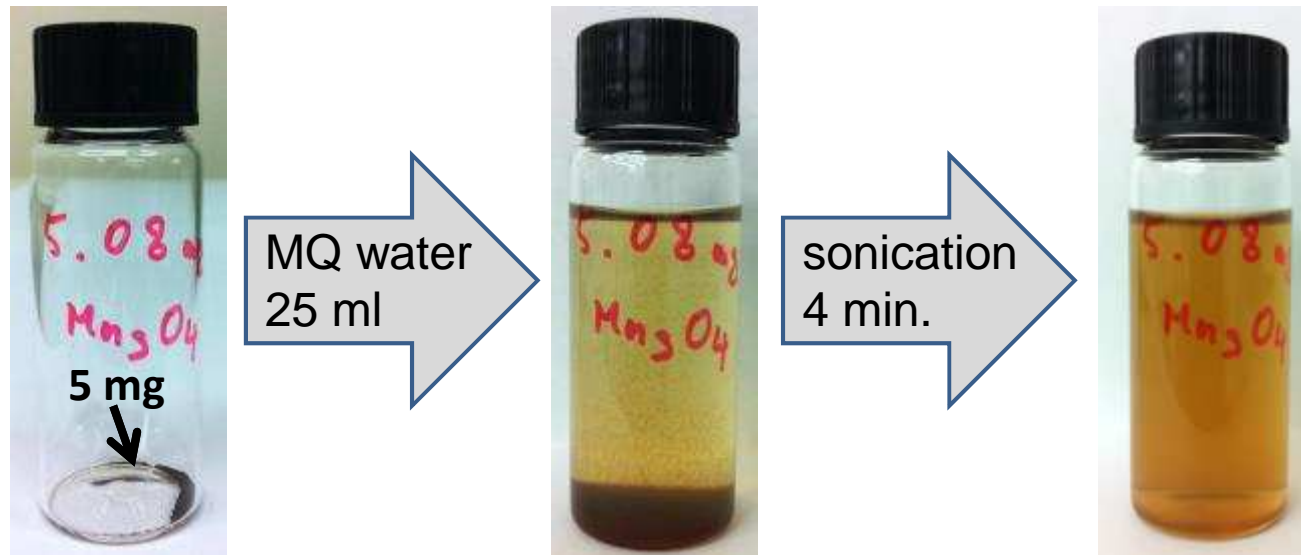




FSP – small and crystalline NPs



Preparation of MOx NPs suspensions

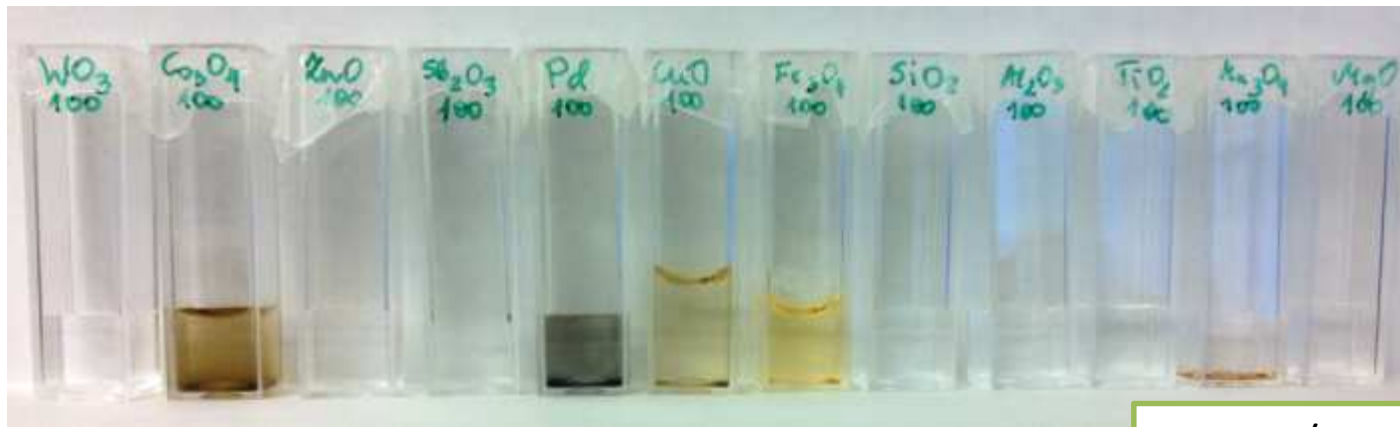


Size, hydrodynamic size

Sample	Specific surface area (SSA); m ² /g	BET size (d_{BET})		DI water				Algal growth medium (pH=8.0)			
		nm	z-average hydrodynamic size, nm	ζ -potential, mV	pH	Metal solubility		z-average hydrodynamic size, nm	ζ -potential, mV	Metal solubility	
						% at 10 mg/l (AAS or ICP-MS*)	% at 200 mg/l (TXRF)			% at 10 mg/l (AAS or ICP-MS*)	% at 100 mg/l (TXRF)
ZnO	53	20.4	171	16.4	6.6	56.1*	5.0	696	-13.1	25.7	3.18
Pd	33	15.1	127	-27.8	6.1	<0.5	NA	151	-18.6	0.24	NA
CuO	72	13.1	130	17	6.2	5.14	0.88	769	-6.2	1.16	0.26
Co ₃ O ₄	85	11.5	99	23	6.1	1.25	6.8	916	10.7	0.18	0.82
TiO ₂	123	12.2	171	-13.6	6.2	<0.83	0.10	717	-15.1	0.42	0.01
Mn ₃ O ₄	81	15.2	395	-14.4	7.0	11.1	4.8	920	-9.8	9.45	6.62
Fe ₃ O ₄	120	9.7	128	22.2	5.9	<1.38	7.1	1005	-12.1	1.66	0.17
Al ₂ O ₃	134	11.4	95	39.2	6.0	0.40*	NA	1232	8.9	0.42	NA
SiO ₂	289	7.8	148	-33.2	6.0	NA	NA	154	-19.8	NA	NA
WO ₃	79	10.6	63	-45.3	5.0	63.2*	2.3	191	-20.4	66.7	75.7
MgO	123	13.6	1964	6.9	9.6	38.1	NA	1581	6.4	87.9 [†]	NA
Sb ₂ O ₃	56	20.5	125	-24.3	4.2	56.3	NA	414	-15.9	21.2	NA

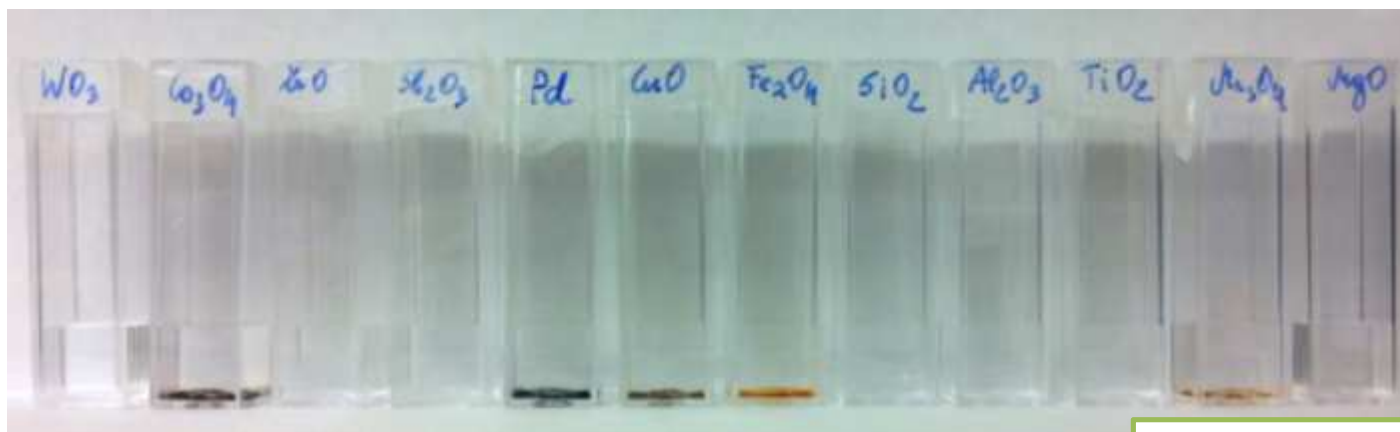
Stability of suspensions in MQ water vs algal test medium

In MQ water: 1 week



100 mg/L

In algal test medium (OECD 201): 1 day.



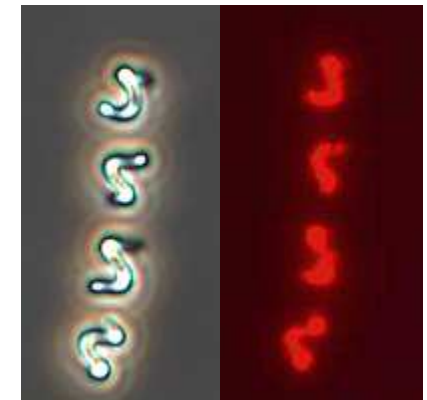
100 mg/L

Composition of the algal test medium.

	Component	mg/L
1	NH ₄ Cl	15
2	MgCl ₂ *6H ₂ O	12
3	CaCl ₂ *2H ₂ O	18
4	MgSO ₄ *7H ₂ O	15
5	KH ₂ PO ₄	1,6
6	NaHCO ₃	50
7	Na ₂ EDTA*2H ₂ O	0,1
8	FeCl ₃ *6H ₂ O	0,08
9	H ₃ BO ₃	0,185
10	MnCl ₂ *4H ₂ O	0,415
11	ZnCl ₂	0,003
12	CoCl ₂ *6H ₂ O	0,0015
13	Na ₂ MoO ₄ *2H ₂ O	0,007
14	CuCl ₂ *2H ₂ O	0,00001

Algal growth inhibition assay (OECD 201)

- Primary producers
- Sensitive to toxicants
- Very sensitive to heavy metals



100 μm

EC₅₀

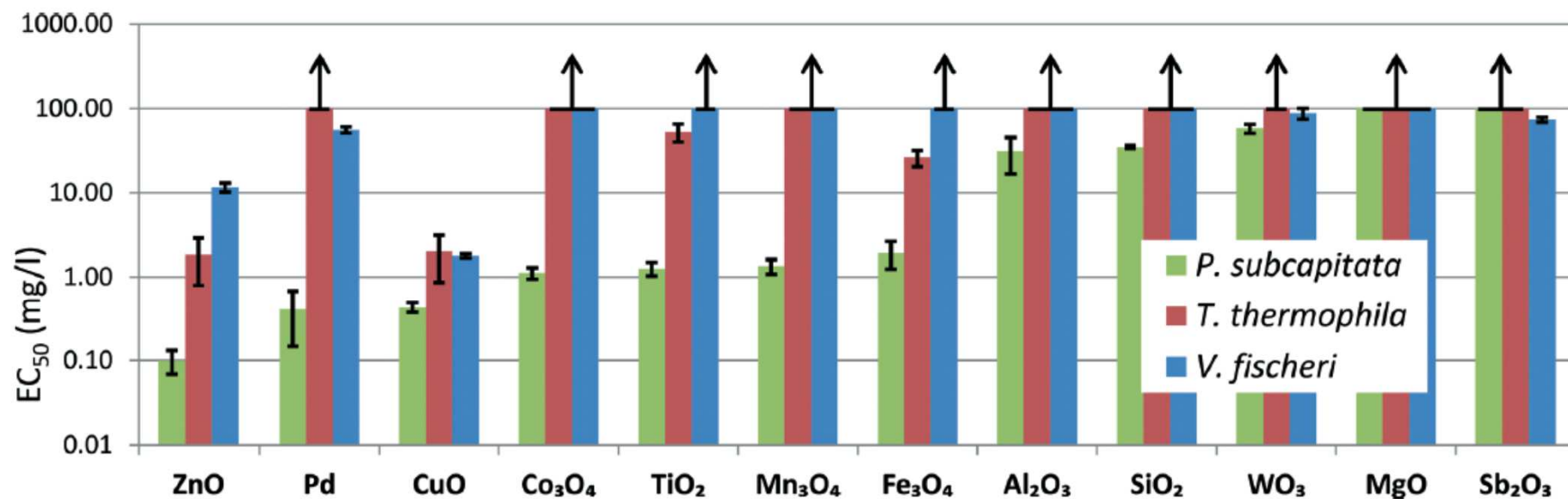
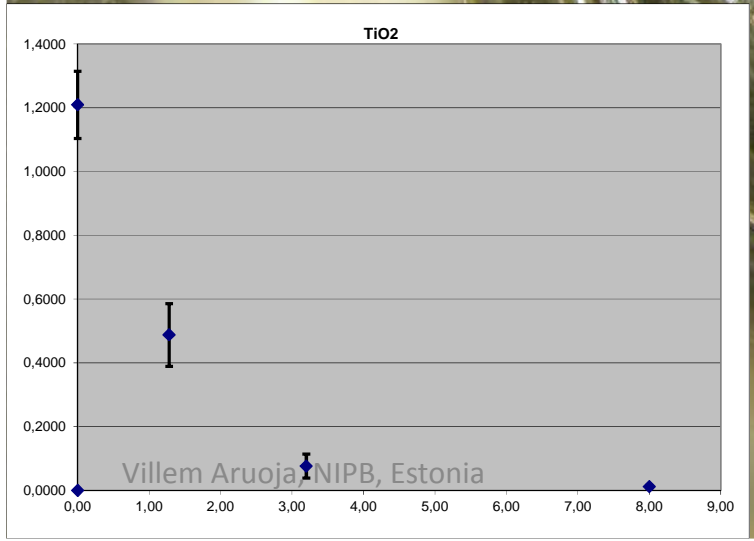
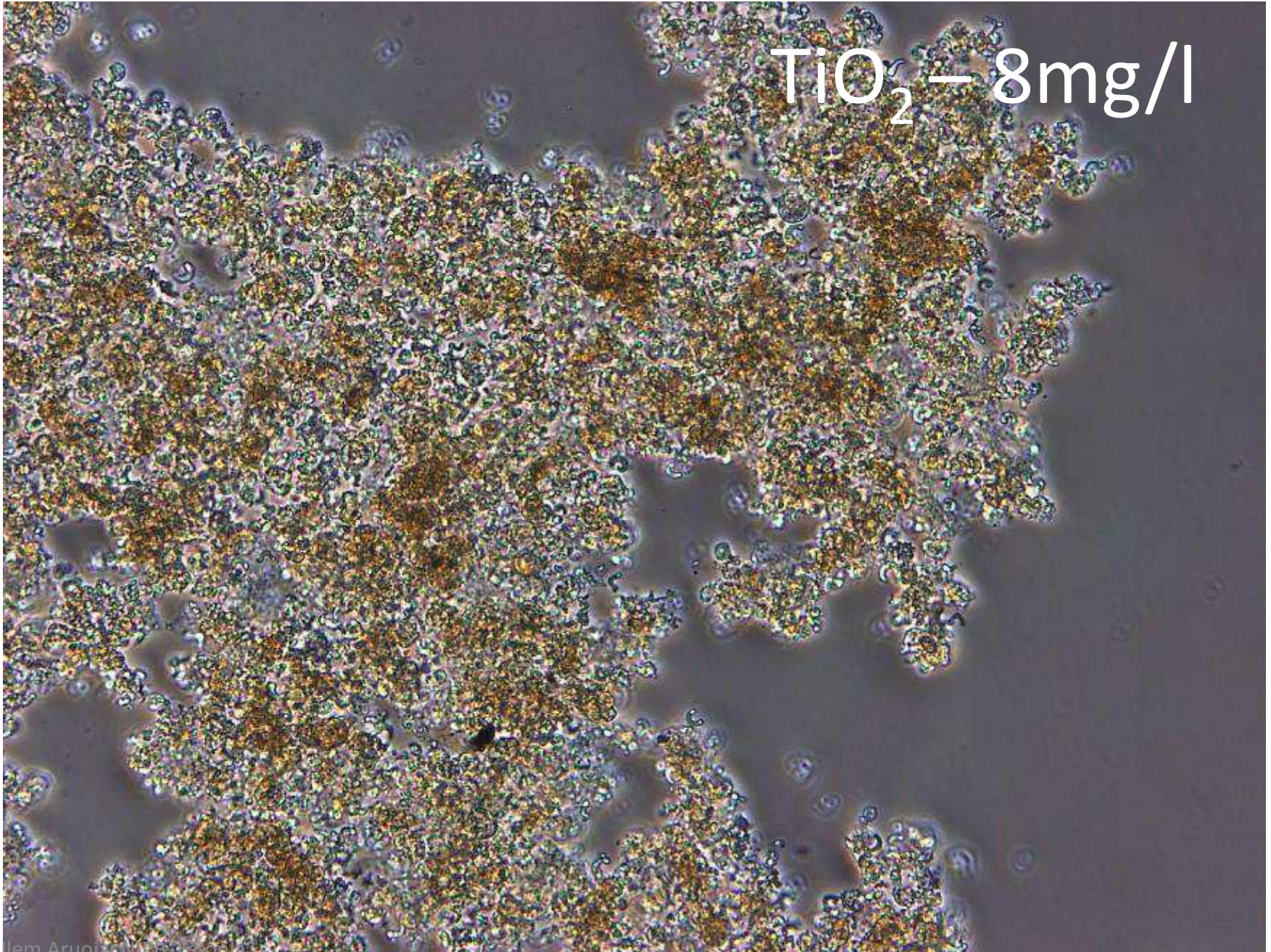


Fig. 2 Toxicity of 12 nanoparticles to alga *Pseudokirchneriella subcapitata*, protozoa *Tetrahymena thermophila* and bacterium *Vibrio fischeri*. EC₅₀ values were obtained from 72 h algal growth inhibition assay, 24 h *T. thermophila* viability assay and 30 min *V. fischeri* luminescence inhibition assay (Table S3†). Arrows indicate EC₅₀ values above 100 mg l⁻¹. Concentrations are nominal.

TiO₂ – 8mg/l

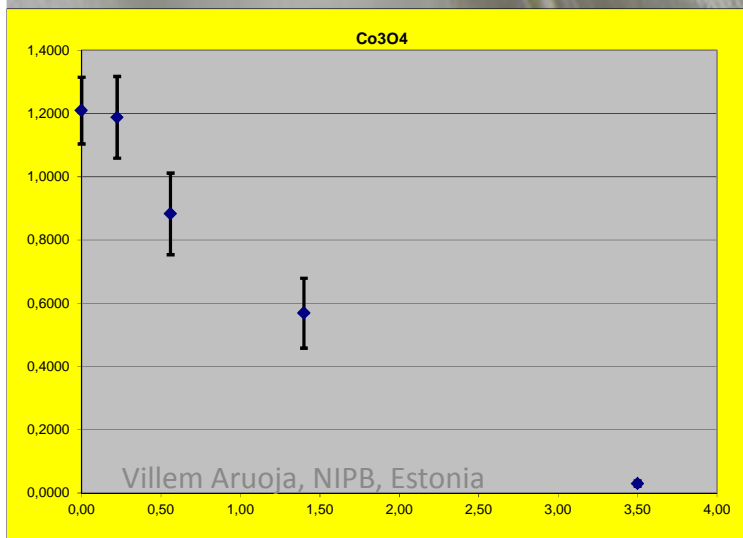


TiO_2 – 8mg/l

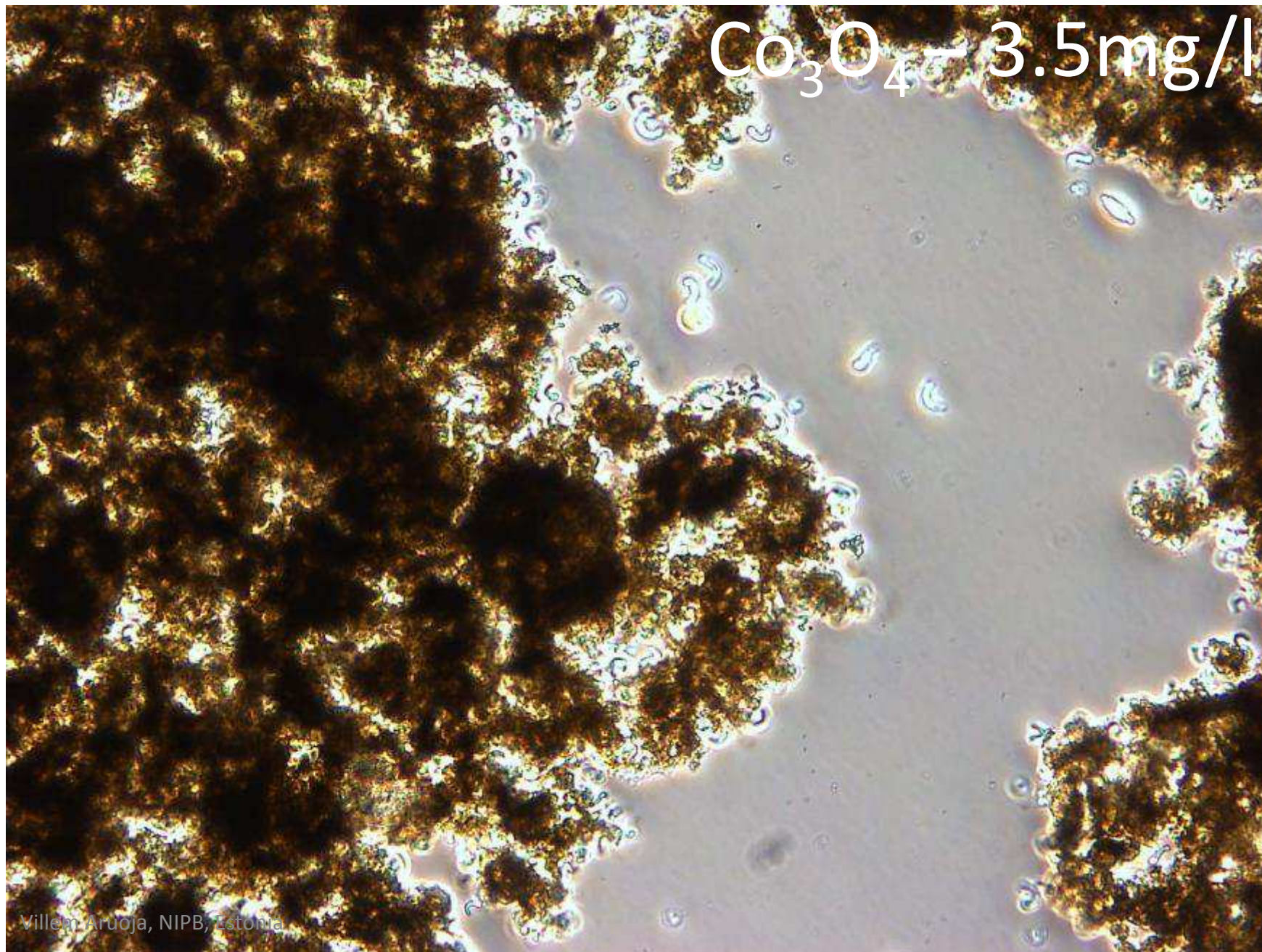


TiO_2 – 8mg/l

Co_3O_4 – 3.5mg/l

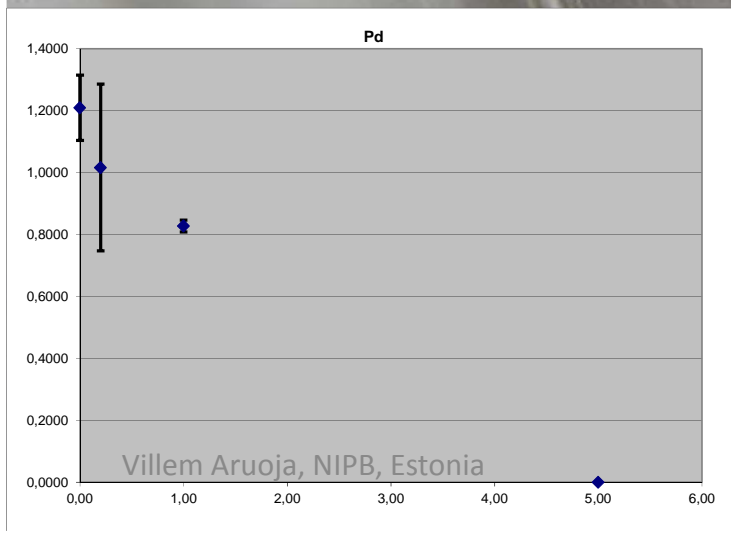


Co_3O_4 - 3.5mg/l

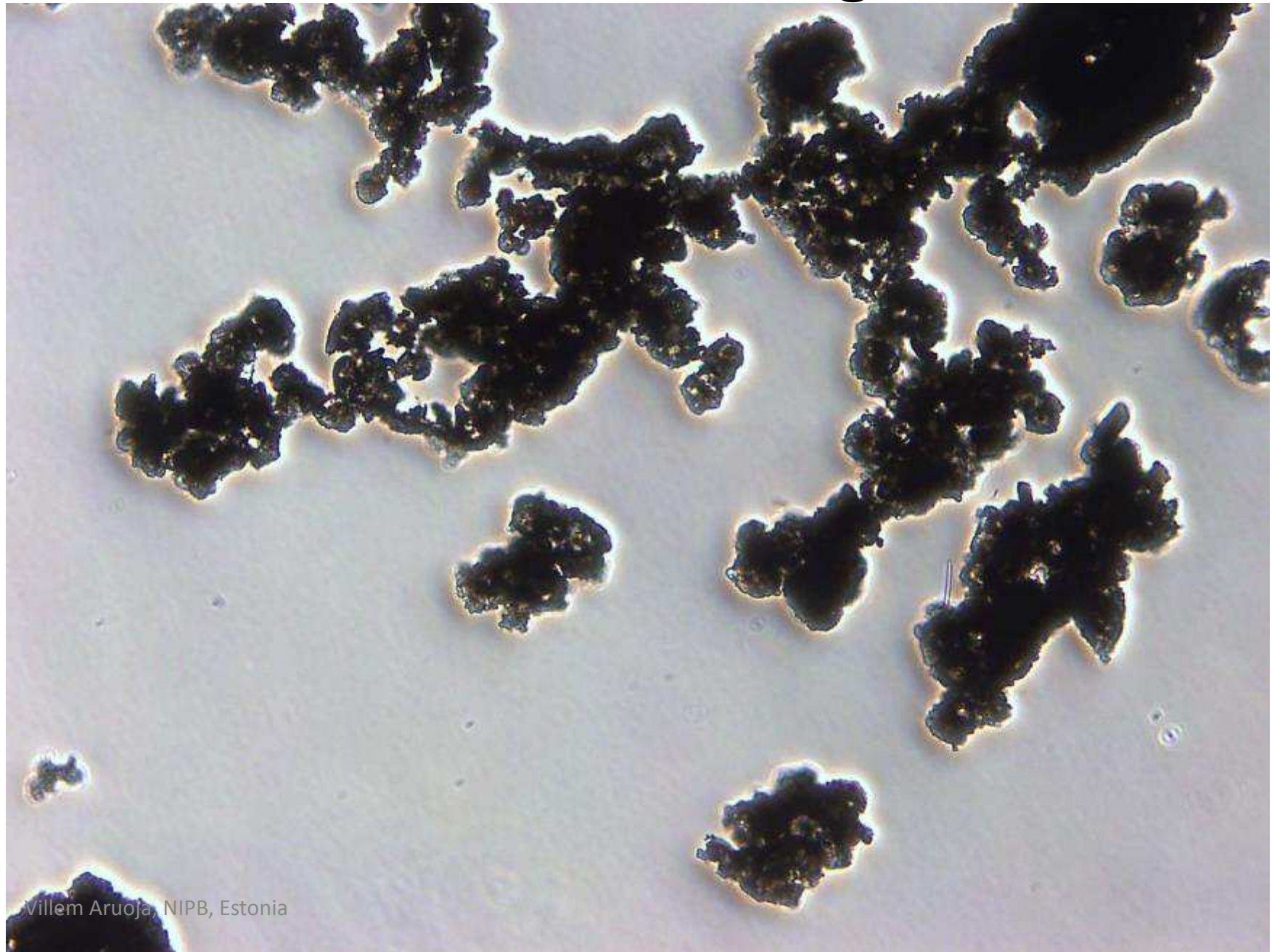


Co_3O_4 – 3.5mg/l

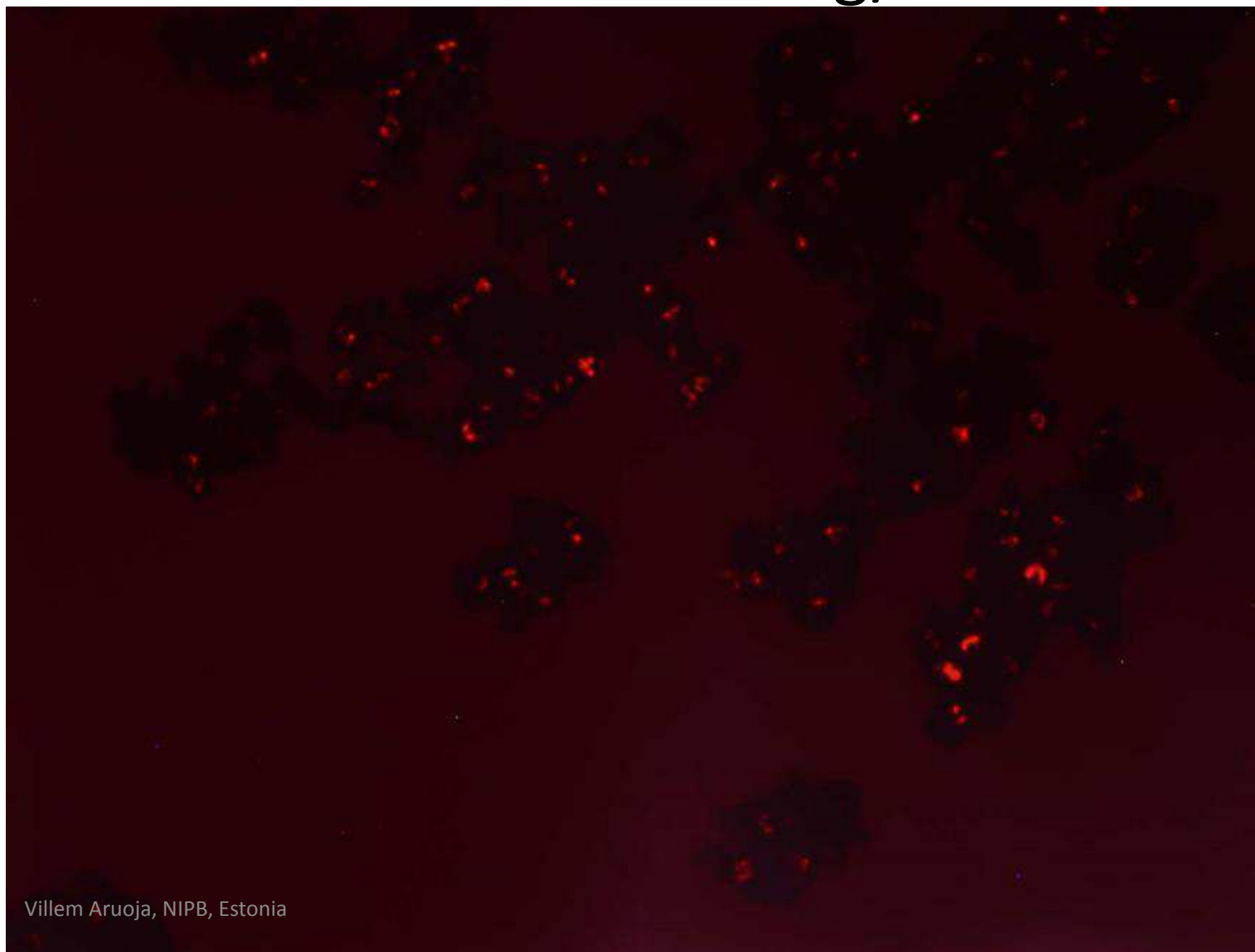
Palladium – 5mg/l



Palladium – 5mg/l

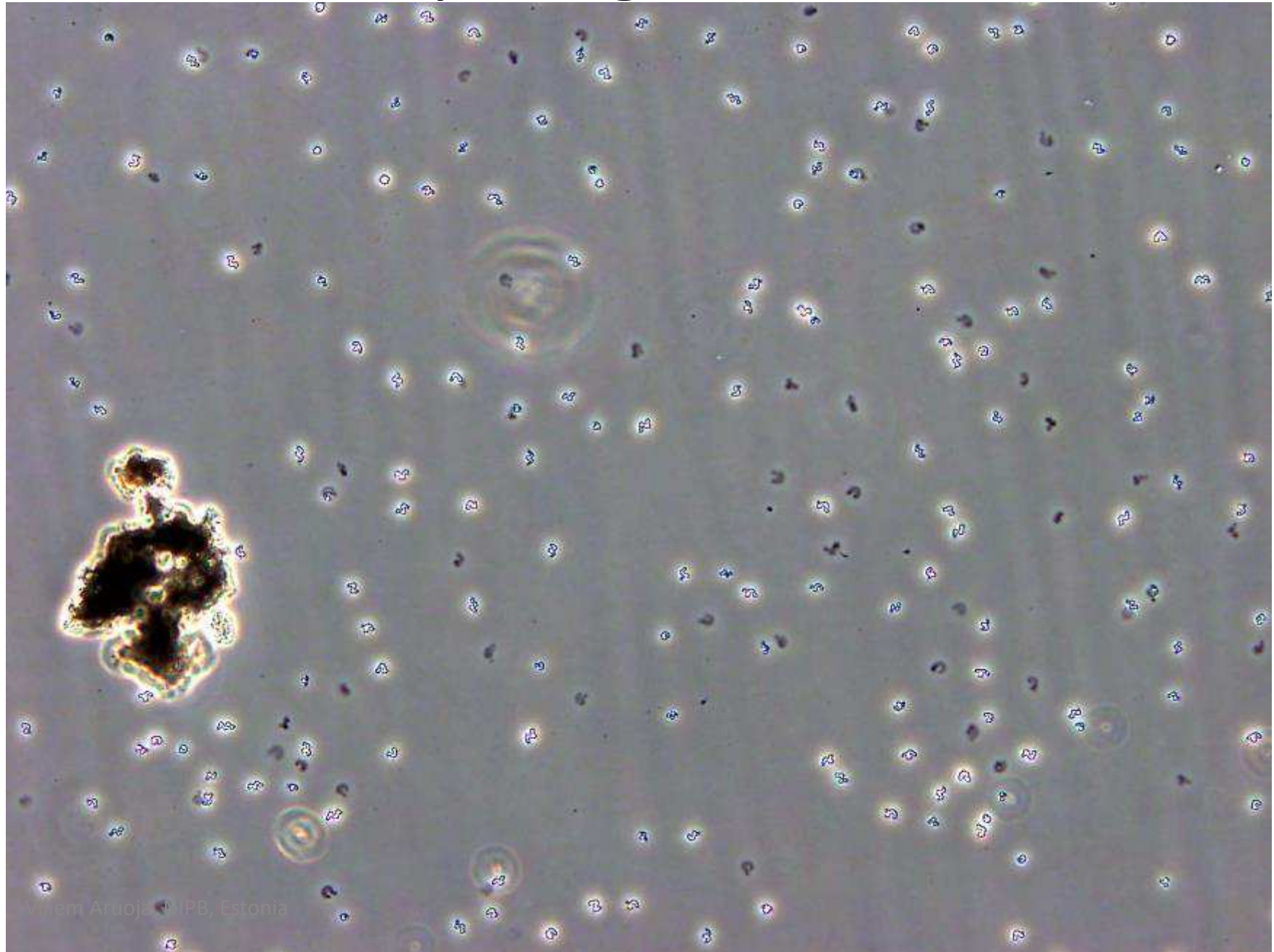


Palladium – 5mg/l



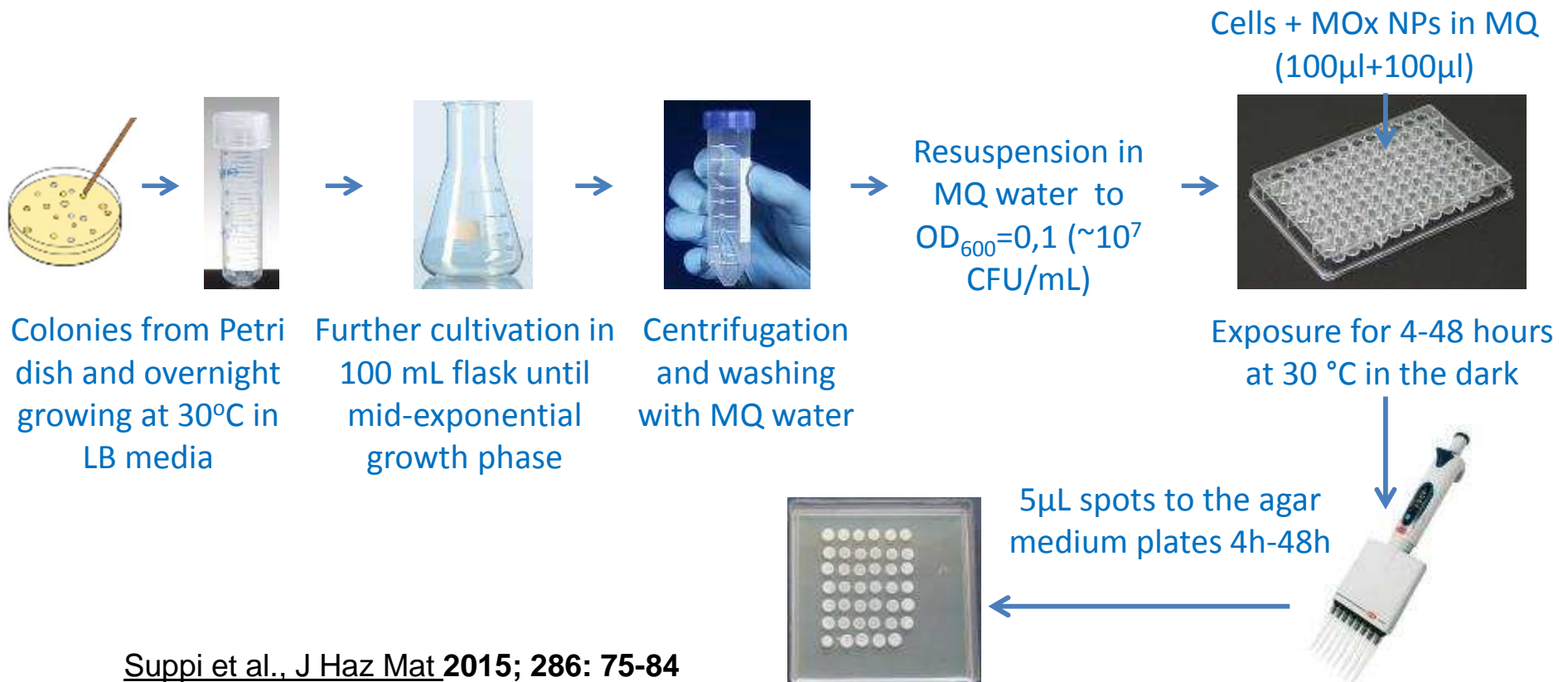
Villem Aruoja, NIPB, Estonia

Palladium – passage to clean medium



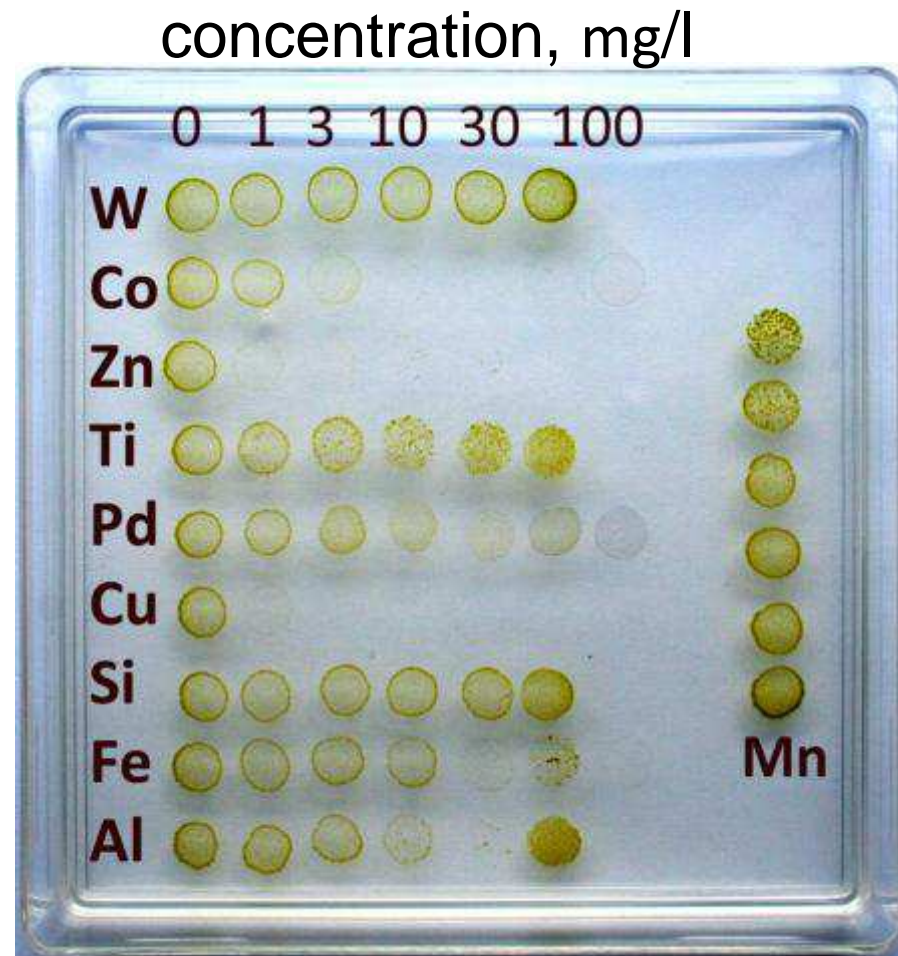
'Spot'-test

Inoculation	Bacterial growth to log phase	Washing with MQ water	Exposure to MOx NPs	4-48h spot
Bacterial colonies from Petri dish are transferred to 3ml liquid LB growth media and grown overnight in 14 mL round-bottom polypropylene culture tube at 30 °C on a shaker (200 rpm).	Dilution (1:50) of the overnight culture in 20 mL of fresh medium (in 100 mL culture flask) and further cultivation at 30 °C on a shaker at 200 rpm for 4-4.5 hours until bacteria reach mid-exponential growth phase ($OD_{600nm} \sim 0.6$).	The cells are harvested by centrifugation at 7000 rpm for 7 min at 20 °C in 50 mL centrifuge tubes. After that the cells were washed twice with MQ water and resuspended in MQ water to a density of $\sim 10^7$ CFU/mL ($OD_{600}=0.1$).	100 μ L of cells suspension and 100 μ L of MOx NPs suspension in MQ water were pipetted into the 96-well microplates. MQ water without test chemicals was inoculated with test strains in parallel and served as a control culture (not treated). The microplates were incubated for 4 hours or 24 hours at 30 °C in the dark, without shaking.	After 4 h or 24 h of exposure, 5 μ L of the cell culture from each well (treated or not treated) was pipetted ("spotted") onto the LB agar medium plates and incubated at 30 °C for 24 h. The growth of bacteria (formation of colonies) was evaluated visually on LB agar.

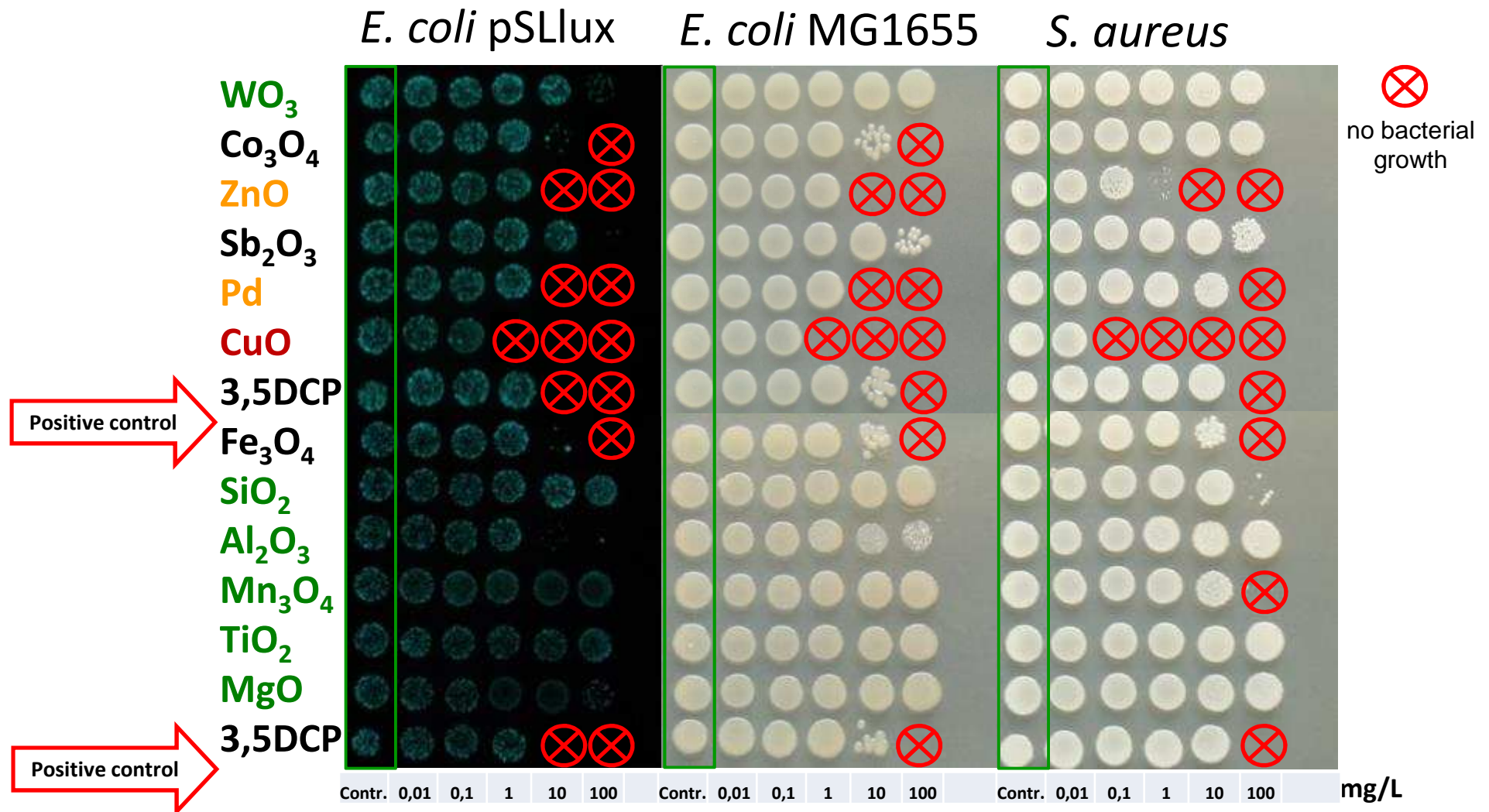


„Spot test“

Algal cells after
24h incubation
with NPs in
deionized water



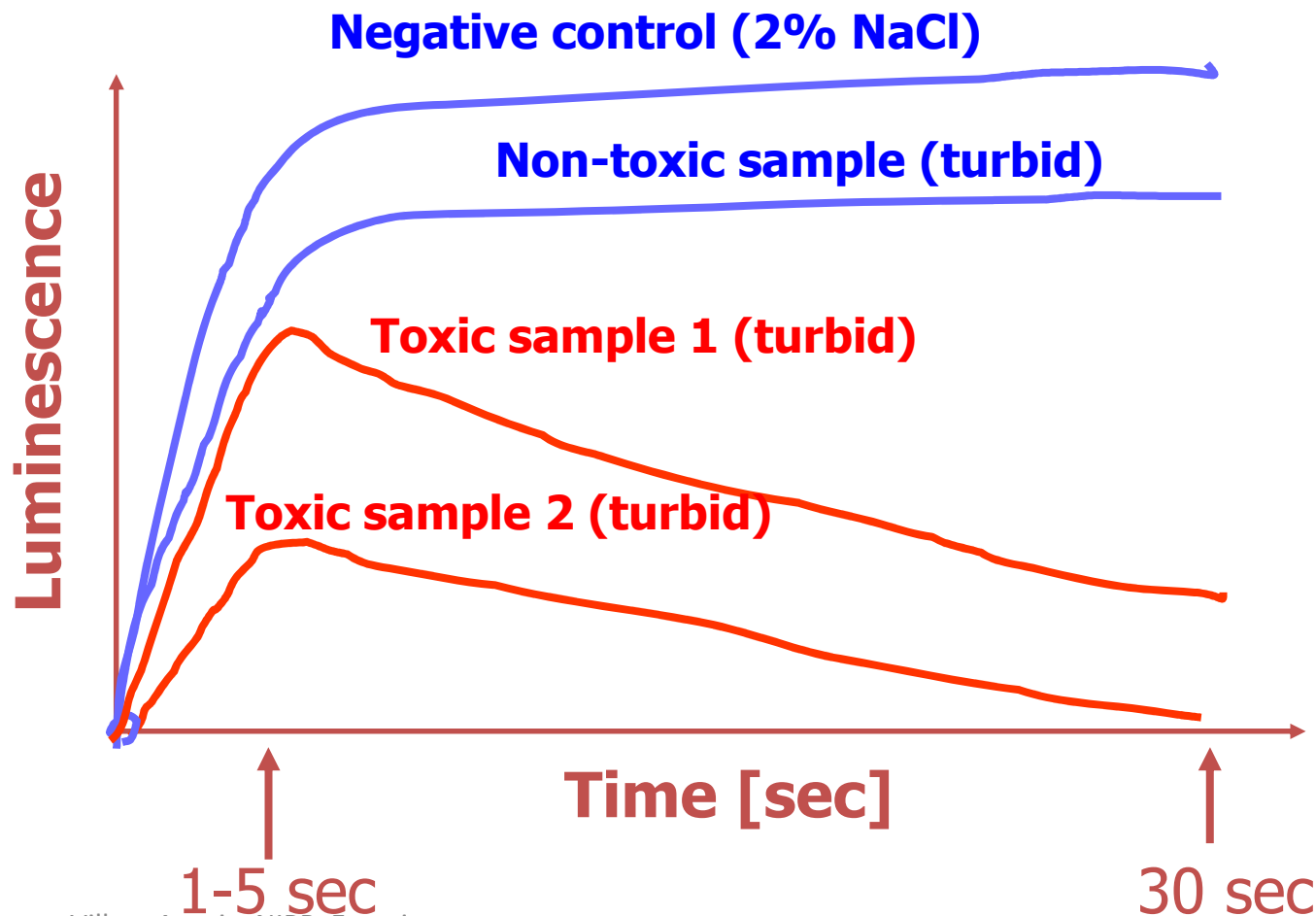
Results from 24-h 'spot'-test with bacteria



→ Co₃O₄, CuO, ZnO, Fe₃O₄ and Mn₃O₄ inhibited bacterial colony forming ability in the 'spot'-test at concentrations ≤ 100 mg/L.

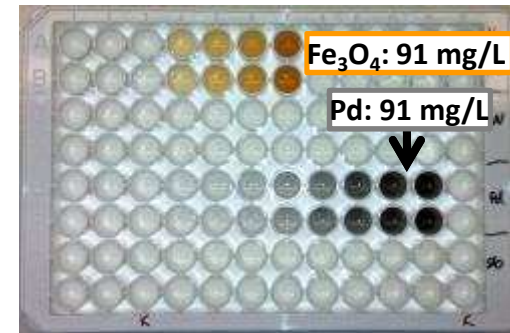
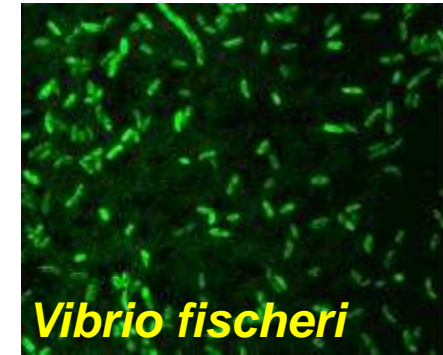
Villem Aruoja, NIBB, Estonia

Kinetic bioluminescence inhibition test with *Vibrio fischeri* (ISO 21338:2010)



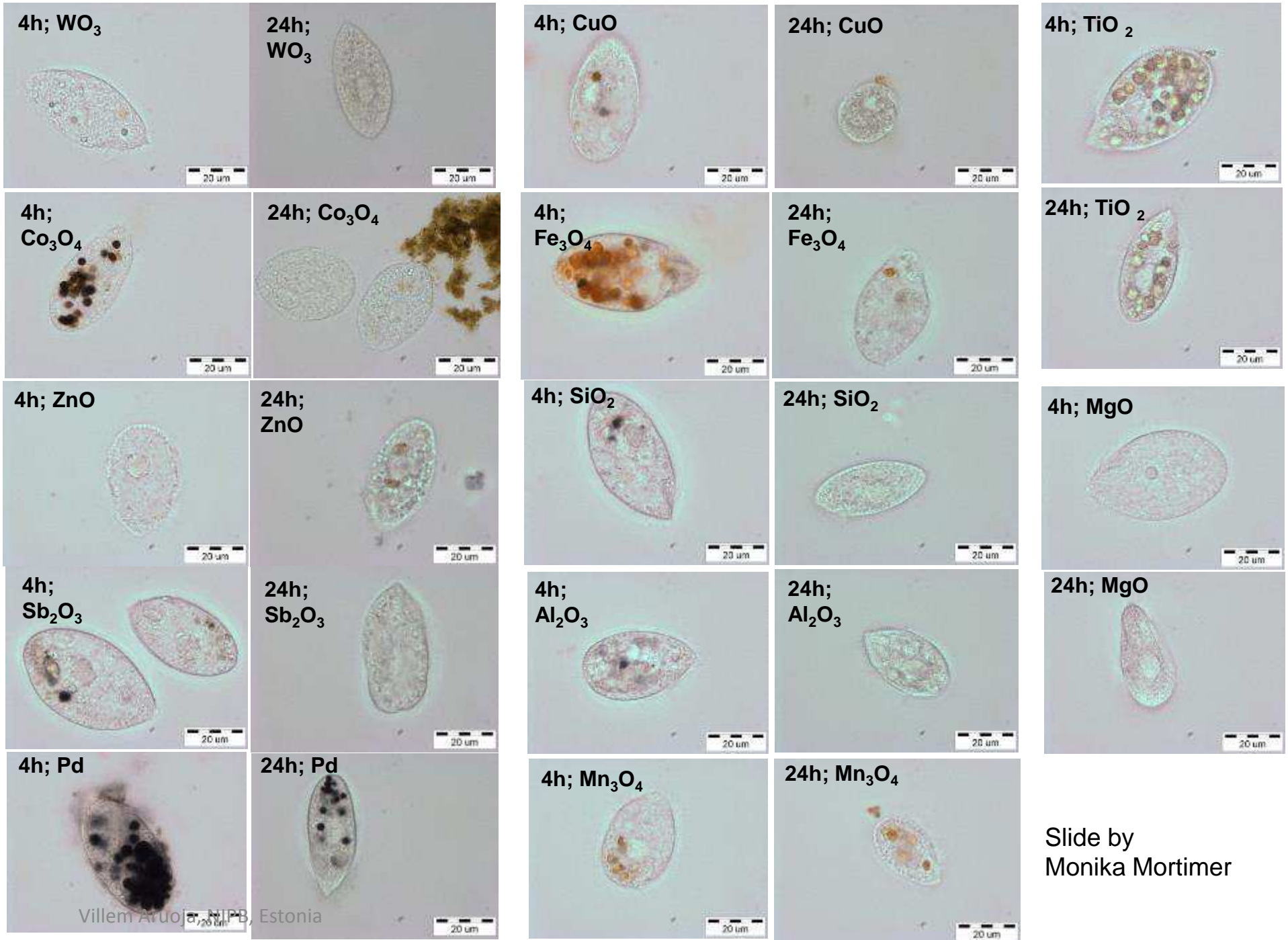
Villem Aruoja, NIPB, Estonia

Slide by Anne Kahru



96-well microplate





Villem Aruoja, NIPB, Estonia







Slide by
Monika Mortimer

Table 1. Toxicity of metal oxide and Pd nanoparticles to protozoa (*Tetrahymena thermophila*) and bacteria (*Vibrio fischeri*, *Escherichia coli*, *Staphylococcus aureus*). The presented toxicity values are based on nominal initial exposure concentrations used in testing.

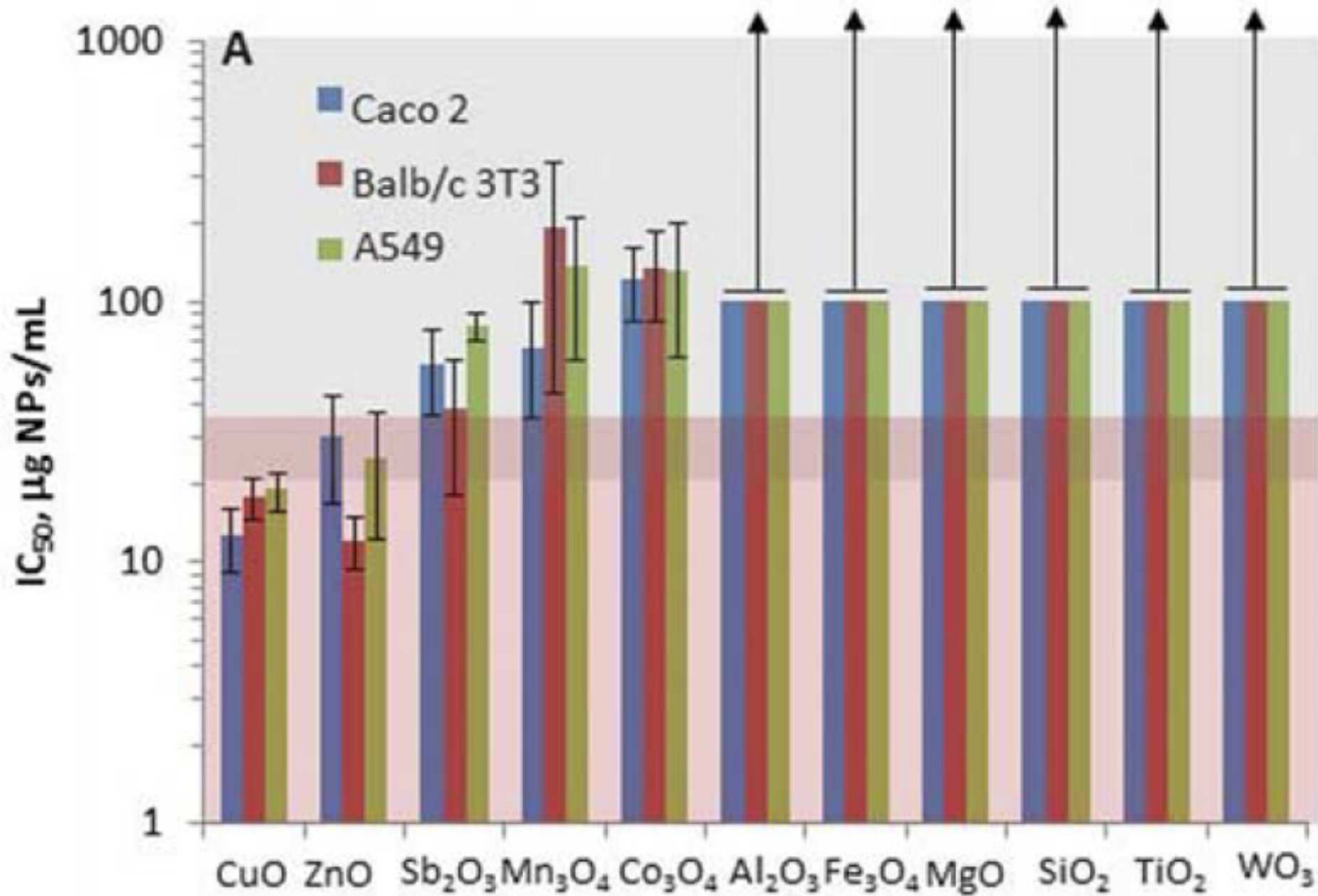
NPs	24-h EC50 ¹ <i>T. thermophila</i>	30-min EC50 ¹ <i>V. fischeri</i> (gram –)	24-h MBC ² <i>E. coli</i> pSLux (gram –)	24-h MBC ² <i>E. coli</i> MG1655 (gram –)	24-h MBC ² <i>S. aureus</i> (gram +)
	mg/L (compound)				
CuO	2.0	1.8	1	1	0.1
ZnO	1.8	11.6	10	10	10
Fe ₃ O ₄	20.1	>100	100	100	100
Mn ₃ O ₄	22.0	>100	>100	>100	100
Al ₂ O ₃	61.3	>100	100	>100	>100
TiO ₂	69.1	>100	>100	>100	>100
Pd	>100	56.3	10	10	100
Co ₃ O ₄	>100	>100	100	100	>100
Sb ₂ O ₃	>100	74.5	>100	>100	>100
WO ₃	>100	92.8	>100	>100	>100
SiO ₂	>100	>100	>100	>100	>100
MgO	>100	>100	>100	>100	>100

¹ **EC50** - half effective concentration; ² **MBC** - minimum bactericidal concentration. The lowest tested concentration that completely inhibited the visible growth of bacteria on the agarized test medium at 30°C in the dark after 24-h of incubation to MOx NPs. **Colour code:** ≤0.1 mg/L (■); >0.1–10 mg/L (■); 10–50 mg/L (■); 50–100 mg/L (■); =100 (■) mg/L; >100 mg/L (■).

Mechanism: solubility, ROS

Trophic level		Primary producer		Consumer	Degrader				
Cell type		Eukaryote		Eukaryote	Prokaryote				
Internalizes nanoparticles		No?		Yes	No?				
Nano-particle [¥]	Solubility (%) [§]	ROS (HPF) [§]	ROS (DCF) ⁺	Algal 72 h growth inhibition OECD 201	Protozoan 24 h viability	Bacterium 30 min. luminescence ISO 21338	Mechanism of toxicity	Classification [†]	
ZnO	56.1	-	-	< 1.0	1 ...2	10 ... 50	Zn ions	 Acute aquatic hazard	
Pd	<0.5	+	++	< 1.0	>100	10 ... 50	ROS		
CuO	5.14	+++*	-	< 1.0	1 ...2	1 ...2	Cu ions & ROS	 Acute aquatic hazard?	
Co₃O₄	1.25	+	+	1 ...2	>100	>100	ROS		
TiO₂	<0.83	+++	+	1 ...2	10 ... 50	>100	ROS	 Acute aquatic hazard?	
Mn₃O₄	11.1	-	+++	1 ...2	>100	>100	ROS		
Fe₃O₄	<1.38	+*	-	1 ...2	10 ... 50	>100	ROS	 Acute aquatic hazard?	
Al₂O₃	0.40	+	-	10 ... 50	>100	>100			
SiO₂	NA	-	-	10 ... 50	>100	>100		 Acute aquatic hazard?	
WO₃	63.2	-	-	10 ... 50	>100	50...100			
MgO	38.1	-	-	>100	>100	>100		 Acute aquatic hazard?	
Sb₂O₃	56.3	+	-	>100	>100	50...100			

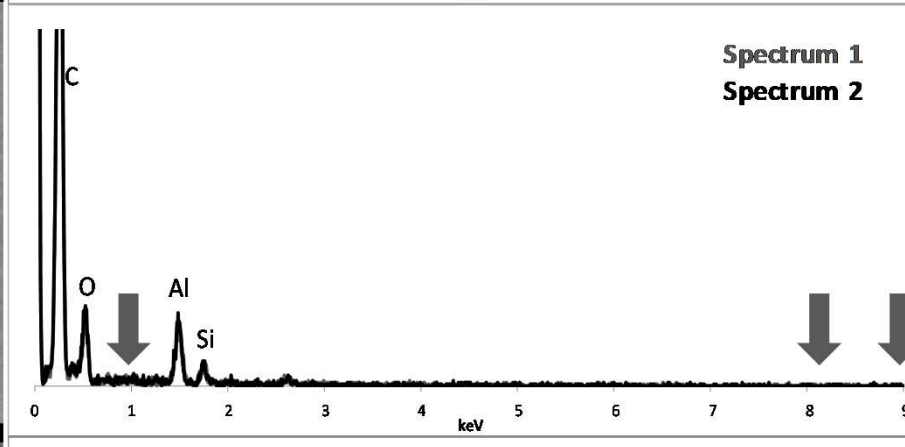
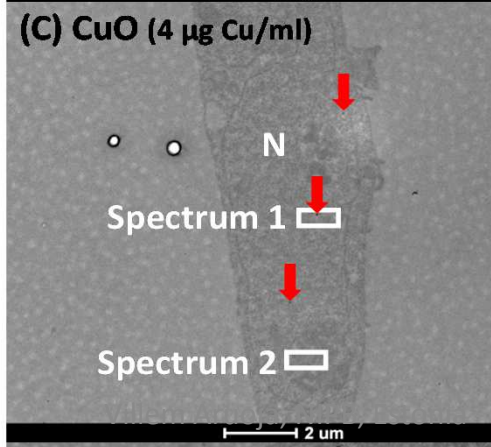
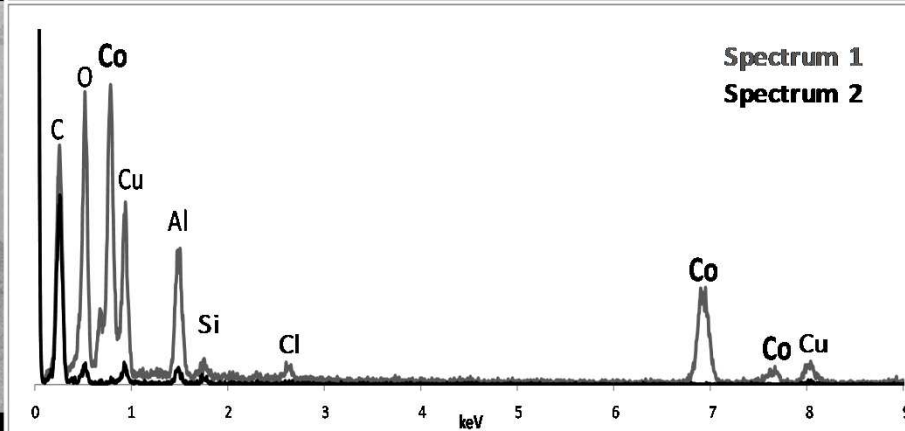
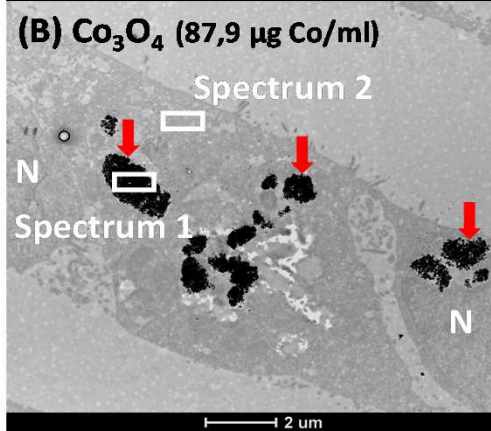
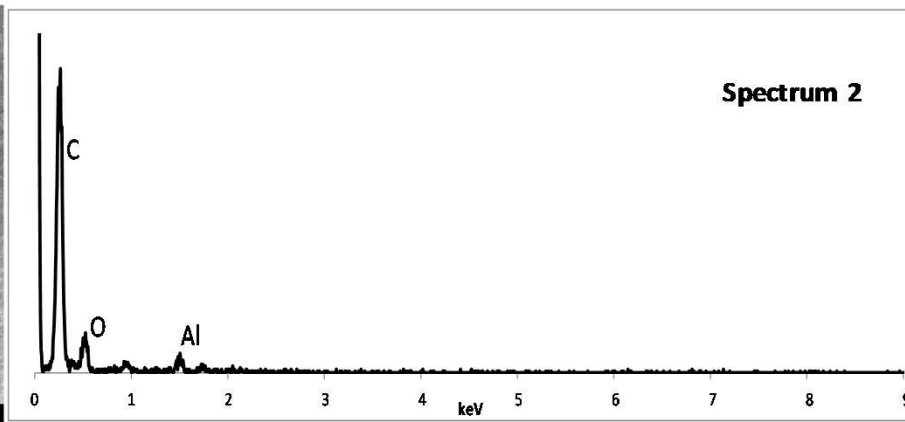
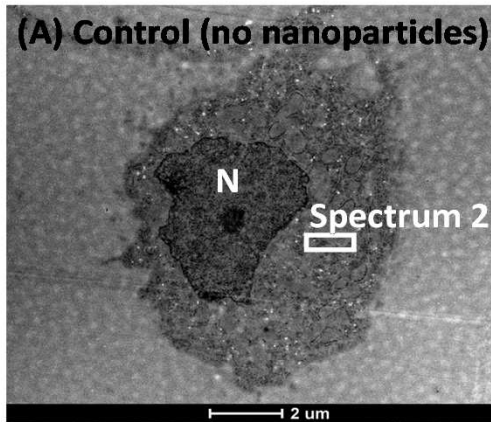
Cell cultures, EC₅₀



TEM images

EDX spectra

Endo- cytosis



Ivask et al. Current Topics in Medicinal Chemistry [2015, 15(18):1914-1929]



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Current Topics in Medicinal Chemistry, 2015, 15, 1-16

1

Toxicity of 11 Metal Oxide Nanoparticles to Three Mammalian Cell Types *In vitro*

Angela Ivask^{1,2,*}, Tiina Titma^{1,3}, Meeri Visnapuu^{1,4}, Heiki Vija¹, Aleksandr Käkinen¹,
Mariliis Sihtmäe¹, Suman Pokhrel⁵, Lutz Mädler⁵, Margit Heinlaan¹, Vambola Kisand⁴,
Ruth Shimmo³ and Anne Kahru¹



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Cite this: DOI: 10.1039/c5en00057b

Toxicity of 12 metal-based nanoparticles to algae, bacteria and protozoa†

Villem Aruoja,^{*a} Suman Pokhrel,^b Mariliis Sihtmäe,^a Monika Mortimer,^a
and Anne Kahru^a



Monte Carlo Modelling of Interaction Processes between Nanoparticles and Biomacromolecules of Variable Hydrophobicity

Fabrice Carnal

October 3rd, 2015



CompinNano, Ljubljana



**UNIVERSITÉ
DE GENÈVE**

FACULTÉ DES SCIENCES

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- pH
- Chain hydrophobicity
- Presence of nanoparticle

BSA protein

- Parametrization
- pH
- Nanoparticle surface charge density

IV. Conclusions and perspectives



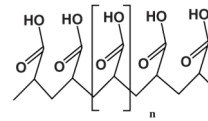
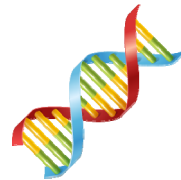
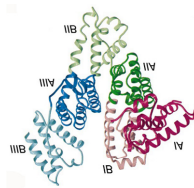
Introduction

I. Introduction

Polyelectrolytes

Charged polymers

- Functional groups are dissociated to charged monomers and counterions (weak polyelectrolyte charge varies in function of pH)

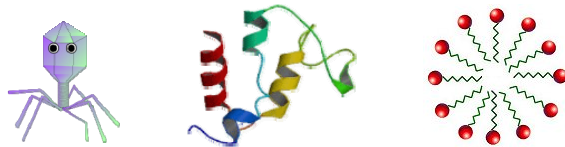


Nanoparticles

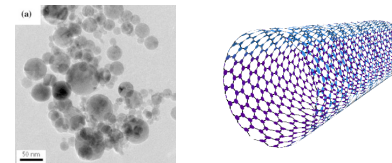
Objects constituted by tens or hundreds of atoms which have structure sizes comprised between 1 to 100 nm in at least one dimension

- Spherical, tubes, needle-like, etc.

Organic nanoparticles



Inorganic nanoparticles



High specific surface → High reactivity

I. Introduction

Electrostatic interactions in polyelectrolyte systems



Self-assembled complexes between polyelectrolytes and nanoparticles, dendrimers, flat surfaces, biomacromolecules, charged polymers

Biology

DNA packaging in eukaryote cells

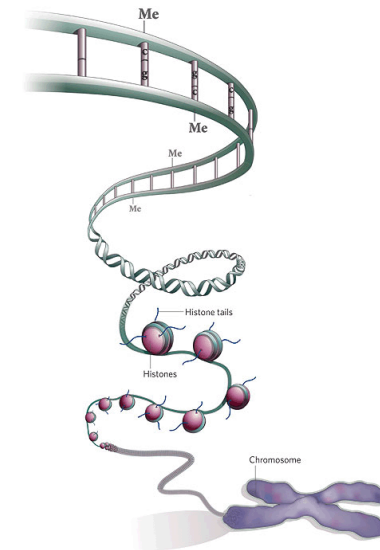
Biomedical

Contrast improvement in Magnetic Resonance Imaging

Cancer therapy by accumulation of active drug

Environment

Wastewater treatment

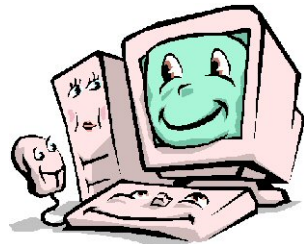


I. Introduction

Goal

How the chain hydrophobicity is playing a role on the conformational properties of biomacromolecules and on the formation of complexes with nanoparticles.

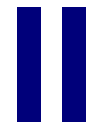
R_m	β
2	0
3	0.3
4	1
5	3



Systematic investigations



Numerical simulations



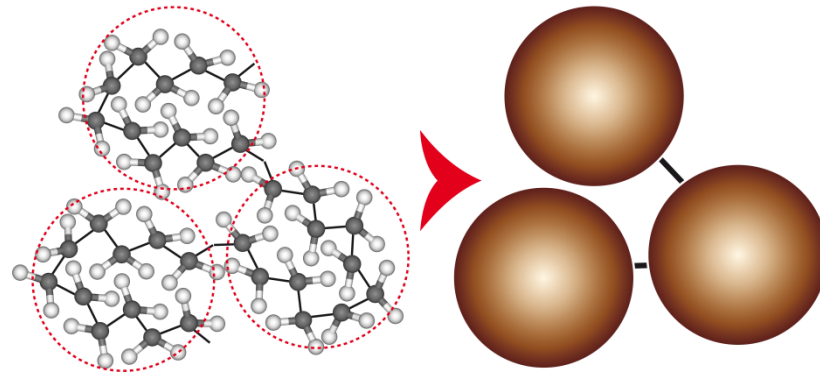
Simulation method and model

II. Simulation method and model

Polyelectrolytes

- Several thousands of atoms !
- CPU consuming

**Coarse-grained
models**



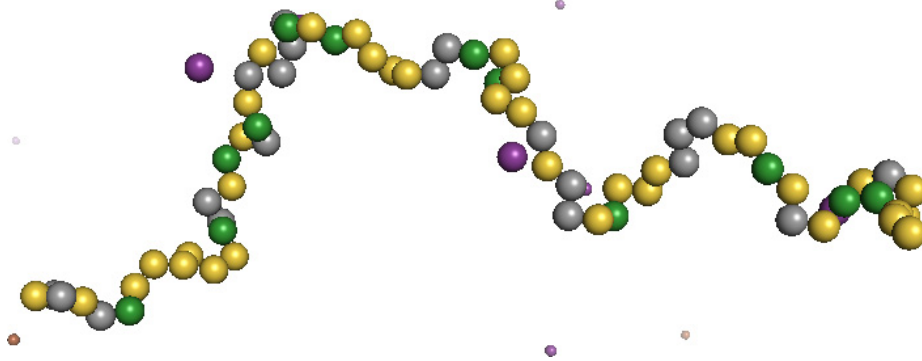
Atomistic details (bond length, vibrations, etc.) are omitted



Group of atoms → effective monomer

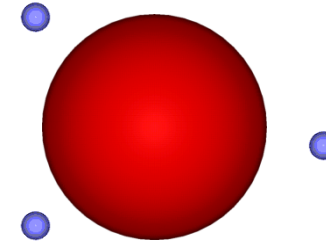
II. Simulation method and model

Biomacromolecules



- Neutral monomers
- Negatively charged monomers
- Positively charged monomers

Nanoparticles



- Nanoparticle counterions
- Chain counterions

- Each object is represented by a hard sphere
- The solvent (water) treated implicitly as a dielectric medium
- Monomer charge varies depending on the solution pH

II. Simulation method and model

Potentials

1) Electrostatic

Each charged objects interact via a full Coulomb electrostatic potential and excluded volume potential

$$U_{el}(r_{ij}) = \begin{cases} \infty, & r_{ij} < R_i + R_j \\ \frac{z_i z_j e^2}{4\pi\epsilon_r \epsilon_0} \frac{1}{r_{ij}}, & r_{ij} \geq R_i + R_j \end{cases}$$

e Elementary charge

ϵ_0 Permittivity of the vacuum

ϵ_r Water dielectric constant

r_{ij} Distance between two particles

R Particle radius

z Particle charge

2) Lennard-Jones

Lennard-Jones potential is used to take into account hydrophobic interactions between monomers

$$U_{vdW}(r_{ij}) = vdW \left[\left(\frac{r_0}{r_{ij}} \right)^{12} - 2 \left(\frac{r_0}{r_{ij}} \right)^6 \right]$$

r_0 Usually $R_i + R_j$

vdW Minimum depth of the potential curve located at a distance r_0

II. Simulation method and model

Minimum energy investigation

Monte Carlo Metropolis

- Random conformation

→ System energy calculation $E_{initial} = \sum_{ij} U_{ij}$

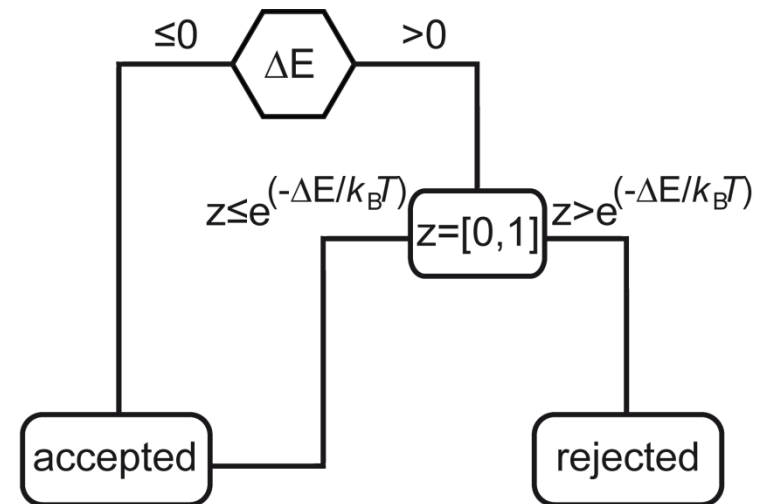
- Elementary Movements

→ System energy calculation E_{final}

- Energy difference

$$\Delta E = E_{final} - E_{initial}$$

- Metropolis test



Several thousands successive trials to achieve equilibrium state



Recording of observables (macroscopic properties) → **average values**

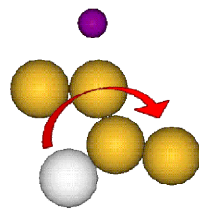
II. Simulation method and model

Elementary movements

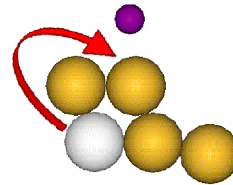
Biomacromolecules

Counterions

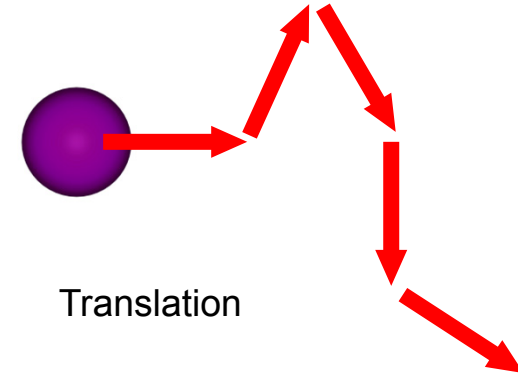
Internal movements



End-bond

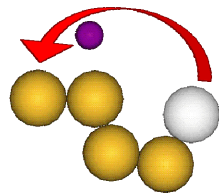


Kink-jump

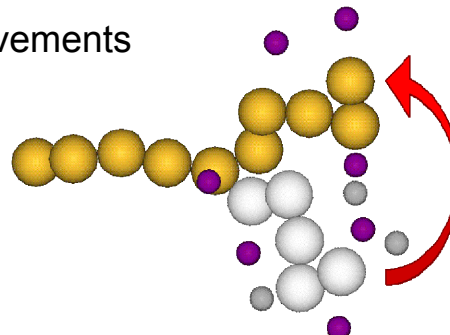


Translation

Global movements



Reptation



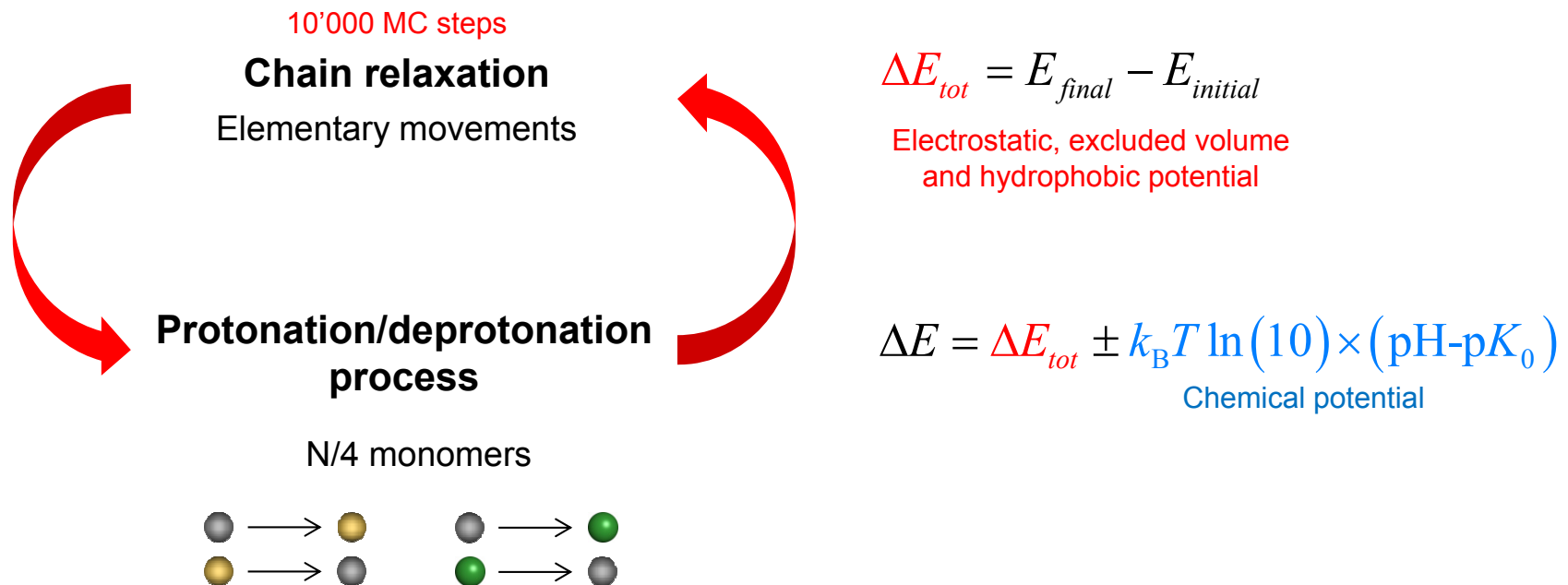
'Partially clothed' Pivot

II. Simulation method and model

pH dependency of biomacromolecules



Protonation/deprotonation cycle



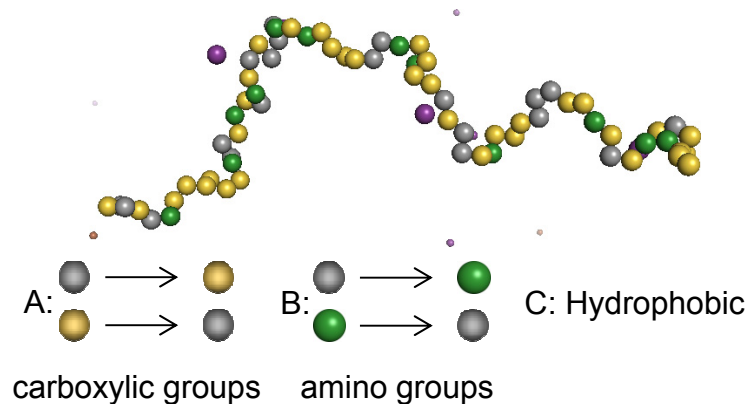
Several million MC steps to obtain a reasonable sampling of low energy conformations

II. Simulation method and model

2 models

Simplified protein structure

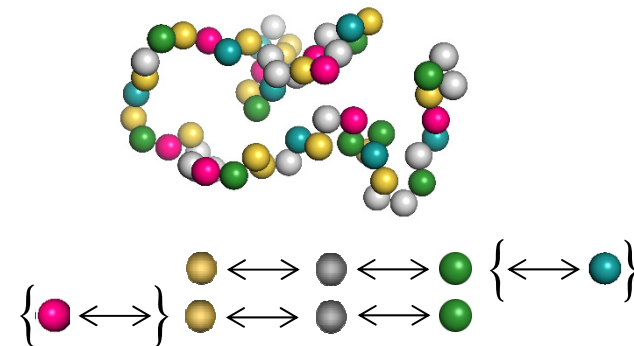
- 0 Neutral monomers
- 1 Negatively charged monomers
- 1 Positively charged monomers



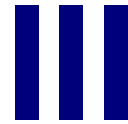
- 100 monomers
- sequence of monomers randomly determined in the beginning of the simulation
- $pK_A = 2.17$ / $pK_B = 9.53$

BSA protein

- 0 Neutral monomers
- 2 -1 Negatively charged monomers
- 2 1 Positively charged monomers



- 583 amino acids
- sequence of amino acids known (x-ray structure)
- each amino acid can be neutral, positively and negatively charged
- pK_A , pK_B and pK_C (if present) different for each amino acid

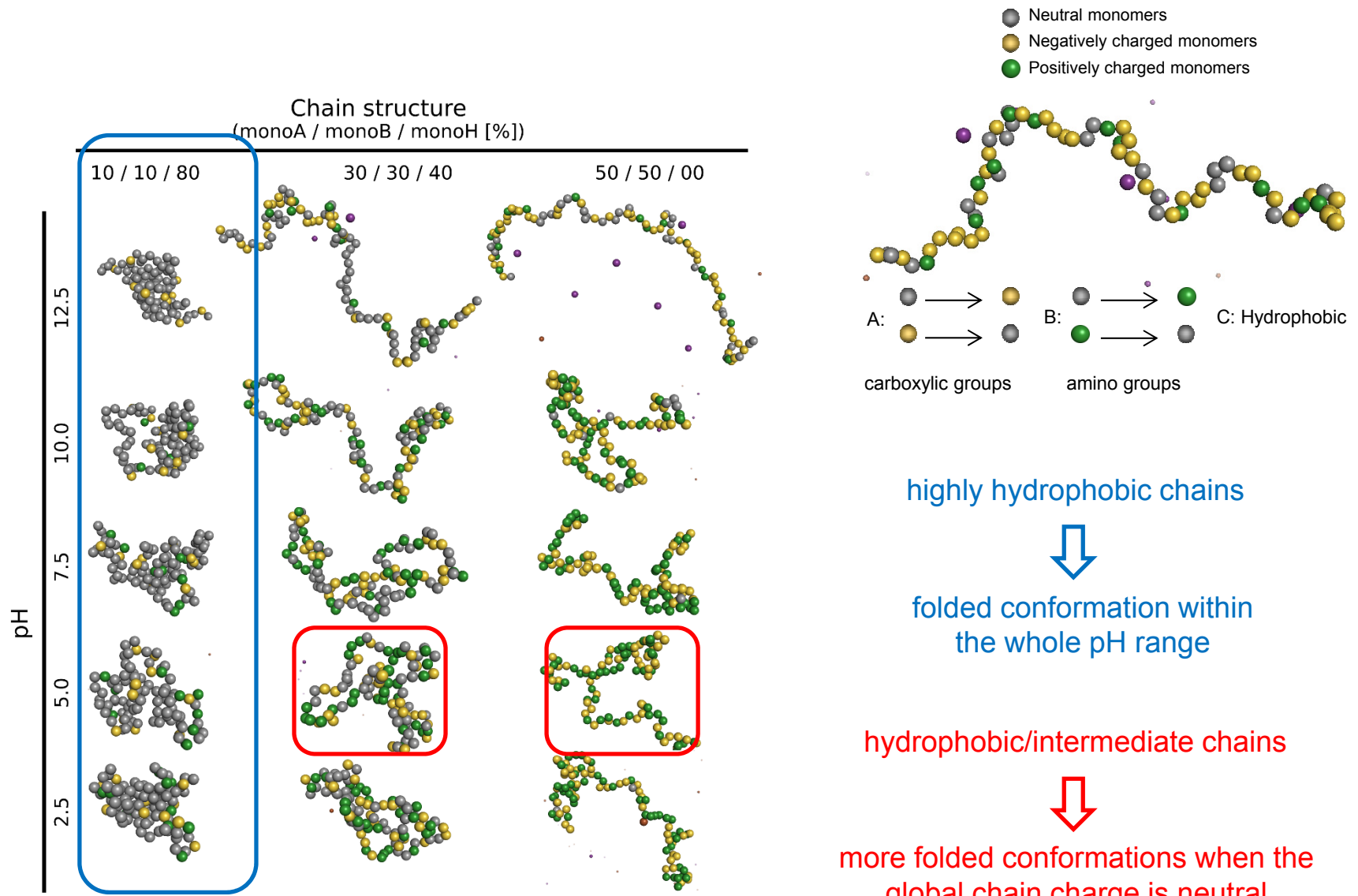


Results

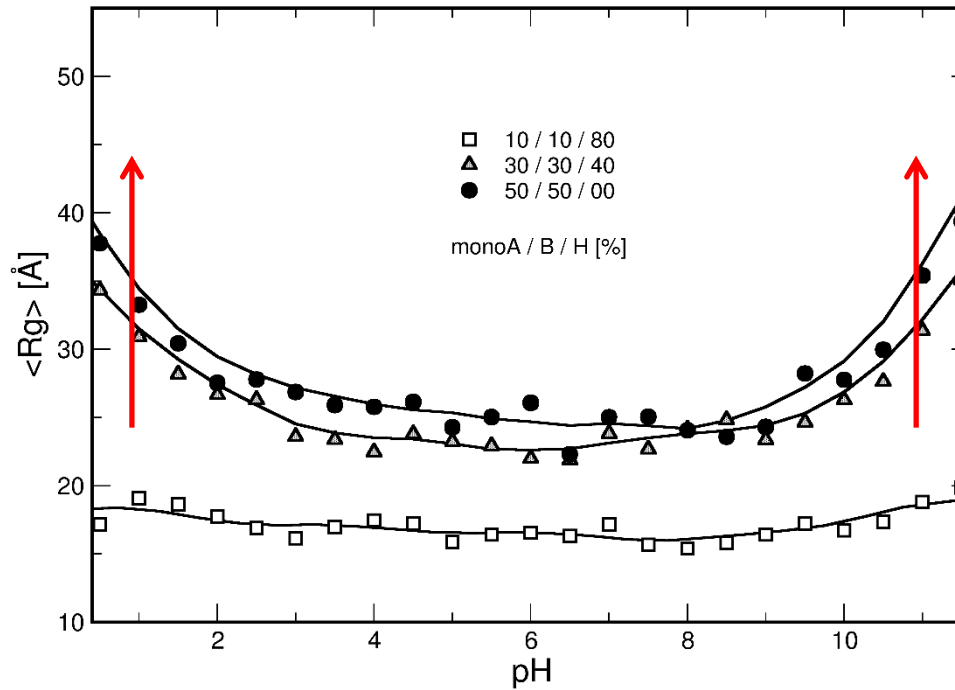
Simplified protein structure

- pH
- Chain hydrophobicity
- Presence of nanoparticle

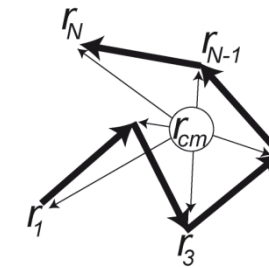
III. Simplified protein structure



III. Simplified protein structure



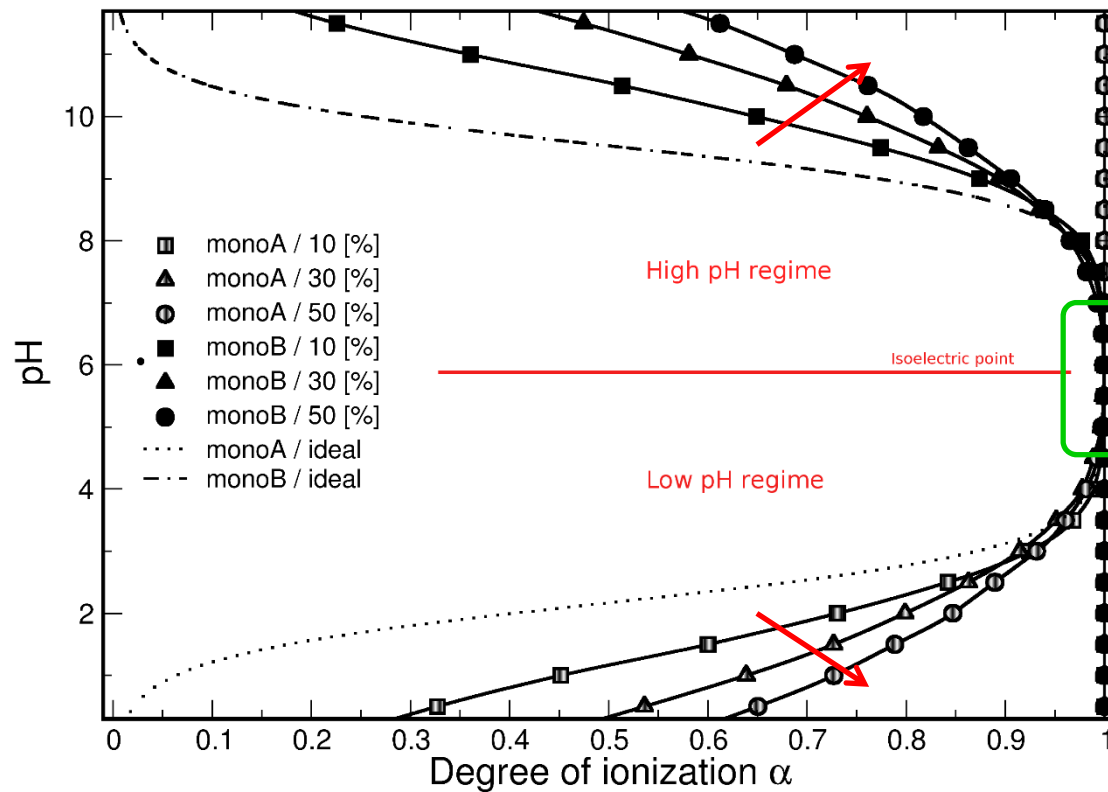
Square radius of gyration



$$R_g^2 = \frac{1}{N} \sum_{i=1}^N (r_i - r_{cm})^2$$

- Extended conformations at low and high pH in the case of hydrophilic and intermediate chains
- Strong repulsive monomer-monomer electrostatic interactions favorise less compact conformations

III. Simplified protein structure



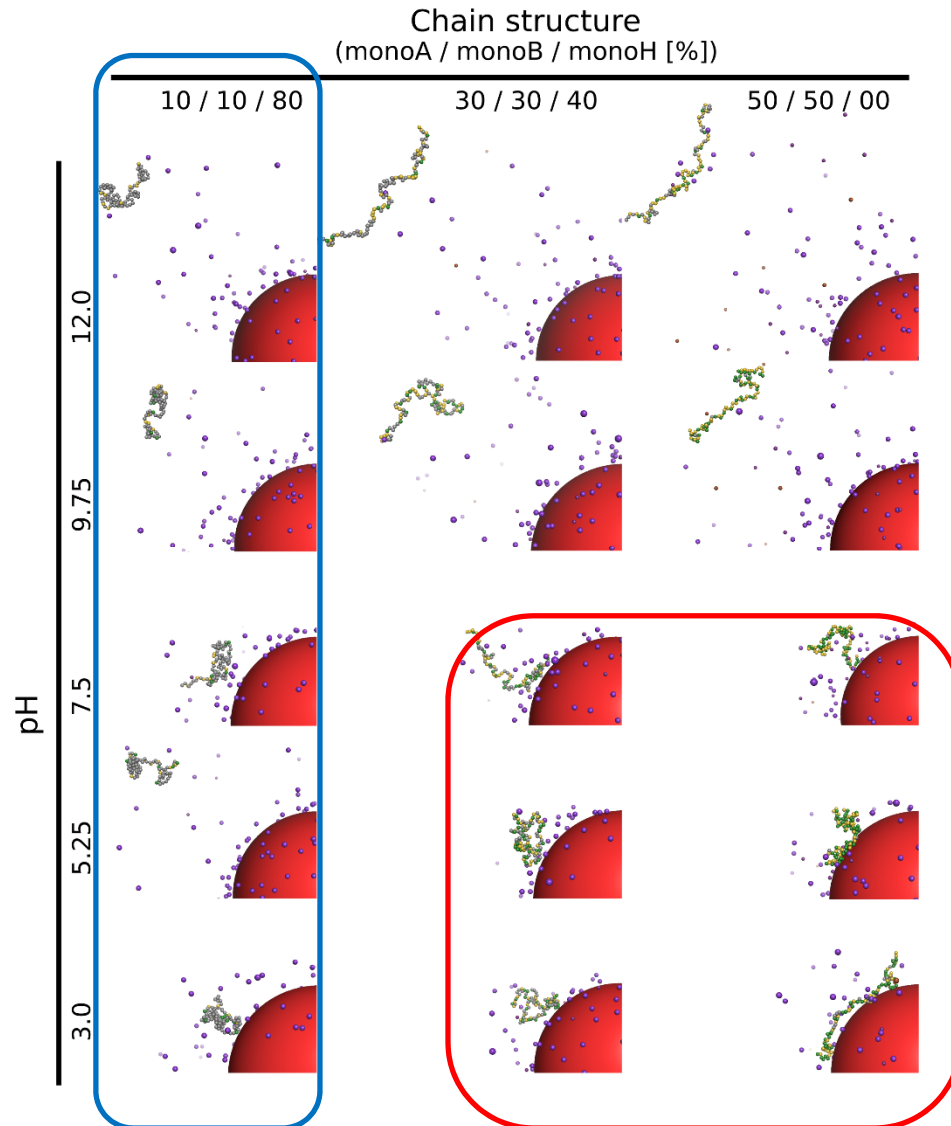
$pK_0^A = 2.17$ (carboxylic groups)

$pK_0^B = 9.53$ (amino groups)

Chain globally neutral!

- Charging process of carboxylic/amino groups promoted by charged amino/carboxylic groups
- Attractive electrostatic interactions weaker for hydrophobic chains resulting in a less efficient charging process

III. Simplified protein structure



$R_{NP} = 100\text{\AA}$
 $\sigma_{NP} = -39.9\text{ mC/m}^2$

highly hydrophobic chains



no elongated conformation
no formation of complexes

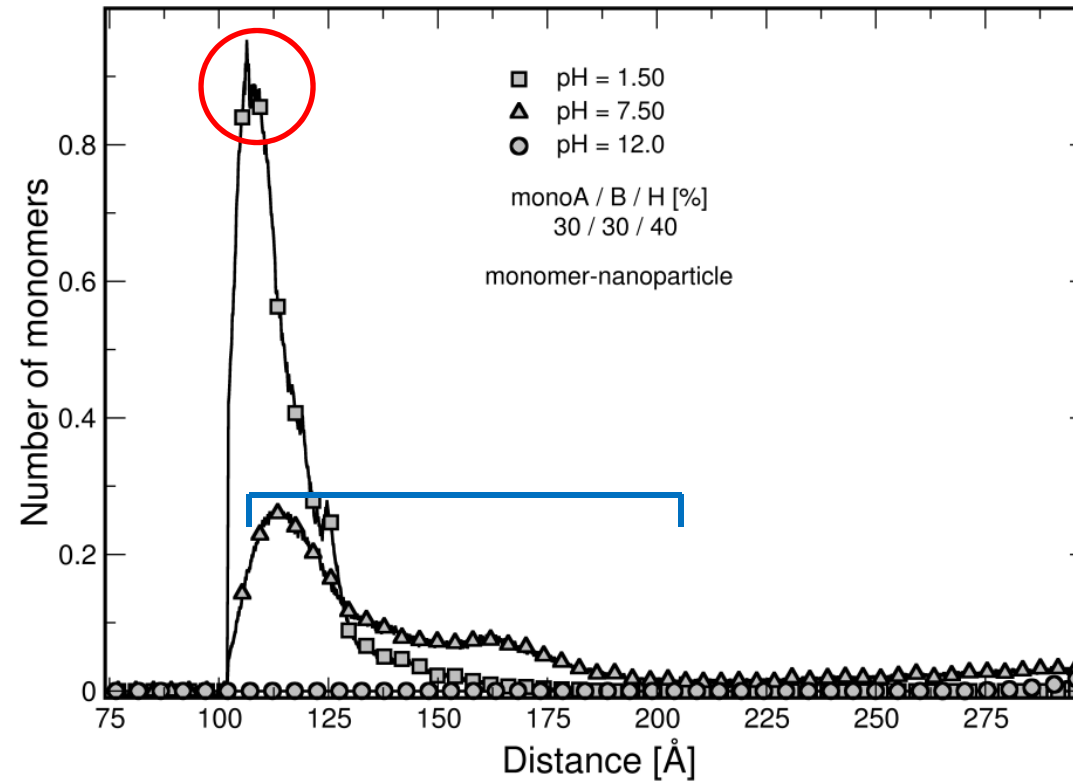
intermediate and hydrophilic chains



stronger electrostatic interactions with
the NP → adsorption at the surface

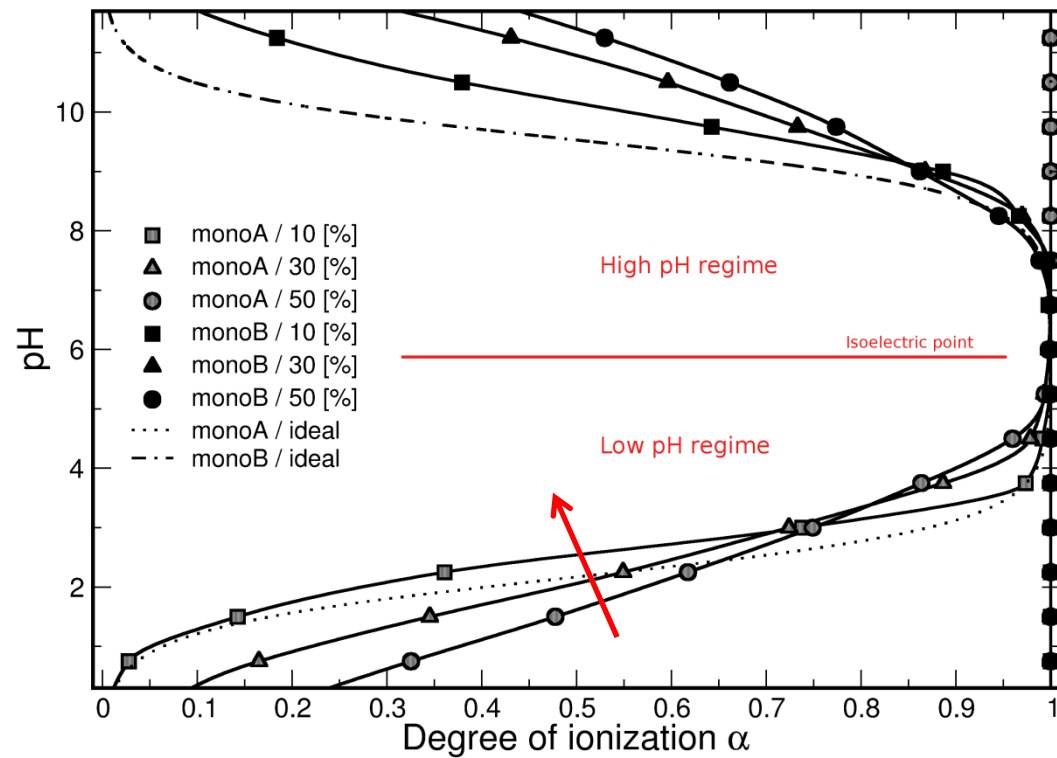
both amino and carboxylic groups
charged at physiological pH resulting in
partial chain adsorption

III. Simplified protein structure

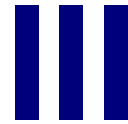


- Strong adsorption at low pH → **high peak**
- At physiological pH, local adsorption resulting in a **larger monomer distribution**

III. Simplified protein structure



- Lost of symmetry at low pH → **intersection** with the ideal curve
- Modification of the apparent pKa hence **promoting** the protonation of carboxylic groups

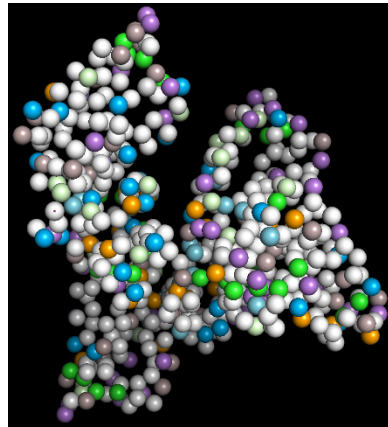


Results

BSA protein

- Parametrization
- pH
- Nanoparticle surface charge density

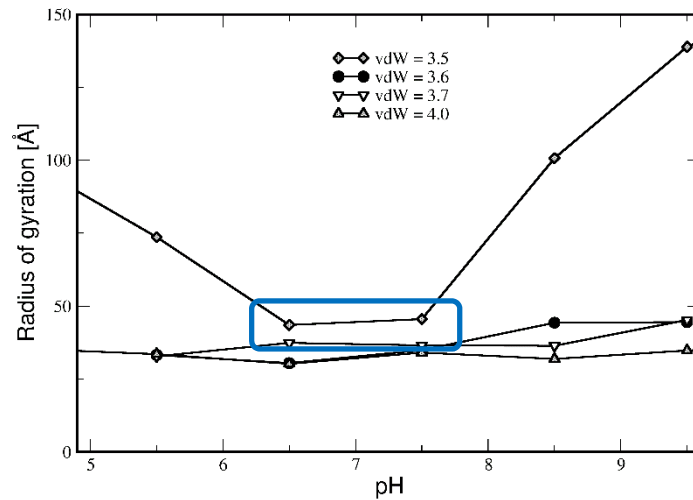
III. BSA protein



BSA x-ray structure

Parameters

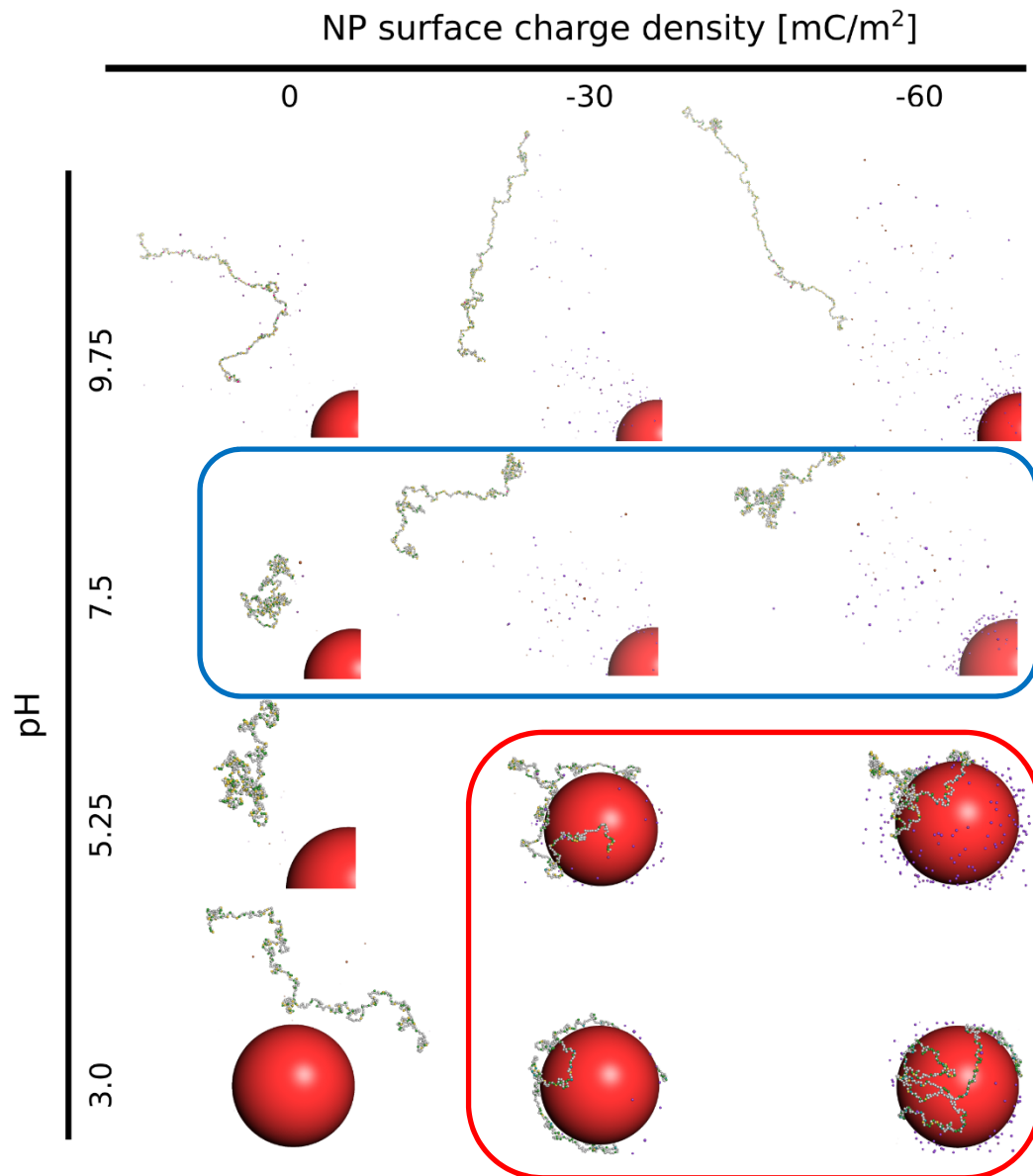
- length and contour size of the protein
- amino acid distribution
- pK_a , pK_b and pK_c values



$$U_{\text{vdW}}(r_{ij}) = \text{vdW} \left[\left(\frac{r_0}{r_{ij}} \right)^{12} - 2 \left(\frac{r_0}{r_{ij}} \right)^6 \right]$$

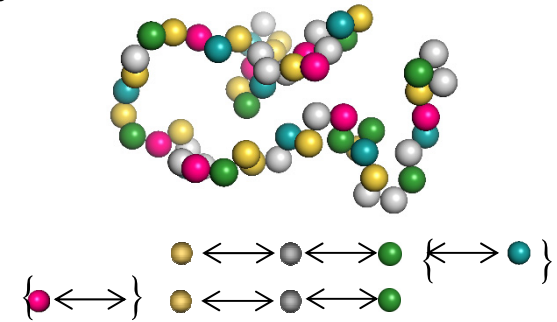
- hydrophobic interactions between amino acids not too strong to observe conformational changes (denaturation)
→ $\text{vdW} = 3.5 K_B T$
- hydrophobic amino acids: Ala, Met, Leu, Val, Ile, Phe

III. BSA protein



$N_{AA} = 583$
 $R_{AA} = 2\text{\AA}$

- Neutral monomers
- Negatively charged monomers
- Positively charged monomers



physiological pH



more folded conformations
 and no adsorption with only
 electrostatic interactions

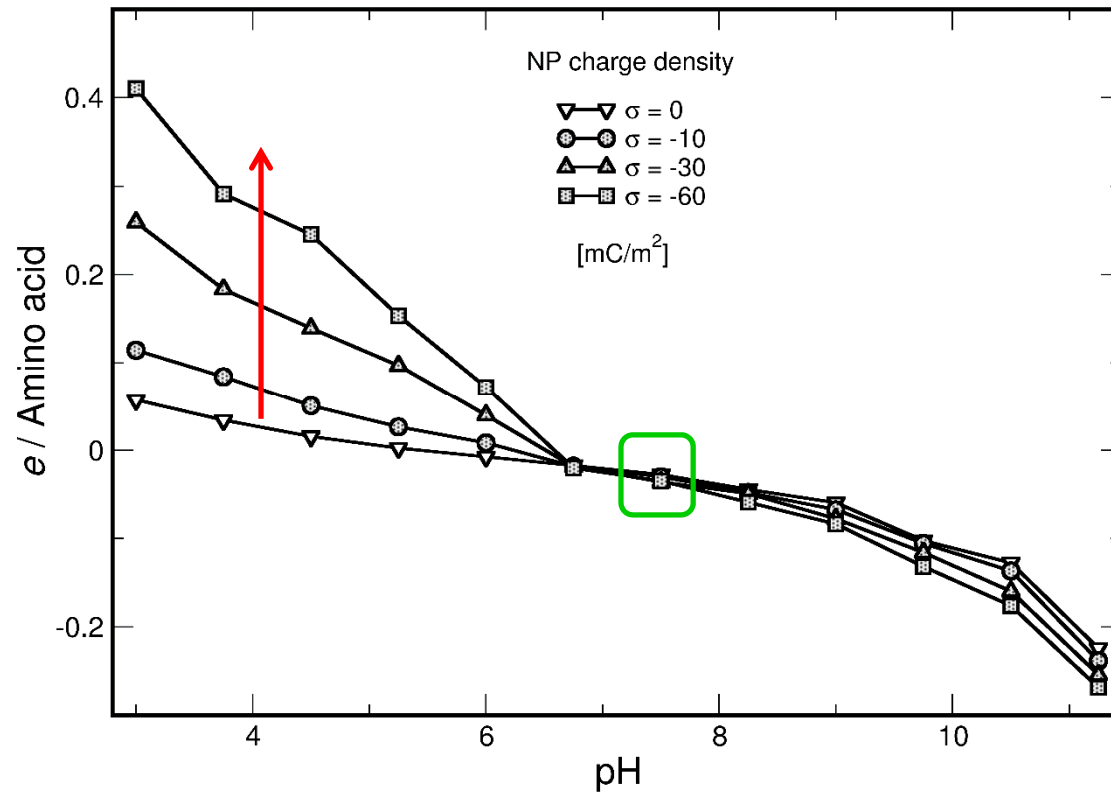
protein positively charged at low pH



adsorption at the surface of the
 nanoparticle

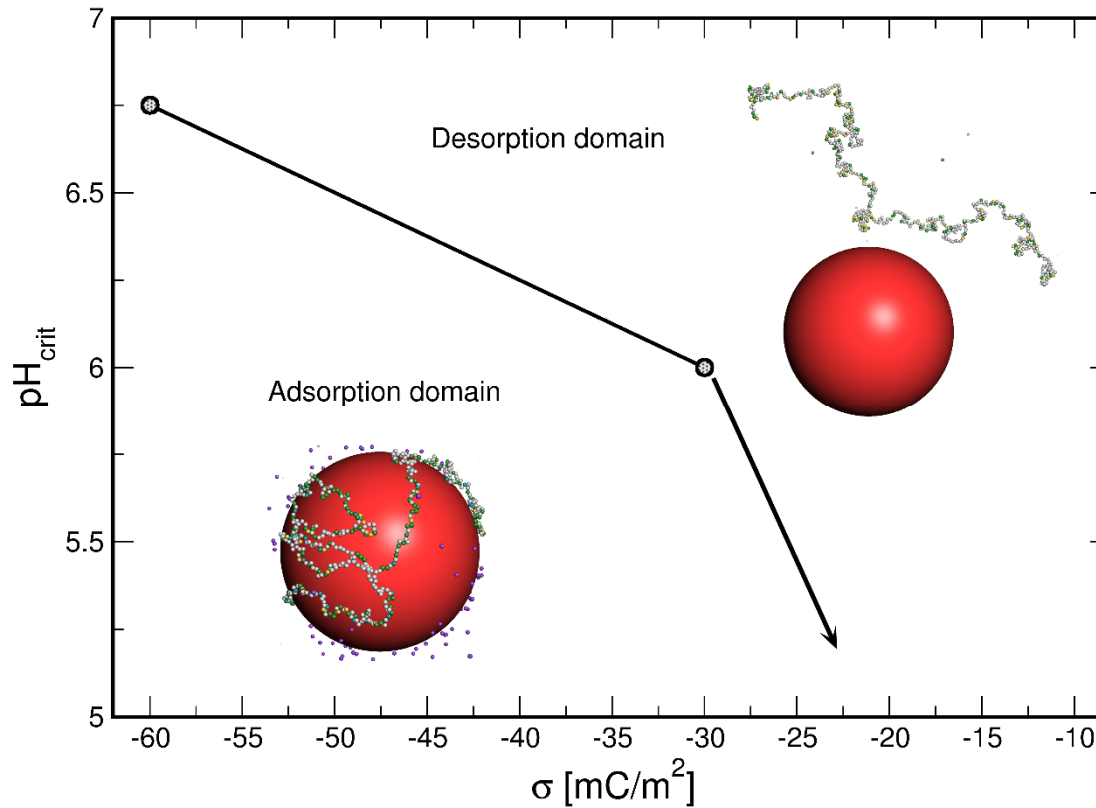
change of the protein conformation and
 modification of the charge of the
 nanoparticle (reactivity)

III. BSA protein



- Global charge of the protein is slightly negative at **physiological pH**
- Presence of the nanoparticle significantly influences the **charging process** of the protein

III. BSA protein



$$R_{NP} + R_{AA} \leq Ads_L \leq R_{NP} + 3R_{AA}$$

Adsorption criterion

One amino acid situated in the adsorption layer (Ads_L) more than 50% of Monte Carlo steps

- Adsorption domain increases with the nanoparticle surface charge density

Prediction of the reactivity of nanoparticles!

IV

Conclusions and perspectives

IV. Conclusions et perspectives

- Structure of strong hydrophobic chains not dependent on pH
 - Denaturation and reactivity limited
- The nanoparticle modifies the acid/base properties of chains, and thus the charging process

BSA protein

- BSA conformations strongly modified at extrem pH and with the adsorption at the nanoparticle surface
 - Denaturation
- No BSA adsorption at physiological pH considering only electrostatic interactions
 - Importance of hydrophobic and structural interactions between the nanoparticle and BSA

Work in progress

- Improvement of the **interactions** between the protein and nanoparticle
- Parametrization and **validation** of the model based on experimental data provided by the University of Ljubljana

A decorative graphic consisting of colored spheres (yellow, green, grey) arranged to form the letters 'T' and 'U'. The 'T' is formed by a horizontal row of five spheres (grey, grey, yellow, green, yellow) and a vertical column of six spheres (yellow, green, green, grey, green, yellow) attached to the center. The 'U' is formed by a vertical column of six spheres (yellow, yellow, yellow, green, green, green) and a horizontal row of three spheres (yellow, grey, grey) attached to the bottom center.

Thank you

for your attention !



Computational Approaches in Nanosciences Workshop

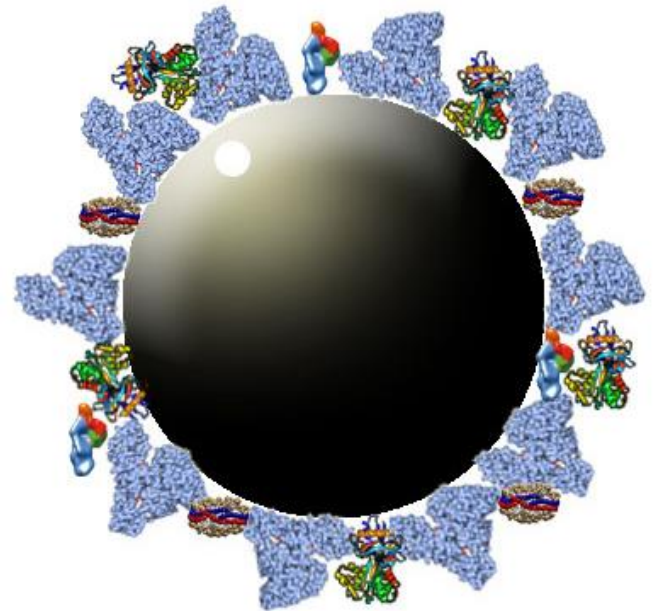
Interaction of carbon-based nanomaterials with cholinesterases and serum proteins

Maja Sopotnik

October 3rd, 2015

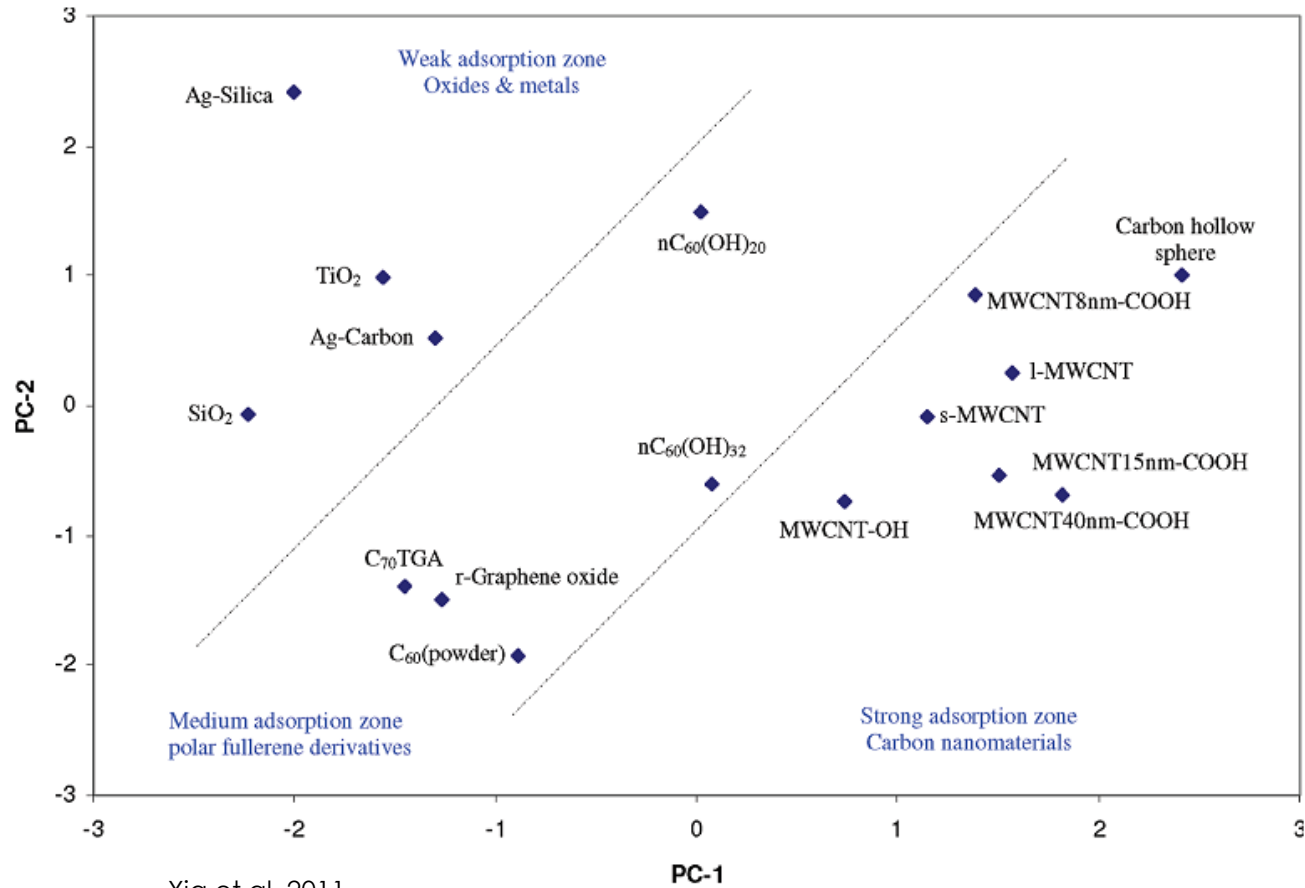
NM in biological media

- › Adsorption of biological molecules on the surface of the NM
- › Protein corona
 - depends on the properties of the NM and the proteins
- › Change in NM characteristics and behaviour
- › The need for biological characterisation of NM



Carbon NM are highly sorptive

- › Surface adsorption index, 5 nanodescriptors



Xia et al. 2011

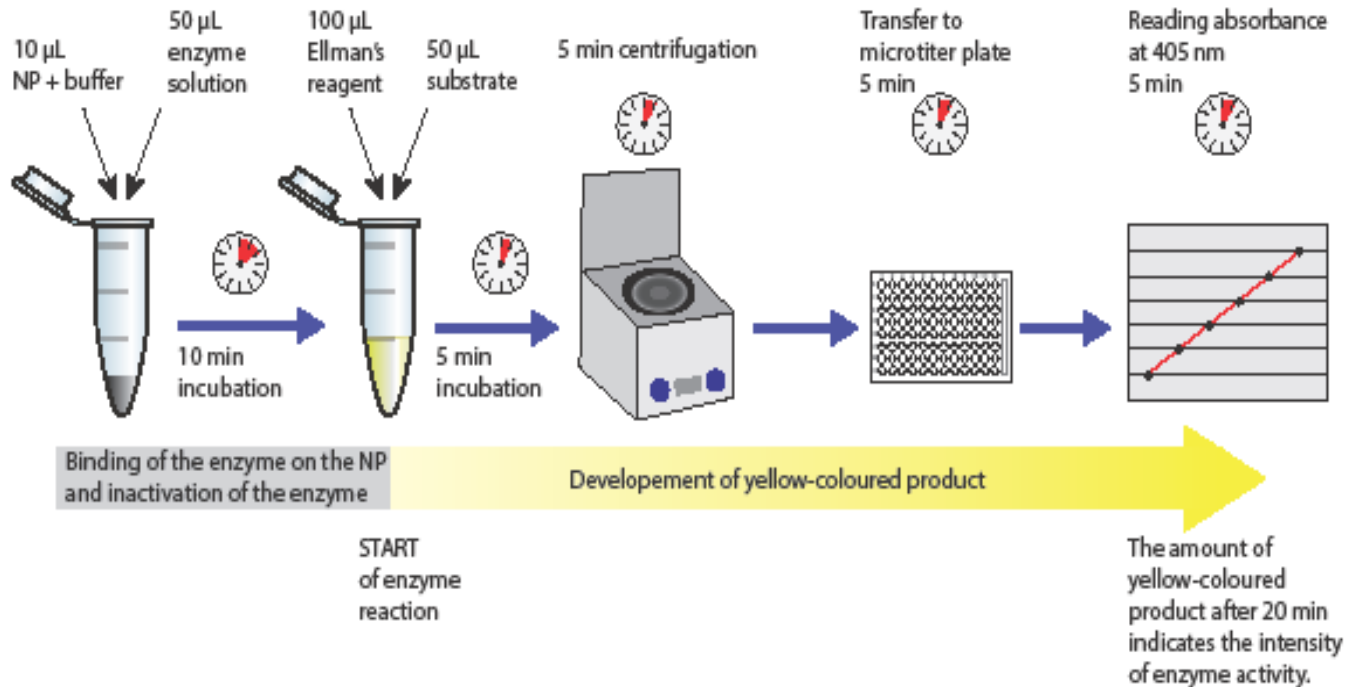




Cholinesterases

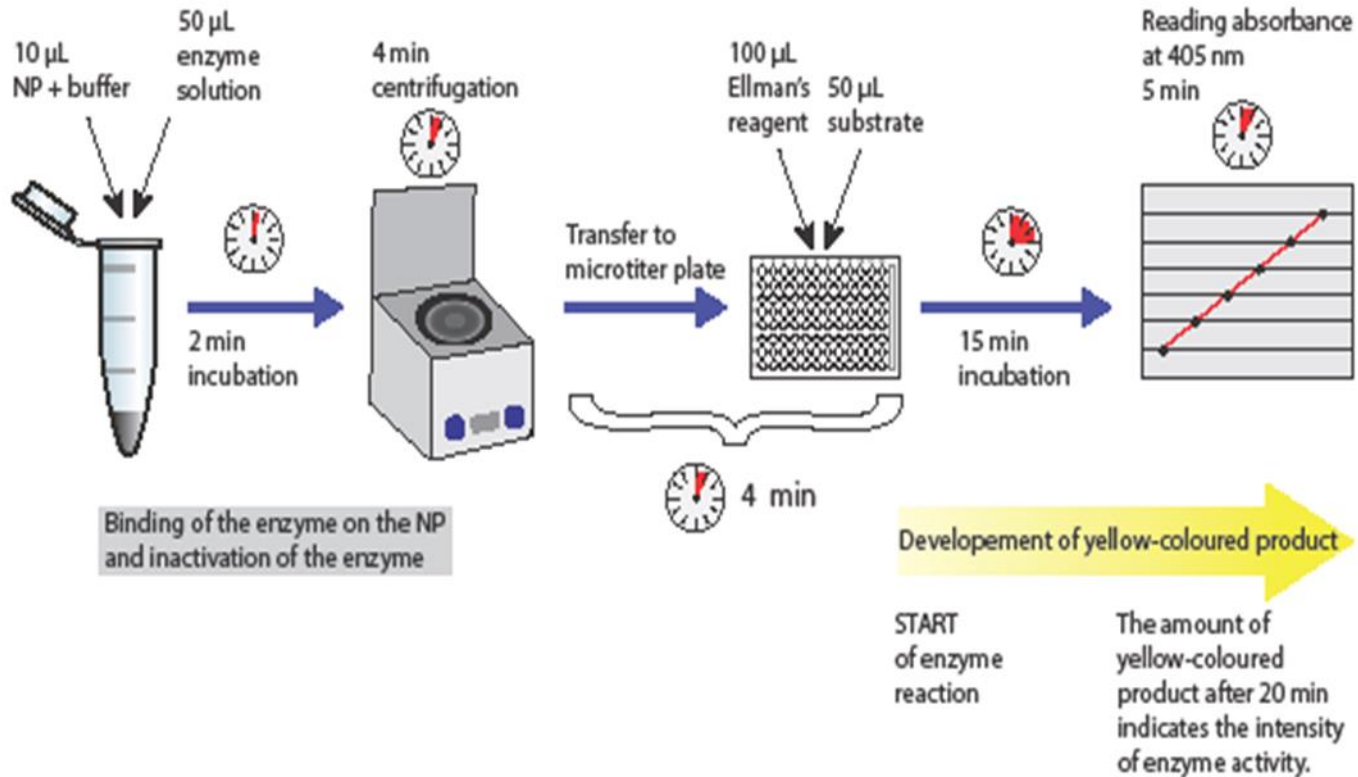
- › Acetylcholinesterase
 - Neurotransmitter acetylcholine degradation
- › Butyrylcholinesterase
 - Backup system in the blood
- › Convenient optical method of measuring the amount of the reaction product – Ellman's method

Measuring the inhibition of enzyme activity by the NM



Jemec et al., in press

Measuring the adsorption of the enzyme on the NM



Jemec et al., in press

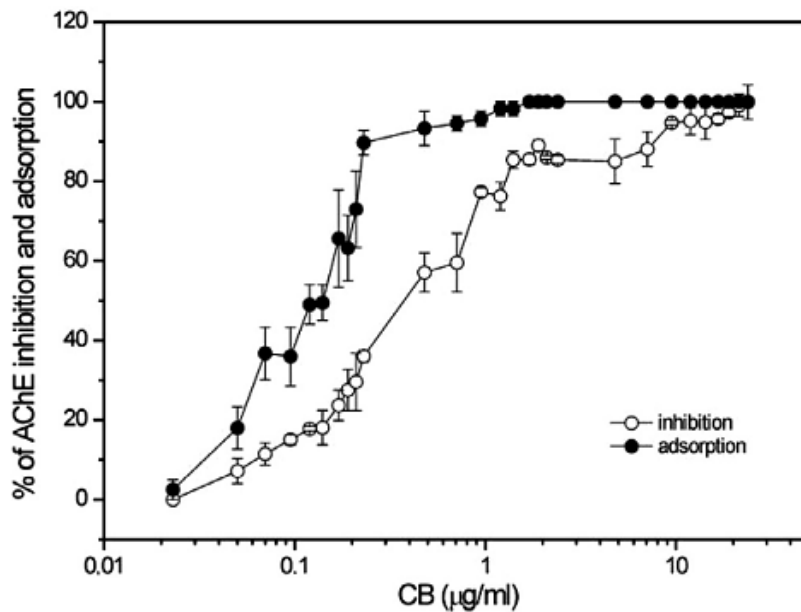




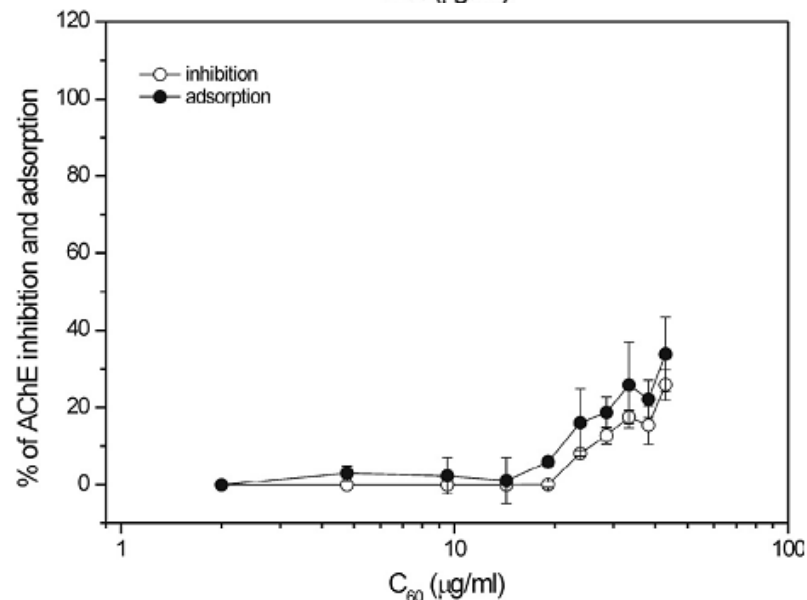
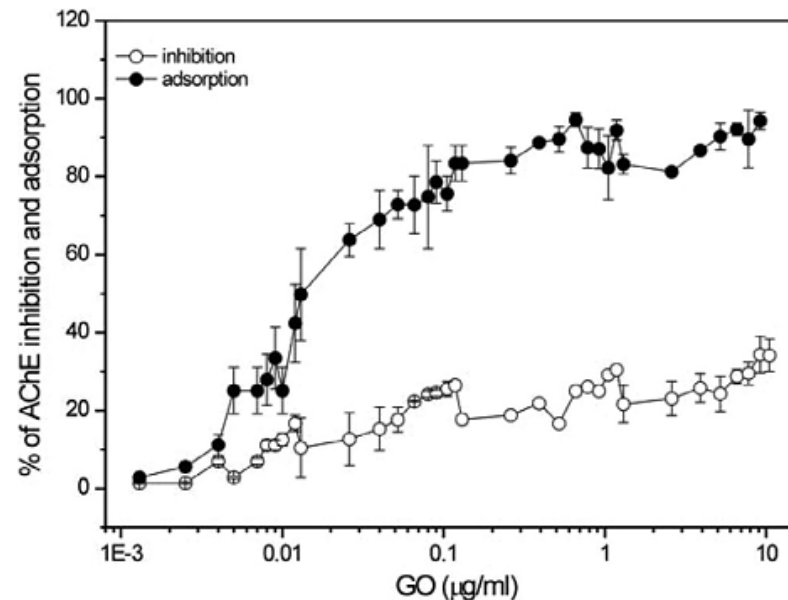
Comparison of different carbon NM

- › Graphene oxide
- › Carbon black
- › C₆₀ – fullerenes
- › Multi-walled carbon nanotubes

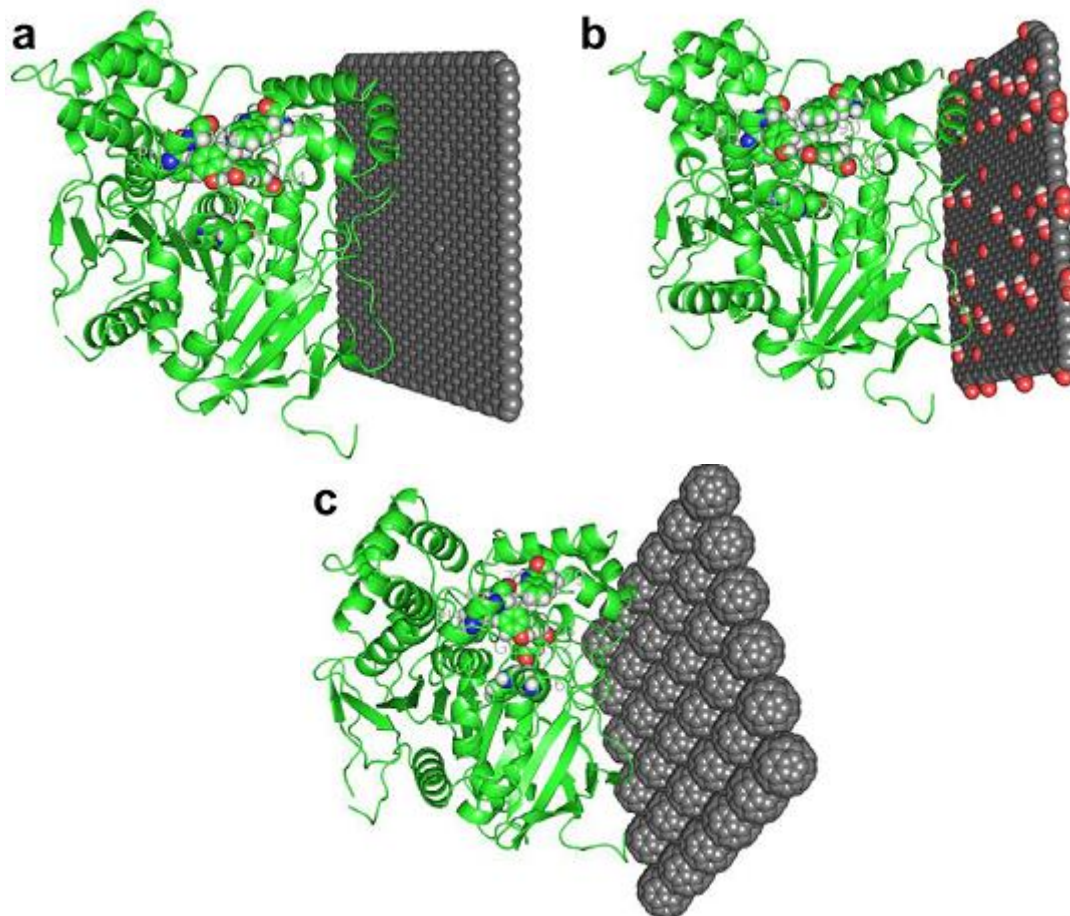
Drosophila melanogaster acetylcholinesterase



Mesarič et al. 2013



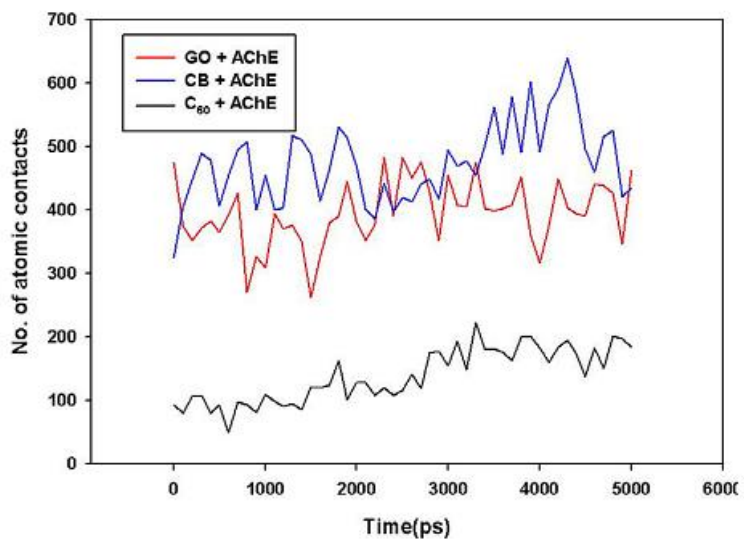
Molecular dynamics simulations



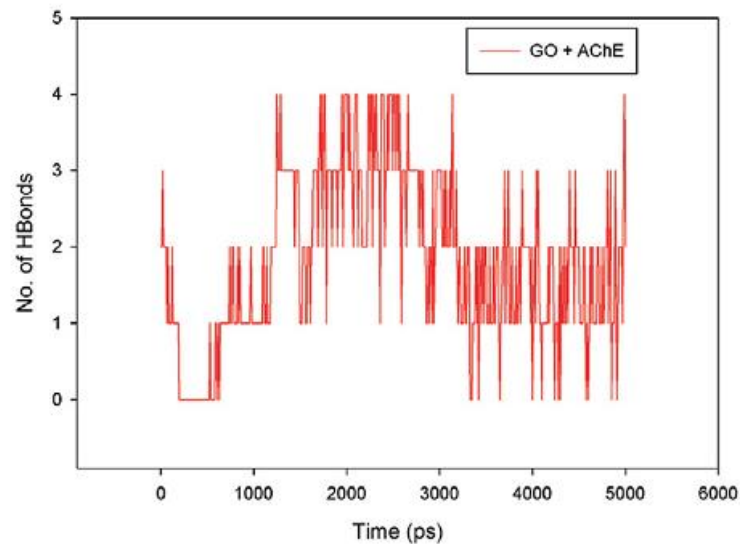
Mesarič et al. 2013



Molecular dynamics simulations



Atomic contacts



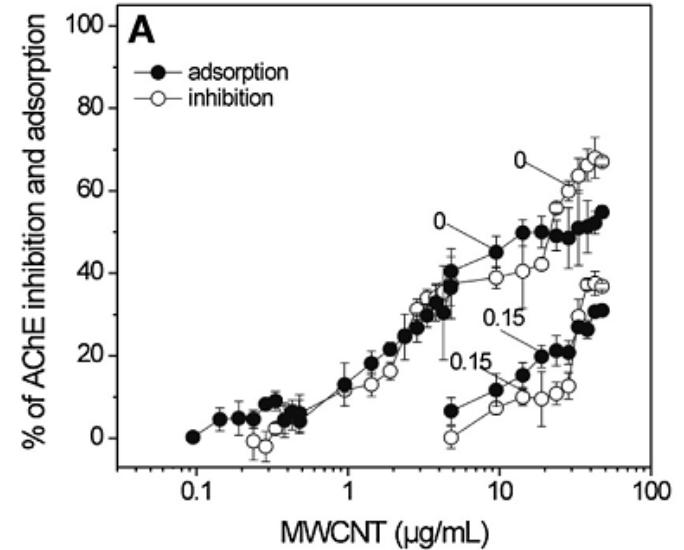
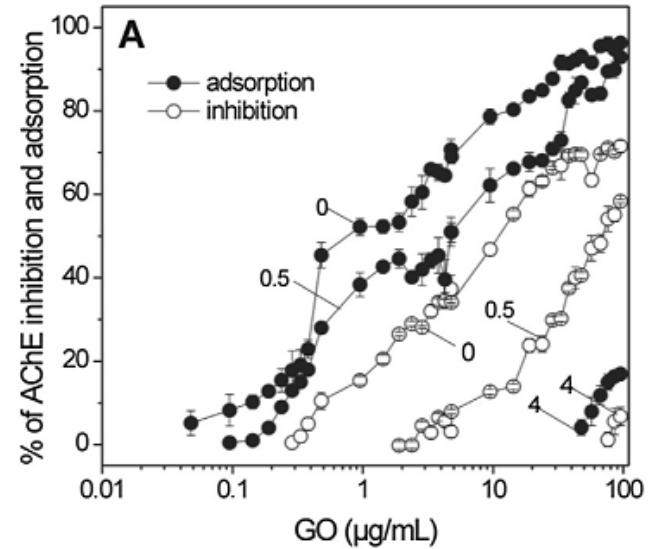
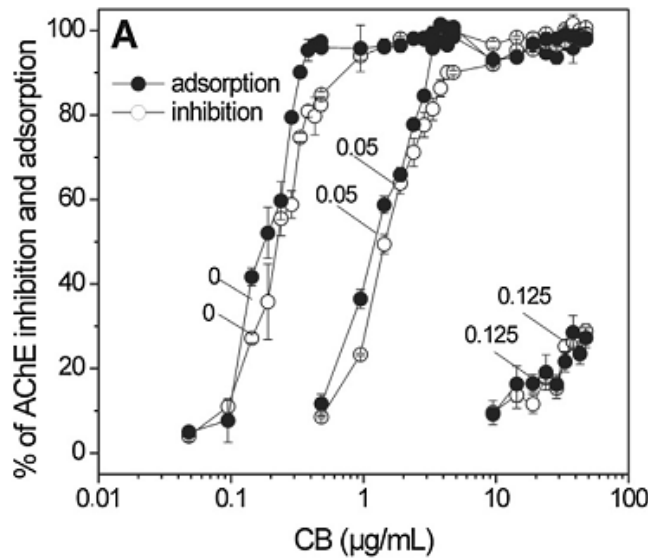
Hydrogen bonds

Mesarič et al. 2013



Electrophorus electricus acetylcholinesterase

> Pre-coating with BSA

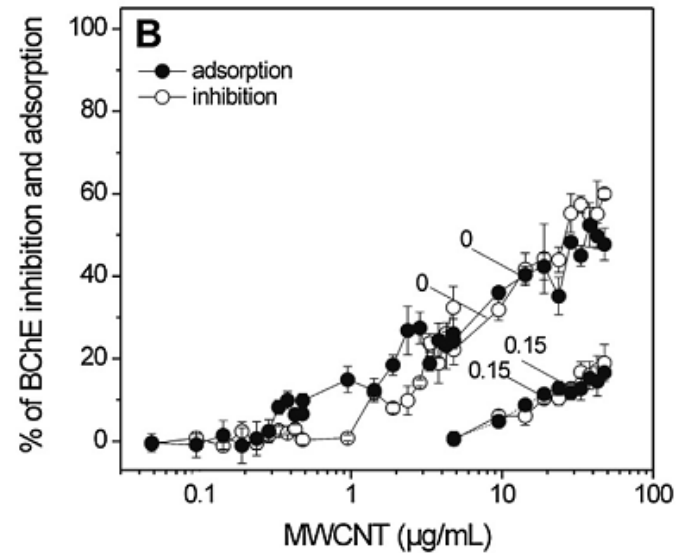
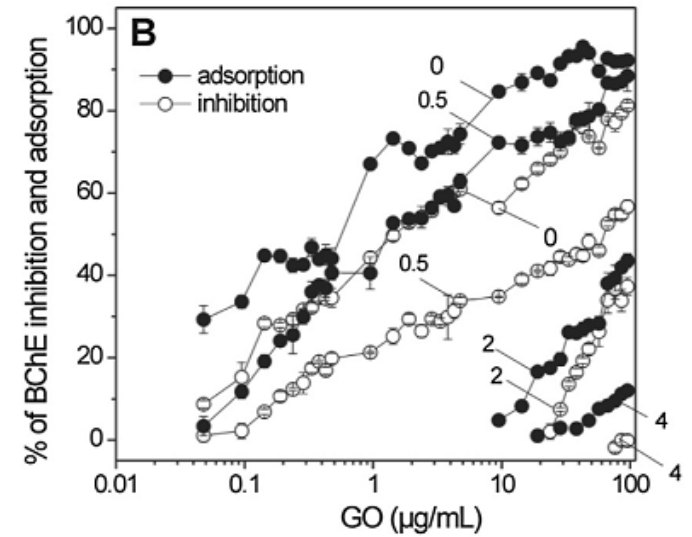
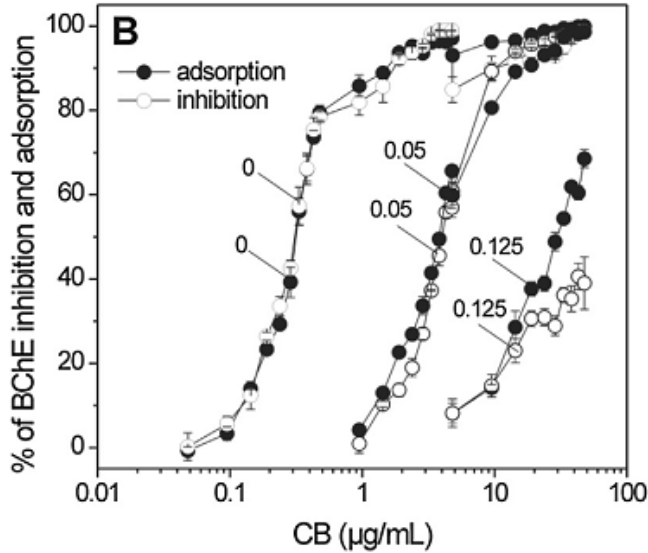


Sopotnik et al. 2015



Equine serum butyrylcholinesterase

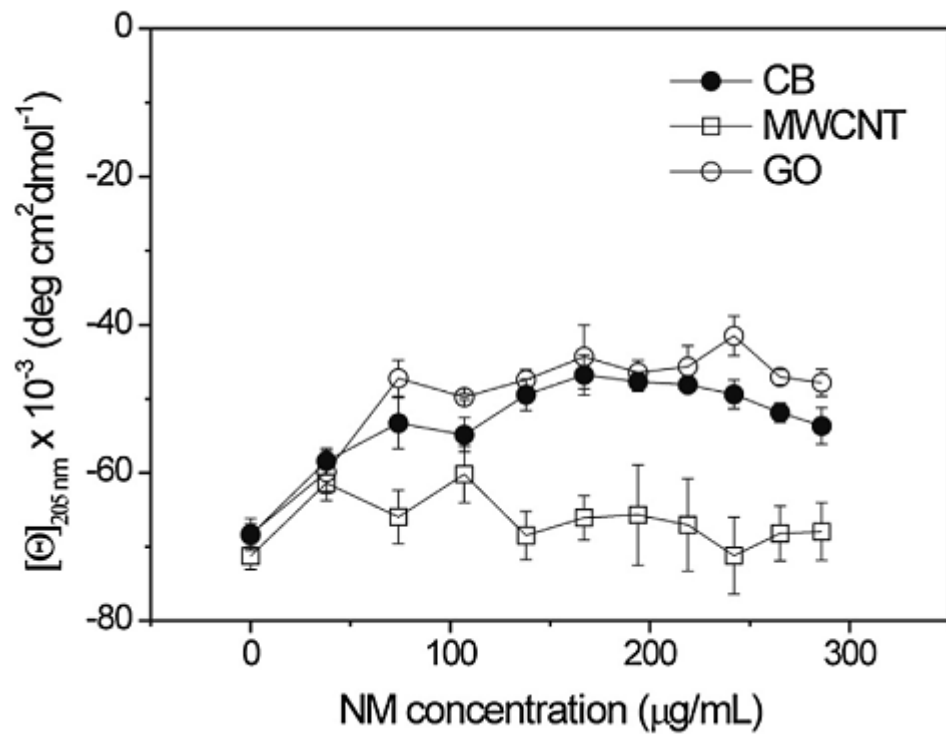
› Pre-coating with BSA



Sopotnik et al. 2015



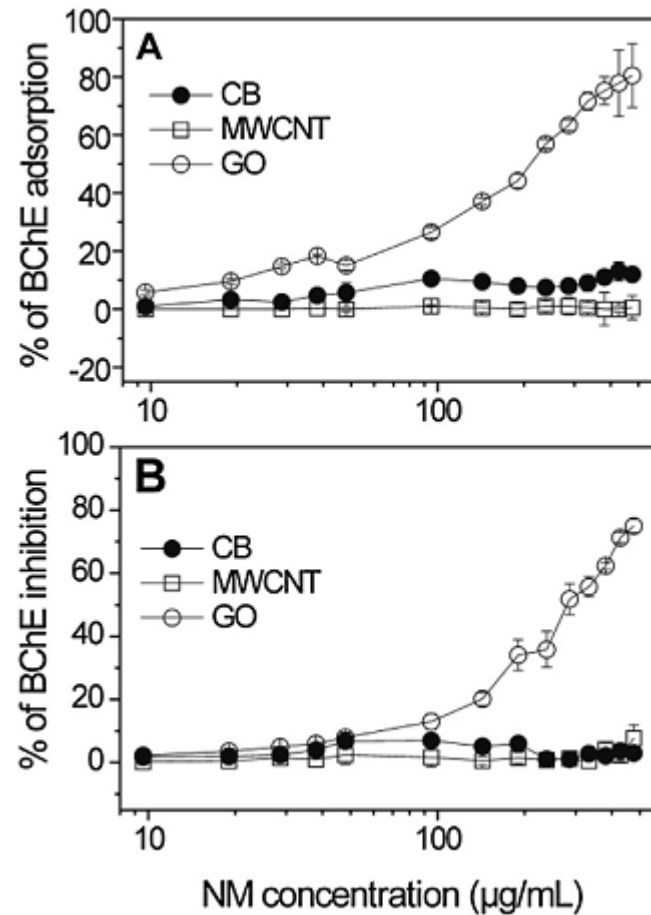
CD-measurements



Sopotnik et al. 2015

Human serum

- > 0,4 % human serum
- > Intrinsic BChE

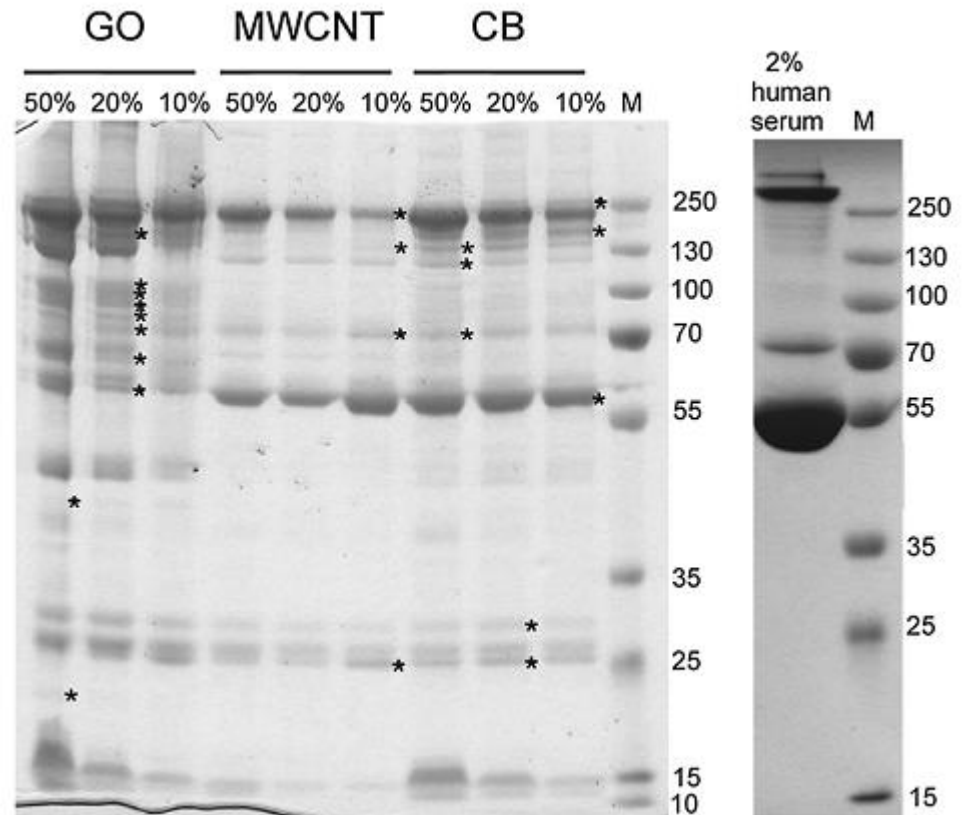


Sopotnik et al. 2015



Human serum

- > Transport proteins
 - Serum albumin
 - Seroransferrin
 - Apolipoprotein A-I
 - Apolipoprotein E
- > Immune system
 - Complement C3
 - Complement C4-A
 - Copmlent C1q
 - Immunoglobulins



Sopotnik et al. 2015





Conclusions

- › AChE inhibition: CB>GO>MWCNT>C₆₀
- › GO : AChE adsorption >> AChE inhibition
- › Binding affinity of NMs for serum albumin was similar in pure albumin solution and in the whole serum
- › CB and MWCNT have a strong affinity for serum albumin
- › GO has a weaker affinity for serum albumin but stronger for other serum proteins, including BChE; is less specific
- › Carbon NM corona is enriched with complement factors and apolipoproteins



Literature:

- › Xia, X. R., Monteiro-Riviere, N A., Mathur, S., Song, X., Xiao, L., Oldenberg, S. J., Fadeel, B. & J. E. Riviere, 2011. Mapping the Surface Adsorption Forces of Nanomaterials in Biological Systems. *ACS Nano* 5(11): 9074-9081.
- › Jemec, A., Mesarič, T., Sopotnik, M., Sepčič, K. & d. Drobne. Biological characterization of nanomaterials. In press.
- › Mesarič, T., Baweja L., Drašler, B., Drobne, D., Makovec, D., Dušak, P., Dhawan, A. & K. Sepčič, 2013. Effects of surface curvature and surface characteristics of carbon-based nanomaterials on the adsorption and activity of acetylcholinesterase. *Carbon* 62: 222-232.
- › Sopotnik, M., Leonardi, A., Križaj, I., Dušak, P., Makovec, D., Mesarič, T., Poklar Ulrich, N., Junkar, I., Sepčič, K. & D. Drobne, 2015. Comparative study of serum protein binding to three different carbon-based nanomaterials. *Carbon* 95: 560-572.

Quality of nanotoxicity data and the importance of harmonization

Dr. Anita Jemec

University of Ljubljana, Biotechnical Faculty, Department of Biology, Večna pot 111, 1000 Ljubljana, Slovenia

Univerza
v Ljubljani
Biotebniška
fakulteta

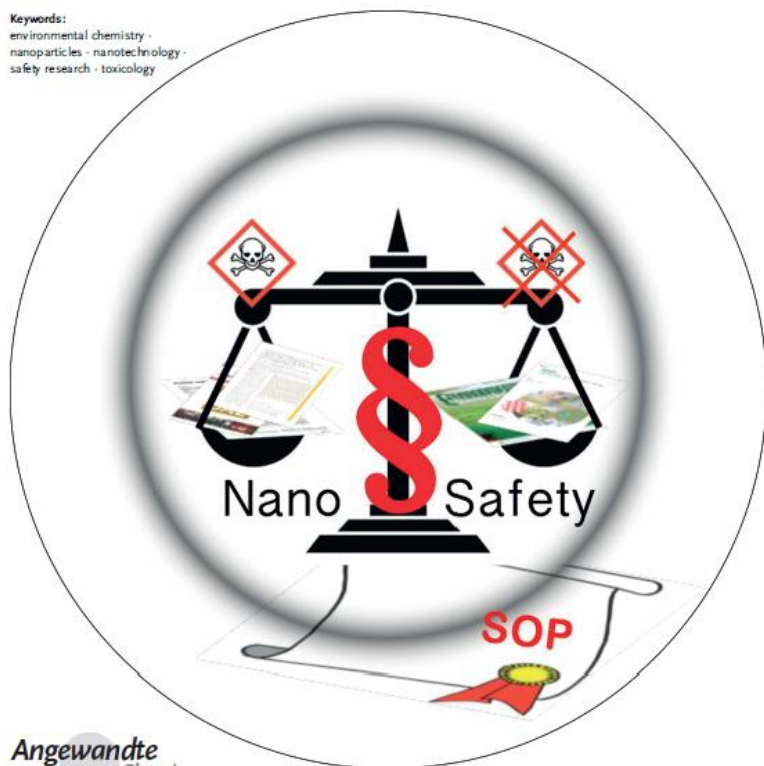


Nanosafety Research—Are We on the Right Track?

Harald F. Krug*

Keywords:

environmental chemistry ·
nanoparticles · nanotechnology ·
safety research · toxicology



Krug et al., 2014.

Screen the nanotoxicity literature of the last 10-15 years
>10 000 publications.

Conclusion:

„ Most of these studies, do not offer any kind of clear statement on the safety of nanomaterials. On the contrary, most of them are either self-contradictory or arrive at completely erroneous conclusions... „



Table 1: Variables in the in vitro toxicity tests during the investigation of ENMs (modified according to Ref. [35]).

Variables associated with the nano-material	Variables associated with the tox assay	Variables associated with the biological model
sample purification for the removal of biologically relevant trace elements	selection of the correct test system regarding the biological end points	selection of the biological system
sample characterization of the raw material: composition and purity size shape agglomeration status etc.	different test systems for the same biological end point	cell lines: selection criteria identification age and storage number of passages etc.
sample characterization regarding biological impurities: endotoxins etc.	controls: adapted negative controls adapted positive controls comparison to reference materials	primary cells/organ systems: donor dependency donor variability culture conditions
dispersion in biological media under relevant conditions: temperature humidity gas concentrations (O ₂ , CO ₂) salinity etc.	testing of possible interferences of the ENM with the biological test system binding of indicator molecules light absorption or fluorescence of the materials etc.	culture conditions during the experiments: temperature humidity gas concentrations (O ₂ , CO ₂) salinity etc.
sample characterization in biological media: size and shape agglomeration status protein corona etc.	measurement uncertainty not considered: round robins calibration with standards or reference material	biological parameter: cell density volume of the medium serum content of the medium compatibility of the solvent or dispersion medium

CRITERIA FOR quality of nanotoxicity data

NanoValid aimed to develop criteria for good quality nanotoxicity data to be used for RA and LCA

-

In order to provide a list of criteria three web-based initiatives, one authority document, one list developed with NanoValid, and two scientific papers (Stefaniak et al., 2013, using various sources, Mills et al., 2014) were used as a basis.

1. MinChar initiative, providing minimal material characterization recommendations for nanotoxicology studies (<http://characterizationmatters.org/parameters/>), see Annex 1
2. DaNa criteria checklist, providing a list to evaluate nanotoxicity studies regarding their scientific value (<http://www.nanopartikel.info/cms/lang/en/Wissensbasis/kriterienkatalog>), (Kühnel et al., 2014))
3. The Nanomaterial Registry's Minimal Information About Nanomaterials (MIAN), <https://www.nanomaterialregistry.org/about/MinimalInformationStandards.aspx> (Mills et al., 2014), see Annex 4
4. Standard information required for Nanomaterials manufactured or imported (REACH) "Nanomaterials and REACH – Background paper on the position of the German competent authorities", (UBA, 2013)
5. List of issues and parameters to be specified for ENPs used in toxicological tests under NanoValid, see Annex 3
6. List of relevant nanomaterials properties provided by Stefaniak et al. (2013, see Table 2, page 1329), see Annex 5

UNIFICATION OF CRITERIA NEEDED!

Physical and chemical properties

- Name of substance (or CAS-No)
- Aggregation / Agglomeration State
- Shape
- Particle Size / Size Distribution (including type of dispersion medium and additives)
- form of delivery (powder, suspension)
- Composition (including chemical composition, elements, element distribution and crystal structure)
- Purity (including levels of impurities)
- Surface Chemistry (including functionalization, reactivity, hydrophobicity)
- Solubility
- Surface Area
- Porosity
- Density
- Defect density
- Surface Charge
- Stability
- Conductivity
- Magnetic properties
- Surface Reactivity
- Consideration of surface modification: exact characterization

Sample preparation

- Adequate characterization of sample: Dissolution for soluble (ion releasing) nanomaterials
- Suitable preparation of the sample: Detailed description of the dispersing procedure
- Determine size in processed sample, do not rely on producer information
- Aggregation / Agglomeration in respective media
- Dispersibility
- Consider age / storage periods of NM powders / suspensions for subsequent testing

Toxicity testing

- Determination of exposure concentration (real vs. nominal)
- Stability—how do material properties change with time, storage, handling, preparation, delivery (aging)
- Behaviour in exposure media over test duration solubility, and the rate of material release through dissolution
- Aggregation / Agglomeration in respective media
- Dose metrics used (mass, surface-area and number concentration in $\mu\text{g}/\text{ml}$, $\mu\text{g}/\text{cm}^2$; N (particle)/cell or pg/cell)
- Controls (positive and negative controls)
- Interferences with test system
- Appropriate methods / endpoints
- Use of reference material



General aspects

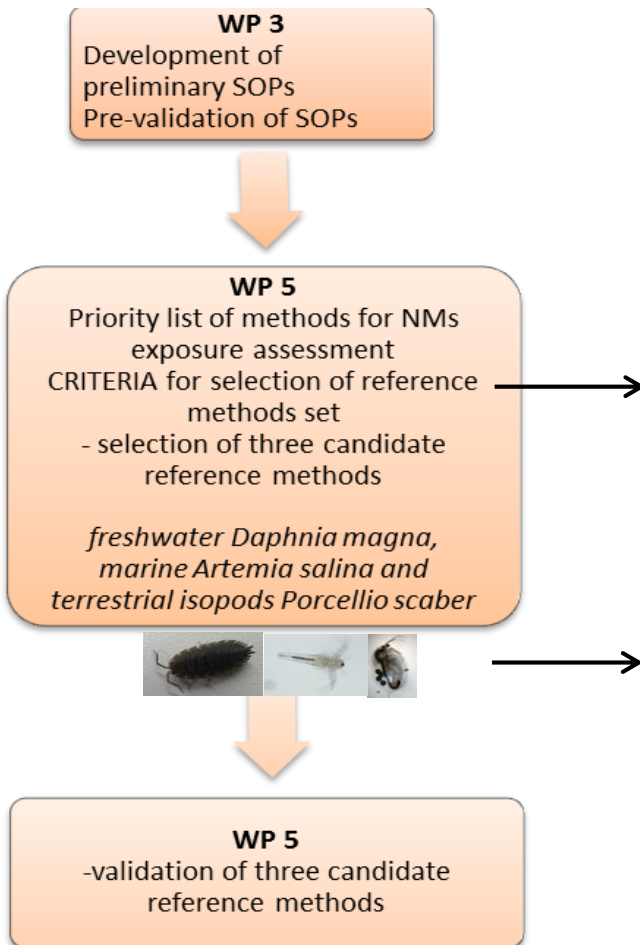
Appropriate data evaluation / statistics

Standardisation criteria (SOPs used, OECD guidelines, decision trees)

Further questions regarding the scientific validity of the data:

- 1) Are raw data provided by the data source?
- 2) Were proper controls used and reported?
- 3) Was the instrument within calibration?
- 4) How many replicates were performed?
- 5) Was the measurement protocol reported?
- 6) Was there a citation to the protocol?
- 7) Were there modifications made to the cited protocol?
- 8) Are there specifications regarding the age of the NM?

STEPS in Establishing reference methods



CRITERIA for reference method:

- sensitive to demonstrate exposure to NPs
- the signal to noise ratio have to be high enough
- the exposure should be linked to effects,
- the test should exhibit good reproducibility, ease of performance, and robustness.

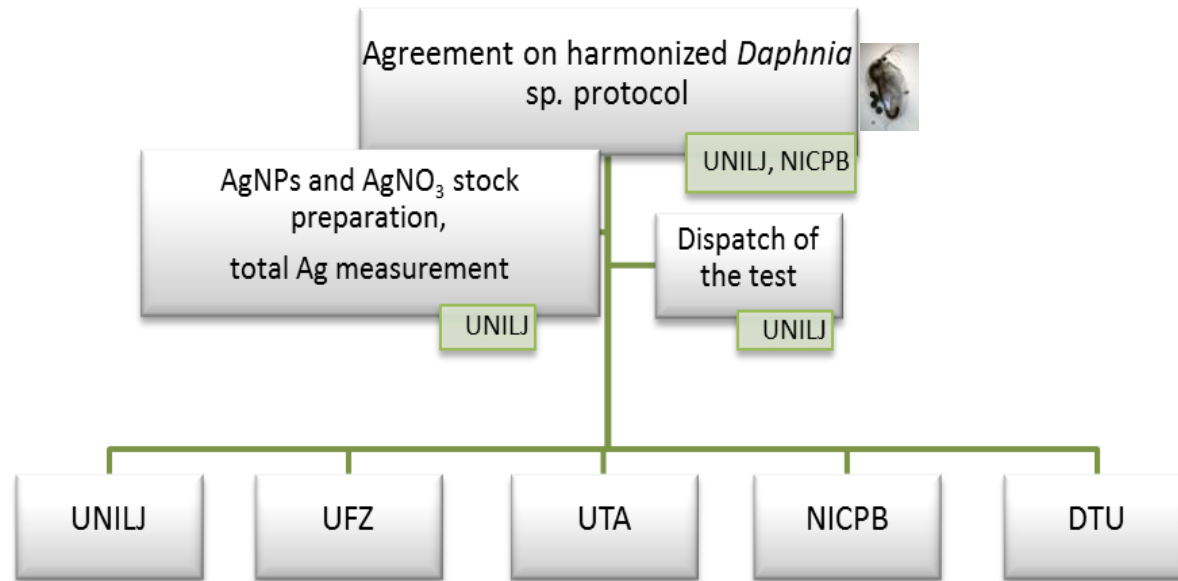
CHOICE:

Daphnia magna
Artemia franciscana
Porcellio scaber

waterflea Daphnia magna



Organization of inter-laboratory comparison study



University of Ljubljana (UNILJ, Biotechnical Faculty, Slovenia) (reference laboratory), National Institute of Chemical Physics and Biophysics (NICPB, Estonia), The Helmholtz Centre for Environmental Research (UFZ, Germany) Technical University of Denmark (DTU, Denmark)



RESULTS: *Daphnia magna*

Exactly after SOP developed by NICPB (OECD 2004: No. 202)

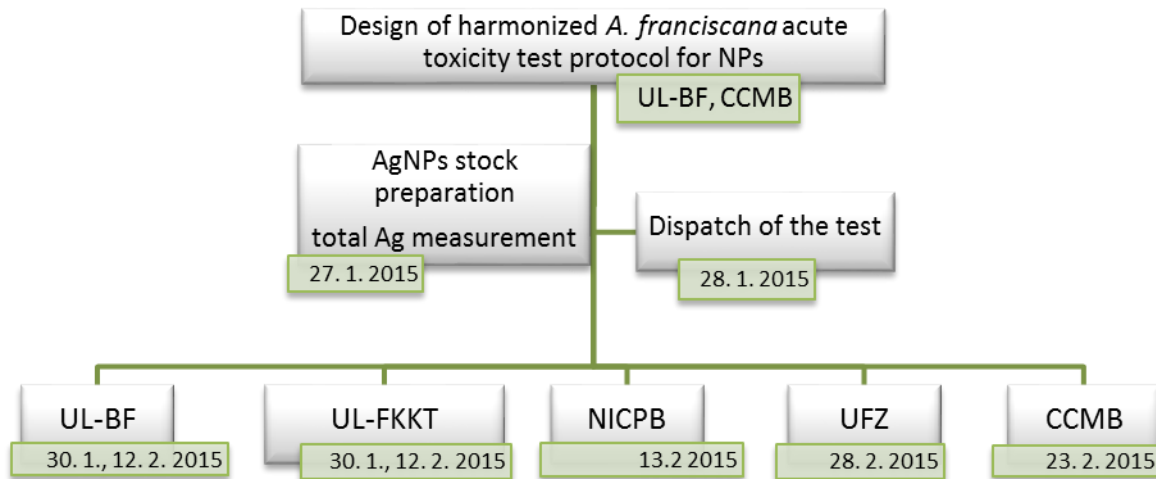
PARTNER	48h EC50, AgNPs µg/L	48h EC50, Ag⁺ µg/L
1: NICPB reference laboratory	2.50	1.00
2: UNILJ	5.37	3.69
3: UFZ	2.39	1.28
4: DTU	4.081	3.21
5: UTA	18.85	2.02

- Good inter-laboratory repeatability

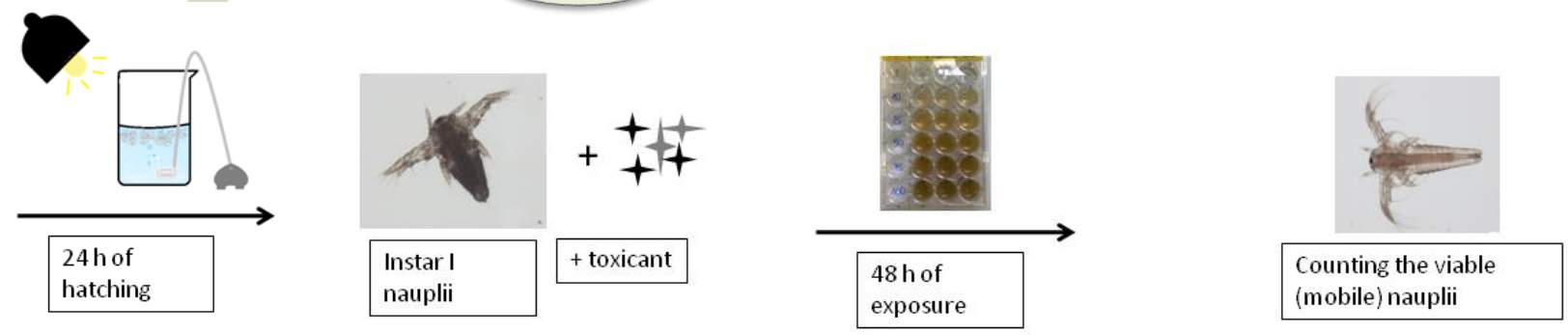
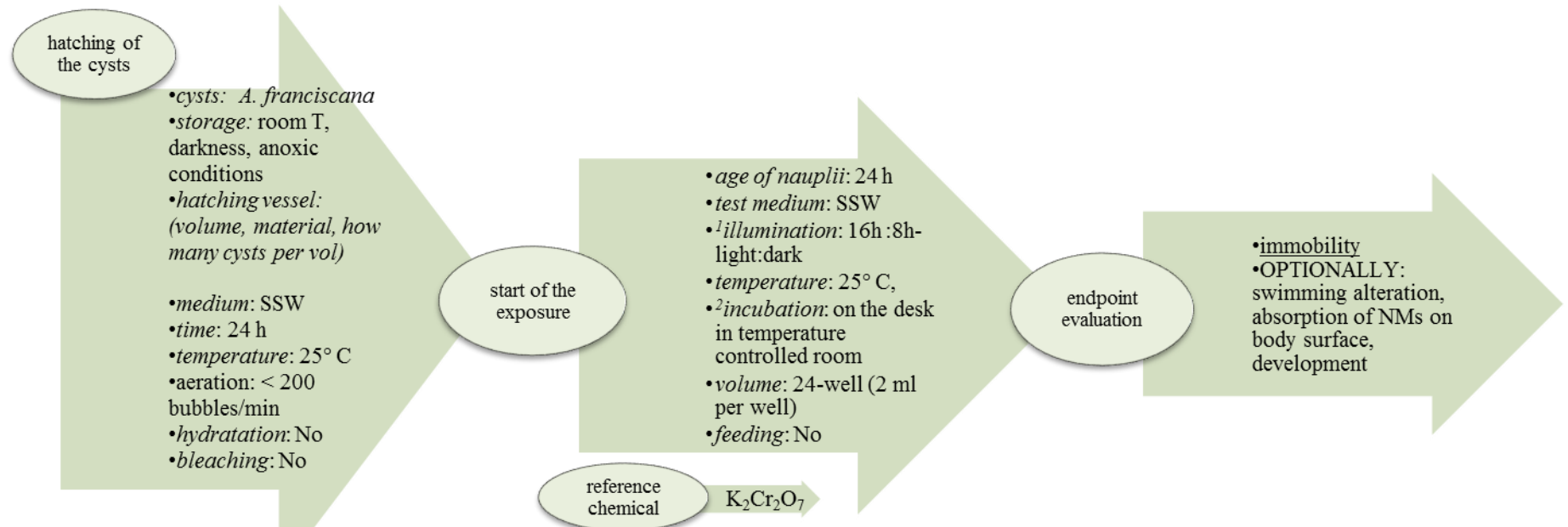
brineshrimp Artemia franciscana



Organization of inter-laboratory comparison study



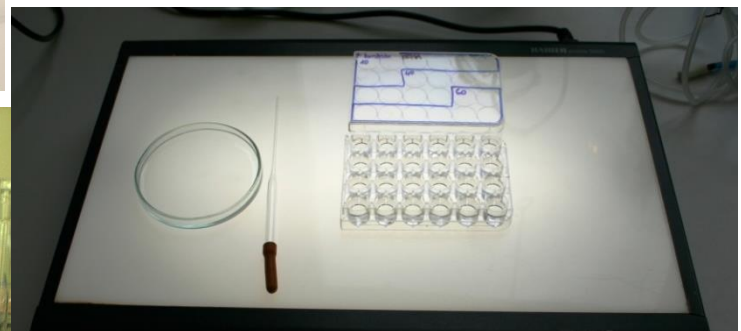
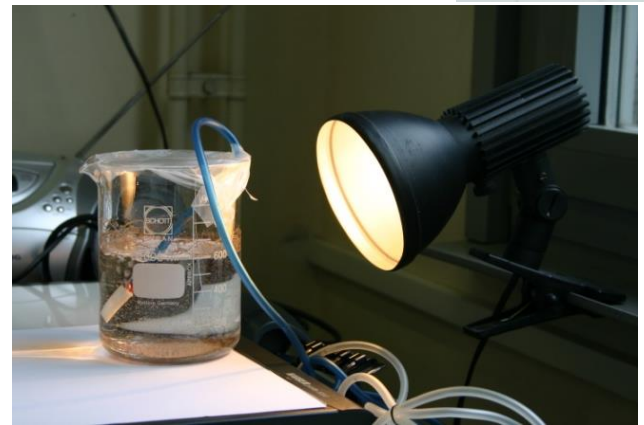
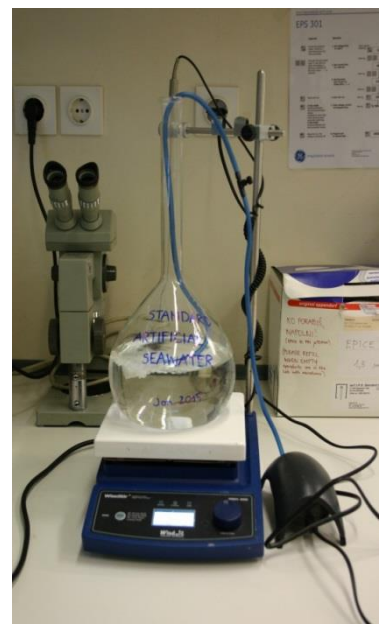
First the harmonised protocol was developed- literature review and experimentation





Experimental set-up.

Negative control: ASW medium
Test compounds AgNPs (mg/L):
25
50
75
100
125
Positive control; $K_2Cr_2O_7$ (mg/L)
15
30
60

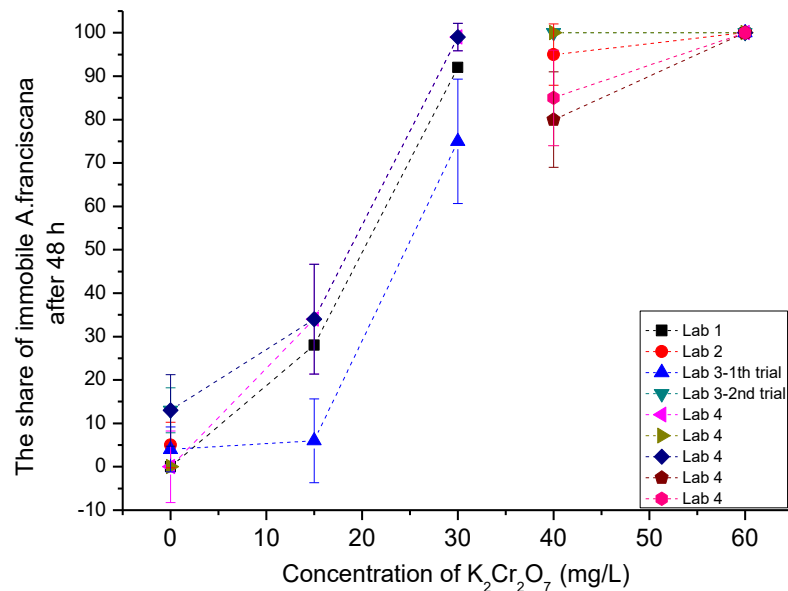




RESULTS: POTASIAM DICHROMATE

Data of the intercalibration study on $K_2Cr_2O_7$.

a.)



CONCLUSIONS:

- 1 partner invalid results-high control mortality (inappropriate cysts storage, prolonged hatching)
- LOW Intra-laboratory variation (UNILJ) (relative coefficient of variation): 9.6 %.
- LOW Inter-laboratory variation: 17%

GOOD REPEATABILITY WITH CHROMIUM

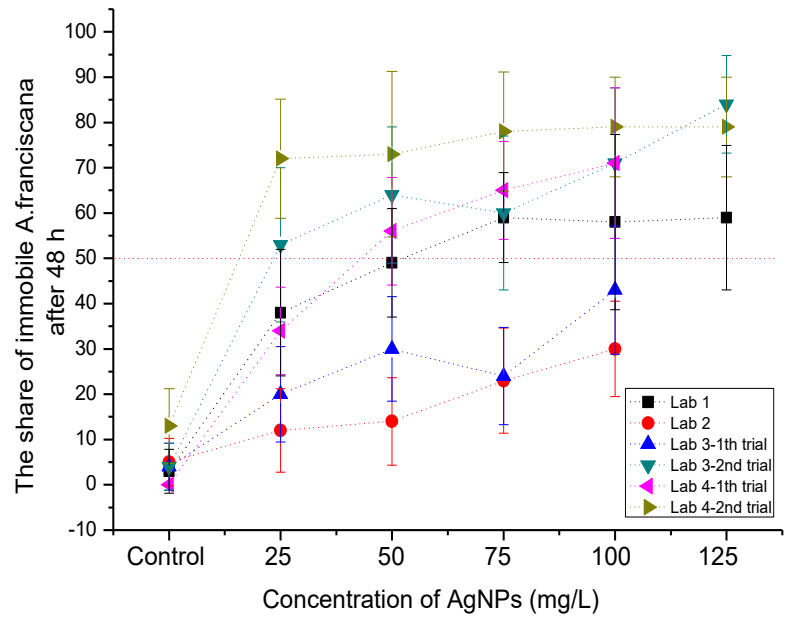
Each data point represents mean \pm SD
(10 repetitions were done for each concentration).



RESULTS: SILVER NANOPARTICLES

Data of the intercalibration study on AgNPs

b.)



Each data point represents mean \pm SD
(10 repetitions were done for each concentration).

CONCLUSIONS:

- 1 partner invalid results-high control mortality (inappropriate cysts storage, prolonged hatching)
- High Inter-laboratory variation (relative coefficient of variation): 37 %
48h EC50 value was 36,48 mg/L

NOT REPEATABLE.
SOURCES OF VARIABILITY?

IDENTIFICATION OF SOURCES OF VARIABILITY

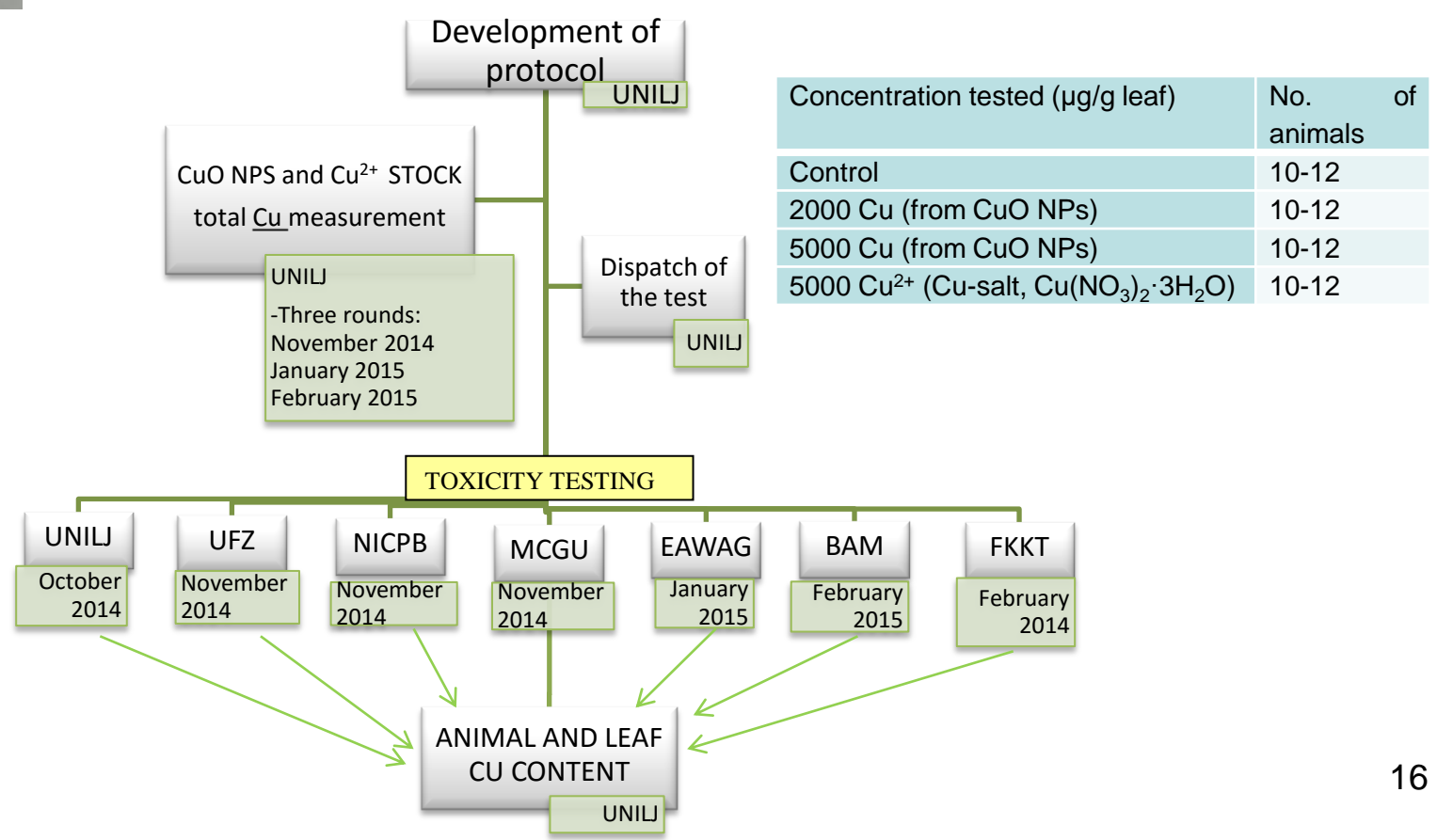
- hatching conditions
- test plate incubation
- illumination regime
- Special AgNPs properties
- light dependent Ag⁺ speciation

	Source of variability	Actions taken to diminish the variability	Previous practises in nanotoxicity assays
HATCHING PHASE	Species of Artemia	Defined species, <i>A. franciscana</i>	Commonly undefined, mostly <i>Artemia sp.</i> or <i>A. salina</i>
	Origin of Artemia cysts	<ul style="list-style-type: none"> ◦ Producer, which supplies <i>A. franciscana</i>, Great Salt Lake, USA ◦ Further suggestion: use of certified cysts 	Very variable sources, commonly undefined
	Hatching medium	Defined composition, artificial salt water (SSW)*	Undefined composition: sea water or Instant Ocean®, Red Sea Salt®, Synthetica Sea salts®, Tropic Marin® – a synthetic sea salt mix
	Illumination	Continuous light	Continuous light, or not reported
	Temperature	Defined temperature: 25 °C	25-30 °C, often not reported
	Duration	Defined duration: 24 h	24 h or 48 h
	Aeration	Defined rate: < 200 bubbles/min, partners reported that hatching was unsuccessful at very high rates	Not defined
	Hydratation/bleaching	Not applied, hatching was successful without this step	Various practises, but usually not applied
TOXICITY TEST	Test medium	Defined composition, synthetic salt water (SSW)*	Undefined composition: sea water or Instant Ocean®, Red Sea Salt®, Synthetica Sea salts®, Tropic Marin® – a synthetic sea salt mix
	Age of cysts	Additional experiments done, 24 h old nauplii (stage I) ensure sufficient survival of controls during 48 h exposure	Various practises: 24 h, 30 h, 48 h, not reported
	Illumination	Additional experiments done, illumination is extremely important in nanotoxicity studies, we suggest	Often not reported, Various practises: dark, 16 h/8 h light/dark regime

isopods *Porcellio scaber*



Organization of inter-laboratory comparison study



RESULTS: *Porcellio scaber*



Detailed instructions, video/audio material



Animal Cu content- Bioaccumulation
Leaf Cu content





- Internationally recognized toxicity test assay the “Feeding assay with *Daphnia magna*” fulfils the criteria for the reference nanomaterial exposure and effect method.
- The assay with *Artemia franciscana* has a number of advantages as a test organism and fulfils a number of criteria as a reference method. The reproducibility of the assay with the reference chemical $K_2Cr_2O_7$ was good, but this was not the case with AgNPs. We attribute this to specific properties of these NPs.
- The “Feeding assay with isopod *Porcellio scaber*” has proved to be a reliable and reproducible assay and we therefore suggest it as reference method for terrestrial nanomaterial exposure and effect. We therefore suggest further steps to standardise the protocol.



harmonisation

quality

Data
sharing

GOOD PRACTISE EXAMPLE:



Information about nanomaterials
and their safety assessment

<http://www.nanoobjects.info/en/>



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