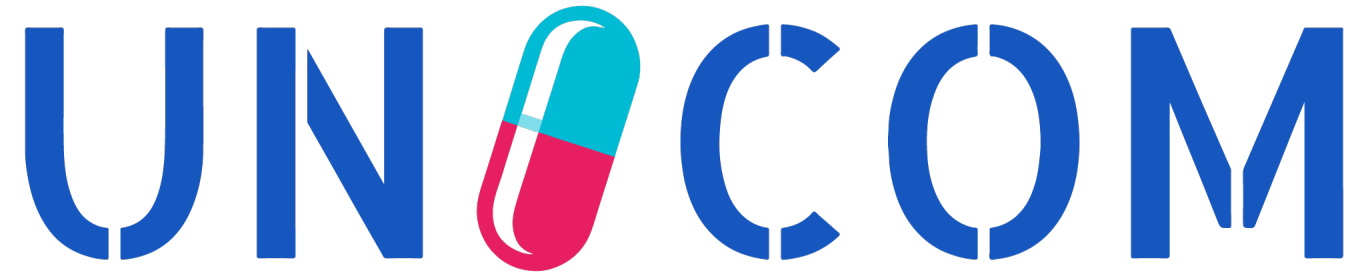


WP-1 / community of expertise

January 2022





Vaccine challenges - cleansing, confidentiality and vaccine naming

Annet Rozema
Olof Lagerlund
Jean-Gonzague Fontaine



SOME RULES FOR THE VIRTUAL MEETINGS

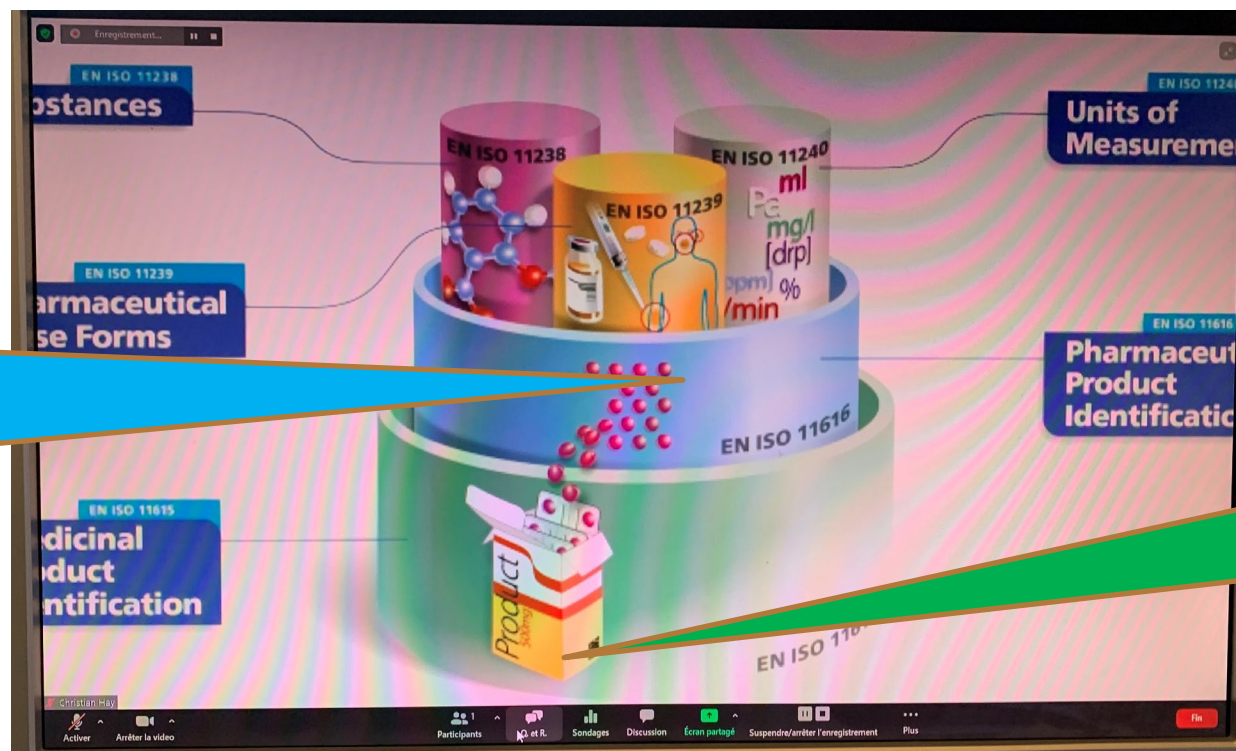
- ✓ **Everybody is on mute**
- ✓ **You post your question in the Q&A facility**
- ✓ **When you speak, please keep concise**
- ✓ **You may show your approval !**

After (and during) the introduction presentations, any UNICOM related question / comment may be shared with Q&A



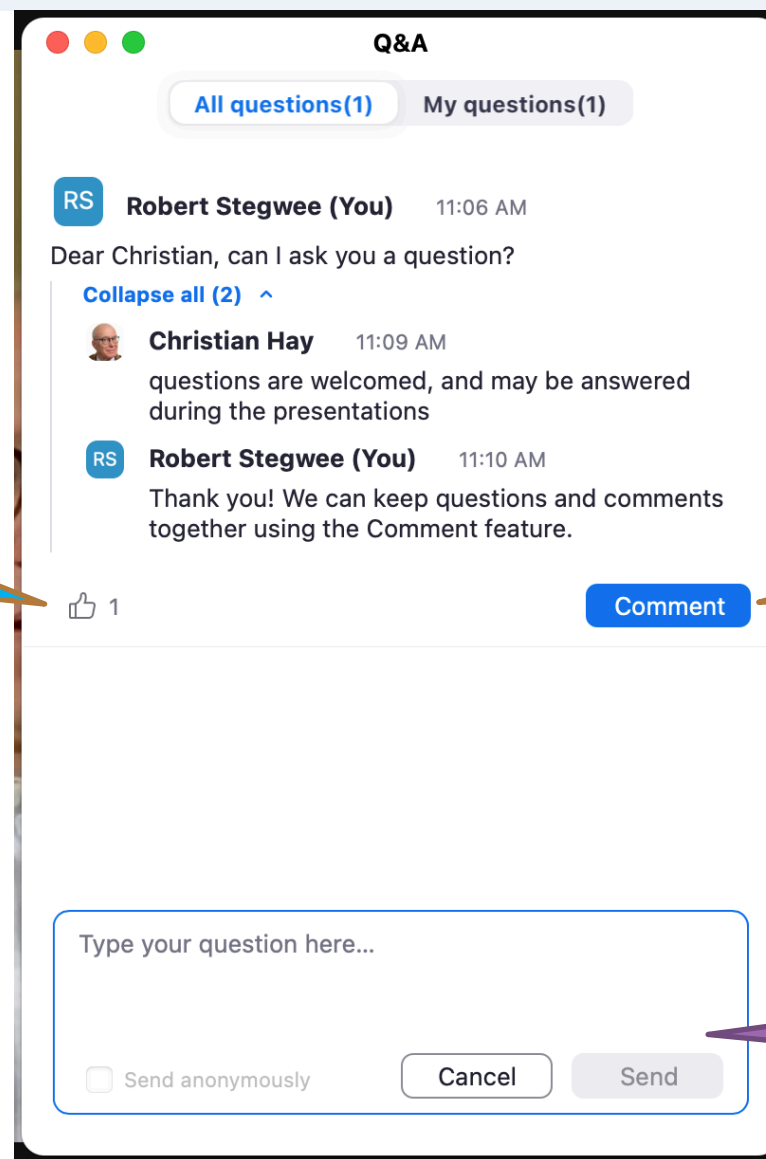
Asking a question or making a comment: please use the Q&A facility

1. Move the mouse on the screen to have the options bar appearing



2. You then select «Q&A» and write your question

You can support a question by clicking the «thumbs up» which moves it up on the list for the presenters



You can comment on a question or answer to engage in a conversation

Typing and sending a new question does not retain the context of your comment

- ▶ Security is our priority
- ▶ This session is password protected



Recording of this session is made available on UNICOM's youtube channel.

At the end of the virtual session, a questionnaire will be sent to the participants, to help us understand participant's reactions and needs



Introductions to our esteemed colleagues and today's speakers



Annet Rozema



Olof Lagerlund



Jean-Gonzague Fontaine

...and our panellist



Raffaella Balocco



Malin Fladvad

This project has received funding from the European Union's
Horizon 2020 research and innovation programme under grant agreement No 875299



Vaccines – A global perspective

Olof Lagerlund,
WP1, SVG and UMC

2022-01-14

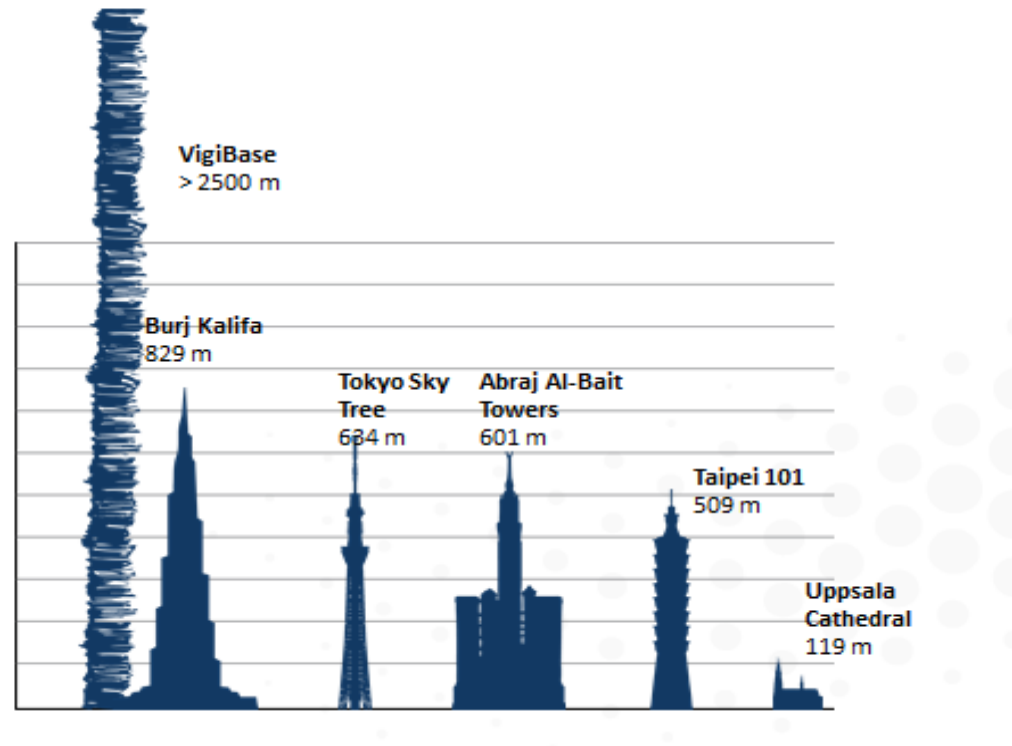
Outline

- Global Vaccine Pharmacovigilance
- Global PhPID's
- Global vaccine initiative
- Final remarks


WHO Program for international drug monitoring (PIDM) and Uppsala Monitoring Centre

- The WHO PIDM was created in 1968 to ensure that evidence about harm to patients was collected from as many sources as possible
- The WHO PIDM had more than 170 full members and associate members in 2021, the program covers about 99% of the world's population.
- UMC has been responsible for the technical and operational aspects of the WHO PIDM since 1978.

VigiBase- the global data base of ICSR's



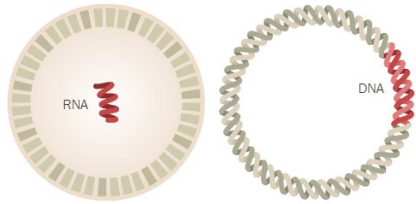
COVID-19 Vaccine ICSR's in VigiBase



COVID-19 vaccine	Number of ICSRs	%
All vaccines	2 720 548	100
AstraZeneca	676 966	25
Comirnaty, Pfizer BioNTech	1 315 056	48
Convidecia	649	0
CoronaVac, Vero Cell, Sinovac	44 583	2
Covaxin, Bharat Biotech	1 566	0
Janssen	114 889	4
Moderna	504 906	19
mRNA, Unspecified	341	0
Sputnik V	4 422	0
Vero Cell, Sinopharm, Beijing	43 730	2
Vero Cell, Unspecified	2 458	0
Other	167	0
Unspecified	10 815	0

Genetic Vaccines

Vaccines that deliver one or more of the coronavirus's own genes into our cells to provoke an immune response.



Comirnaty
COVID-19 mRNA Vaccine BNT162b2
Pfizer BioNTech COVID-19 vaccine
Vacuna COVID-19 Pfizer BioNTech

Tozinameran (INN)
PF 07302048
BNT162b2

Spikevax
Moderna COVID-19 vaccine
COVID-19 Vaccine Moderna
Vacuna COVID-19 Pfizer BioNTech

Elasomeran (INN)
CX-024414
MRNA 1273
TAK 919

Vaxzevria
Covishield
AstraZeneca COVID-19 vaccine
Vacina COVID-19 (recombinante)
COVID-19 vacuna AstraZeneca

ChAdOx1 nCoV-19
AZD 1222

Sputnik
Sputnik light
Gam-COVID-Vac
Vacina COVID-19 (recombinante)

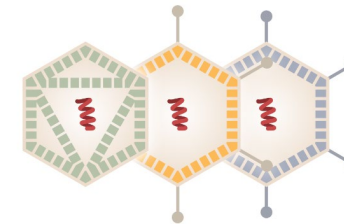
COVID-19 vaccine NRVV Ad26
(Gam-Covid-Vac)
COVID-19 vaccine NRVV Ad5 (Gam-
Covid-Vac)
AZD 1222

Convidecia
Ke wei sha
PakVac
Recombinant COVID-19 Vaccine
(Adenovirus Type 5 Vector)

COVID-19 vaccine NRVV Ad5 (Ad5-
nCoV)
Ad5-nCoV

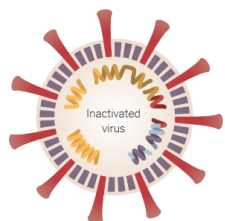
Viral Vector Vaccines

Vaccines that contain viruses engineered to carry coronavirus genes. Some viral vector vaccines enter cells and cause them to make viral proteins. Other viral vectors slowly replicate, carrying coronavirus proteins on their surface.



Inactivated or Attenuated Coronavirus Vaccines

Vaccines created from weakened coronaviruses or coronaviruses that have been killed with chemicals.



CoronaVac
Covid-19 vaccine (vero cell),
inactivated
Vaksin COVID-19 bio farma

COVID-19 vaccine inact (Vero)
CZ02

CanSino COVID Vaccine

Covilo
COVID-19 vaccine (vero cell),
inactivated
Hayat vax
SARS-CoV-2 vaccine (vero cell),
inactivated
Zhong ai ke wei

COVID-19 vaccine inact (Vero)
HB02

Sinopharm COVID vaccine, Beijing

Covilo
Covid-19 vaccine (vero cell),
inactivated
Zhong kang ke wei

COVID-19 vaccine inact (Vero)
WIV04

Sinopharm COVID vaccine,
Wuhan

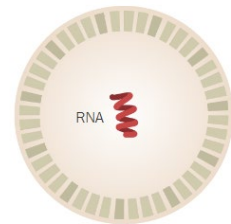
COVID-19 product and substance nomenclature

Regional differences - Dose Form

Covid-19 vaccine	
Authority of approval	AMD dose form
EMA	dispersion for injection
UK	solution for injection
US	suspension for injection



Pfizer BioNTech COVID-19 vaccine
Tozinameran



mRNA vaccine

Regional differences - Strength expression

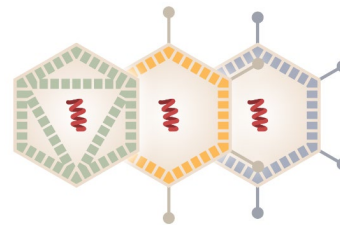
Variation in use of units for strength expression for similar products

Covid-19 vaccine AstraZeneca

Authority of approval	Strength per dose (0.5 ml)
EMA	2.5×10^8 infectious units
UK	5×10^{10} viral particles
Australia	5×10^{10} viral particles



Covid-19 vaccine AstraZeneca ChAdOx1 nCoV-19

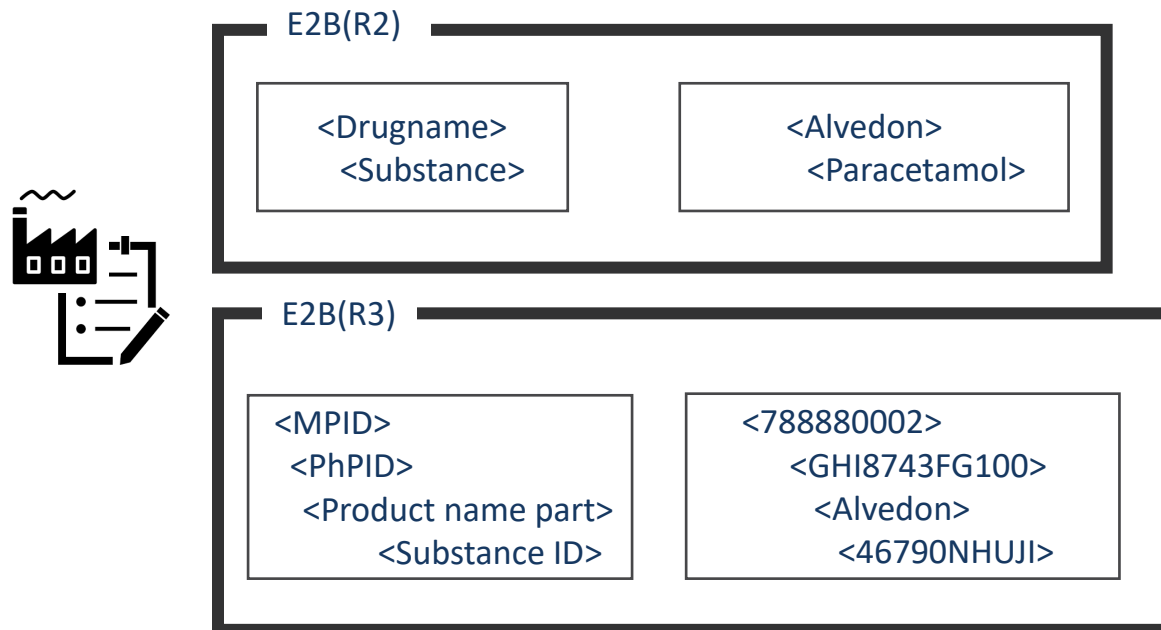


Viral vector vaccine

IDMP in post marketing safety ICH E2B(R3)

Each ICH region to implement their own guidelines for ISO IDMP implementation in ICH E2B (R3)

For EU, the five ISO IDMP standards apply to both authorised and developmental medical products that are regulated in the EU and should be used in ICSR submissions to EudraVigilance when available

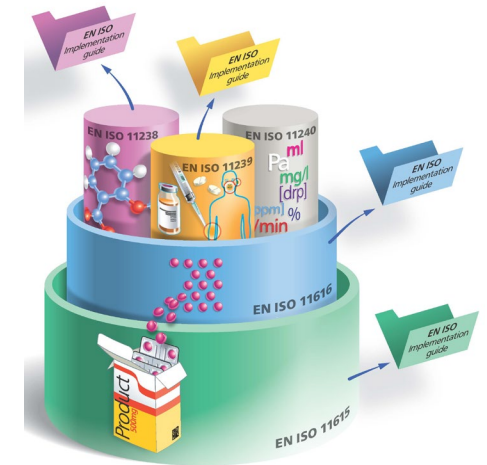


Harmonisation of data using the ISO IDMP suite of standards



PhPID Set

- PhPID Level_**L1** → Substance(s) Term
- PhPID Level_**L2** → Substance Term(s) +Strength+ reference strength
- PhPID Level_**L3** → Substance Term(s) + **Administrable Dose Form**
- PhPID Level_**L4** → Substance(s) Term+ Strength + reference strength + **Administrable Dose Form**



Global PhPID's

- Single database containing global validated PhPID
- Limit unnecessary data redundancies
- Transparent business rules based on the IDMP algorithm for PhPID
- A Global Substance ID is the key for all PhPIDs
- A global Level 3 and 4 PhPID is not possible without a global consensus on Dose Form IDs

Global vaccine initiative

- A collaboration between UMC, US-FDA, NCATS and EU-SRS
- To investigate the feasibility of having “global” substance management with regards to
 - Technical aspects, global system
 - Scientific aspects

Global vaccine initiative – technical

- Test performed with exchanging (non-confidential) vaccines data between NCATS public GSRS and UMC-SRS
- User requirements were discussed, such as, search function
- Vaccines are complex and feedback for improvements for data exchange were given to the GSRS development team
- Industry has expressed an interest in joining the discussions, especially with regards to data exchange
- Global Substance IDs (and PhPID) generated based on the vaccines in UMC-SRS

Global vaccine initiative - Scientific

Agreement on hierarchy (see next slide)

Discussions on naming

Initial comparison of vaccines in SRS (GSRS vs EUSRS)

- COVID vaccine: Tozinameran
- Meningococcal vaccine: Neisseria
- Comparison side-by-side between regions

Conclusion:

- Some differences in how vaccines were built, but no showstoppers (willingness to align)

Next steps?

- Discuss the differences in building vaccines and the possibility to align globally, and/or if regional differences are acceptable
- Discuss minimal fields for the vaccine substances to be assigned a global Substance ID
- Aim for achieving global agreement on naming of vaccines

UMC-SRS Vaccine hierarchy

Organism

Vaccine (substance)

Vaccine adjuvant
(SSG1)

Constituents

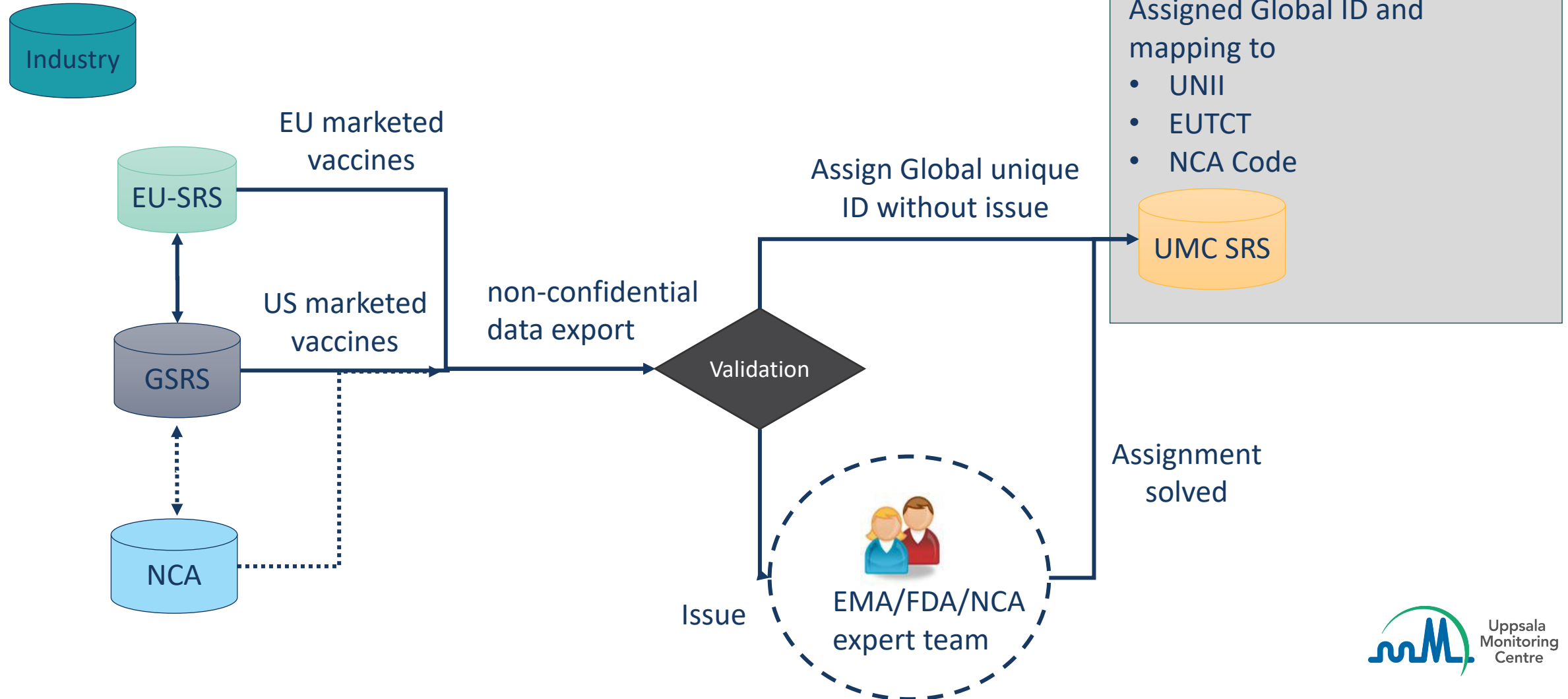
Basic vaccine hierarchy for the vaccines in UMC-SRS to be shown in the substance hierarchy.



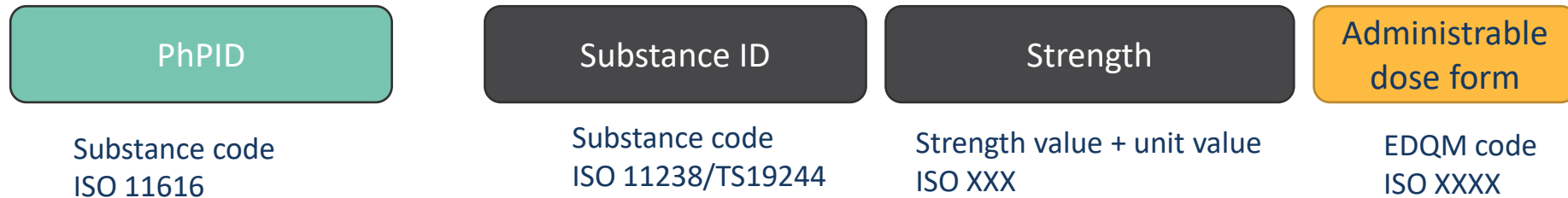
Substance Hierarchy

- 🔍 **Bordetella pertussis (Bergey et al., 1923) Moreno-Lopez, 1952, Whole TEST-OL-IP**
 - 🔍 Bordetella pertussis, Filamentous Haemagglutinin, TEST-OL-IP
 - ▷ 🔍 Bordetella pertussis, Filamentous Haemagglutinin, Inactivated-TEST-OL-IP
 - ▷ 🔍 Bordetella pertussis, Pertactin, TEST-OL
 - ▷ 🔍 Bordetella pertussis, strain Tohama 1, Pertussis Toxin, TEST-OL
 - 🔍 Bordetella pertussis, strain Tohama 1, Pertussis Toxin, Inactivated, TEST-OL
 - ▷ 🔍 Bordetella pertussis, strain 10536, Pertactin, TEST-IP

Possible Global vaccine process



Global Harmonisation of data using the ISO IDMP suite of standards

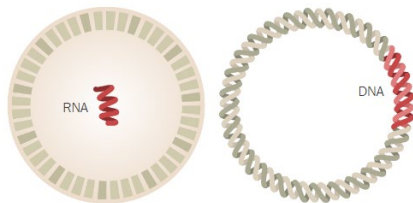


By using the ISO IDMP suite of standards PhPID's could be assigned for some COVID-19 Vaccines

Product	Substance	PhPID
Covid-19 vaccine Astra Zeneca/ Vaxzevria	COVID-19 vaccine NRVV Ad (ChAdOx1 nCoV-19)	0x26F99364EBCD42F53E5E17D89DC3A2FD
Covid-19 vaccine Moderna	COVID-19 vaccine mRNA (mRNA 1273)	0x2AAED87EBAD551B23869BD1B6C13DE0E
Pfizer COVID-19 Vaccine/ Comirnaty	Tozinameran	0xCF85D49CF8730FEBF44268D12CB115F2
Covid-19 vaccine Janssen	COVID-19 vaccine NRVV Ad26 (JNJ 78436735)	0xD21DC8D29D16257E14D6C57B0FE70157

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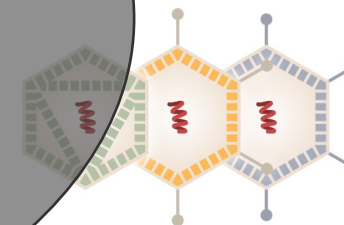
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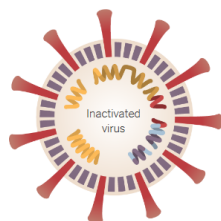
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COVID-19 vaccine inact (Vero) WIV04

Sinopharm COVID vaccine, Wuhan

Global Substance ID
 PhPID

COVID-19 product and
 substance nomenclature

Final remarks

- There is a need to work on the global scale and it is feasible
- Global Substance ID's and PhPID's can be assigned to vaccines and could improve pharmacovigilance
- The need for umbrella/grouper terms for vaccines

Questions in the Q & A
facility, please

Advancing medicine safety *together*

Uppsala Monitoring Centre (UMC)
Box 1051, SE-751 40 Uppsala, Sweden
Email: info@who-umc.org, www.who-umc.org



Vaccines - Substances

WP2: EU-SRS & SMS

On behalf of the experts in the EU-SRS team,
Annet Rozema (project leader EU-SRS)

CoE - 14/Jan/2022



TOPICS

- EU-SRS implementation project
- Cleansing (SMS) & building vaccines substances (EU-SRS)
- Vaccines naming and structure
- Collaboration

TODAY'S TOPIC

- Vaccine challenges:
 - cleansing
 - confidentiality
 - vaccine naming
- This is all very relevant to the substance experts working on vaccines in the EU-SRS implementation project

EU-SRS IMPLEMENTATION PROJECT

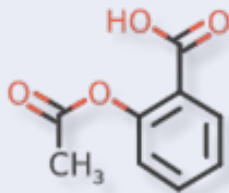
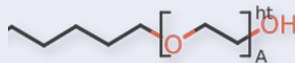


EU-SRS IMPLEMENTATION PROJECT

- GSRS software is implemented in EU:
 - In line with ISO IDMP
 - Open source software
 - Describe substances in a scientifically sound manner
 - Substance class-specific data capturing
- Key components of the project:
 - Cleansing SMS substance data (SVG provides feedback to EMA)
 - Building substances in EU-SRS, as well as enriching substances
 - Software installation & validation
 - Establish Substance Validation Group
 - Guidance documentation & procedures available (master data management)



IDENTIFYING ELEMENTS PER SUBSTANCE TYPE

Substance type	Chemical	Polymer	Protein	Nucleic acid	Structurally diverse												
Defined by	Chemical structure	Structural repeat unit(s)	Amino acid sequence(s)	Nucleobase sequence	Taxonomic information + part												
Example			<div>>A35X00TA2K RCPGCGQGVQAGCPGGCVEE EDGGSPAEGCAEAEGLRRE GQECGVYTPNCAPGLQCHPP ...</div>	<div>>303159CVH9 TAAACGTTATAACGTT ATGACGTCAT</div>	<table><tr><td>Organism Family</td><td>CANNABACEAE</td></tr><tr><td>Organism Genus</td><td>CANNABIS</td></tr><tr><td>Organism Species</td><td>SATIVA</td></tr><tr><td>Author</td><td>L.</td></tr><tr><td>Infraspecific Type</td><td>SUBSPECIES</td></tr><tr><td>Infraspecific Name</td><td>SUBSP. SATIVA</td></tr></table>	Organism Family	CANNABACEAE	Organism Genus	CANNABIS	Organism Species	SATIVA	Author	L.	Infraspecific Type	SUBSPECIES	Infraspecific Name	SUBSP. SATIVA
Organism Family	CANNABACEAE																
Organism Genus	CANNABIS																
Organism Species	SATIVA																
Author	L.																
Infraspecific Type	SUBSPECIES																
Infraspecific Name	SUBSP. SATIVA																

Vaccine antigens – among the most complex substances to define:

Description starts from pathogen, defined as structurally diverse

- Specific antigens can be described as e.g., proteins or polymers
- Inactivation of toxins and adsorbing to aluminum are described in SSG1

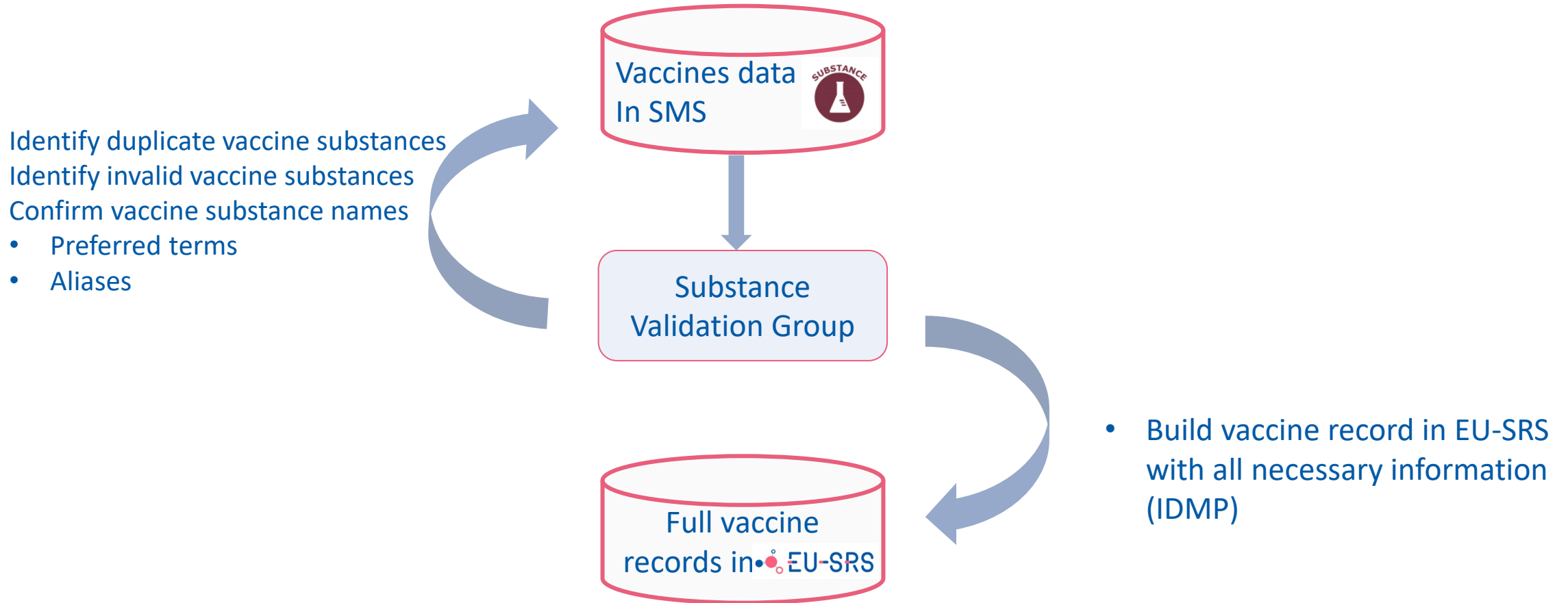
DESCRIPTION OF VACCINES IS IMPORTANT...

- Regulatory assessors benefit from insight in
 - Scientific description of vaccines
 - Relationships
 - References, identifiers
 - Naming
 - Hierarchy
 - Access to information of similar or related vaccines
- Rapporteurs across Europe will have access to the “single point of truth”
- Vaccines described through structured data explains more than many words

CLEANSING (SMS) & BUILDING VACCINE SUBSTANCES (EU-SRS)



CLEANSING + BUILDING VACCINES RECORDS



CLEANSING & BUILDING VACCINES

- ~550 unique vaccine substances received from SMS
- 66 microorganisms
 - 42% Viruses
 - 58% Bacteria
- Each microorganism contains 1 or more vaccine records + 1 or more names (PT/Aliases)
- SVG processes records per microorganism:
 1. Establish the naming rules
 2. Review SMS records
 3. Provide to SMS team
 4. Build example record in EU-SRS
 5. Document how to build the vaccines in EU-SRS
 6. Build records in EU-SRS

VACCINES SUBSTANCES - FACTS & FIGURES (SMS)

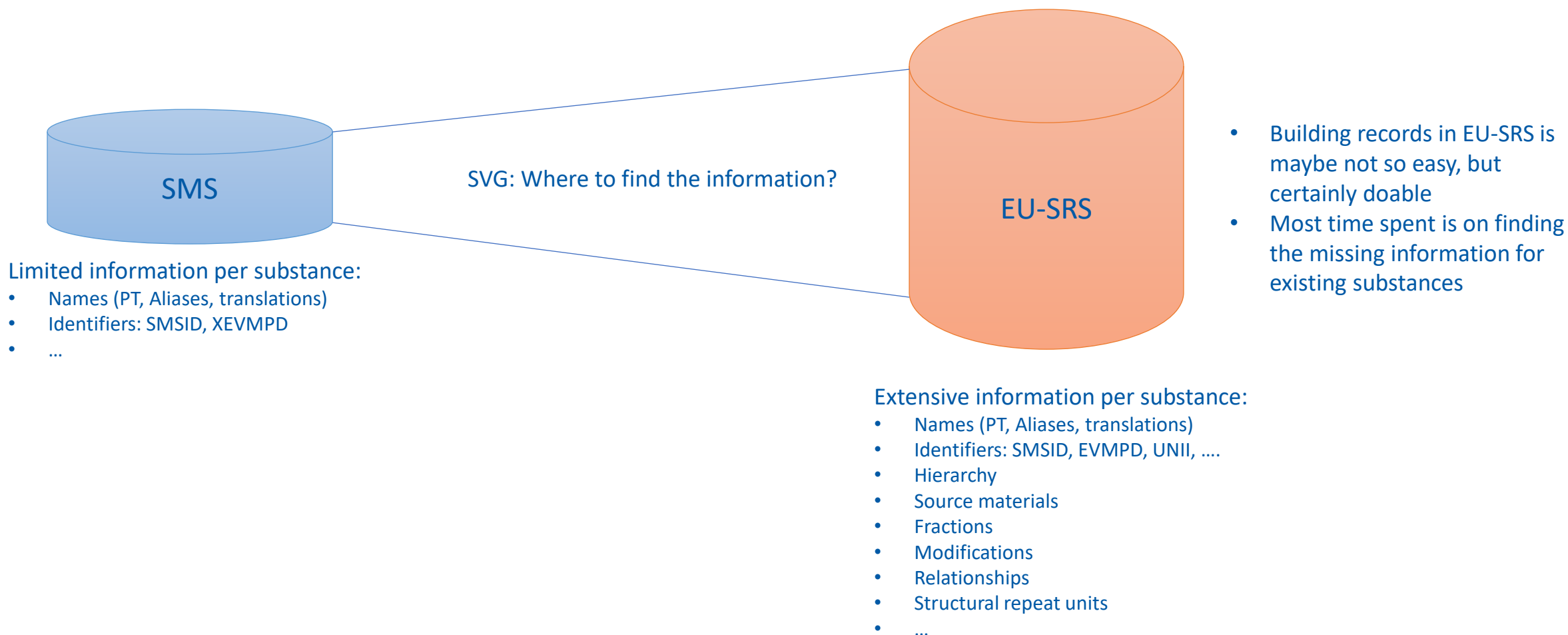
Microorganism (main categories)	Unique substances	Total number of names
Influenza virus	172	191
<i>Streptococcus agalactiae</i>	38	57
<i>Neisseria meningitidis</i>	36	69
<i>Bordetella pertussis</i>	34	40
Poliovirus	23	35
Rotavirus	19	22
<i>Haemophilus influenzae</i>	18	54
Hepatitis B virus	18	25
Hepatitis A virus	14	19

~75% of SMS records processed
Examples created in EU-SRS

Note: Significantly more changes
to SMS data requested compared
to already completed cleansing
of chemicals

Selection of microorganisms with the largest number of unique substances

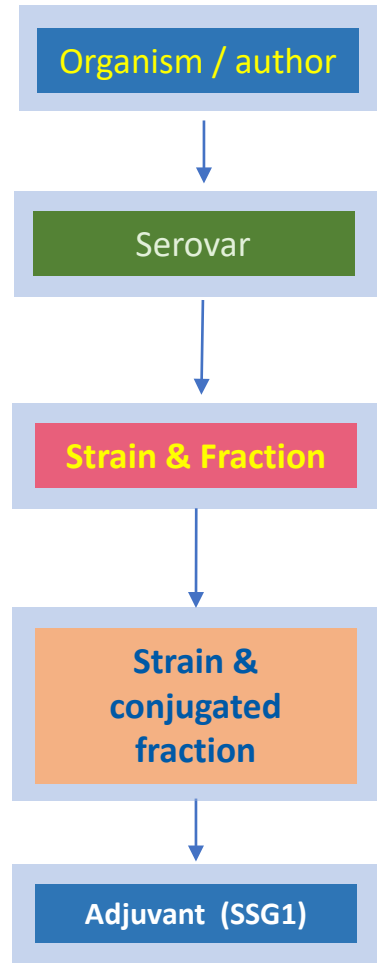
CHALLENGES WITH *CLEANSING*



VACCINES NAMING & STRUCTURE

NAMING SYNTAX OF HUMAN VACCINES BACTERIUM

Example bacterium



Neisseria meningitidis, (Albrecht and Ghon, 1901) Murray, 1929, Whole

Syntax: <Organism>,<Author>,< Year>, <Status> (Whole = Live)

Neisseria meningitidis, Serogroup A, Whole

Syntax: <Organism>,<Serogroup/ Serovar>, <Status>

Neisseria meningitidis, serogroup A, Strain C1, Polysaccharide
(Structurally diverse definition e.g. taxonomic description)

[Addition fields for polysaccharide description
by Alternative Polymer definition by intrinsic relationship of both records]

Syntax: <Organism>,<Serogroup>,<Strain>,<Fraction>

Neisseria meningitidis, serogroup A, Strain C1, Oligosaccharide

C. diphtheriae CRM197 conjugated antigen

(Structurally diverse definition e.g. strain & fraction description)

[Additional fields for the conjugated **protein description** by **Alternative Protein definition by intrinsic relationship of both records**]

Neisseria meningitidis, serogroup A, Strain C1, Oligosaccharide
C. diphtheriae CRM197 conjugated antigen, Adsorbed on Aluminum hydroxide

Syntax: <Organism>,<Serogroup>,<Strain>.<Fraction>, <Adjuvant>

NEISSERIA RECORDS IN EU-SRS

STRUCTURALLY DIVERSE



Names: Neisseria meningitidis, Whole-HD-TEST
Neisseria meningitidis (Albrecht and Ghon, 1901)...

Codes: ITIS: [ITIS- 964013](#) ↗
NCBI TAXONOMY: [NCBI-487](#) ↗

Relationships: 7

Part: **WHOLE**

Created: a year ago

Last modified: a month ago

Status: pending

Version: 11



Substance Hierarchy

▲ 🔍 Neisseria meningitidis (Albrecht and Ghon, 1901) Murray, 1929, Whole-HD-TEST	PENDING RECORD
▶ 🔍 Neisseria meningitidis, Serogroup B, Whole-HD-TEST	PENDING RECORD
🔍 Neisseria meningitidis, Serogroup C, Whole-HD-TEST	PENDING RECORD
🔍 Neisseria meningitidis, Serogroup Y, Whole-HD-TEST	PENDING RECORD
▲ 🔍 Neisseria meningitidis, Serogroup A, Whole-HD-TEST	PENDING RECORD
🔍 Neisseria meningitidis, Serogroup A, Strain C1, Polysaccharide-HD-TEST	PENDING RECORD
▲ 🔍 Neisseria meningitidis, Serogroup A, Strain C1, Oligosaccharide C.diphtheriae ...	PENDING RECORD
🔍 Neisseria meningitidis, Serogroup A, Strain C1, Oligosaccharide C.diphtheria...	PENDING RECORD

POLYSACCHARIDE RECORD

1) STRUCTURALLY DIVERSE DEFINITION

2) POLYMER DEFINITION

Display of the
Monomers

Neisseria
meningitidis,
Serogroup A, Strain
C1, Polysaccharide-
HD-TEST

pending record

Overview

Alternative Definitions 1

Names 1

Identifiers 4

Relationships 2

Modifications 1

Notes 4

Overview

Substance Class Structurally Diverse

Source Materials Class ORGANISM

Source Materials Type BACTERIUM

Source Materials State INACTIVATED

Source Materials Parent



Neisseria meningitidis, Serogroup A, Whole-
HD-TEST

Development Stage MATURE

Part CELL WALL

Fraction Material Type Polysaccharide

ALTERNATIVE
DEFINITION for
[Neisseria
meningitidis,
Serogroup A, Strain
C1, Polysaccharide-
HD-TEST]

alternative record

Overview

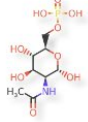
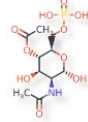
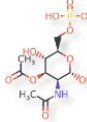
Primary Definition

Display Structure

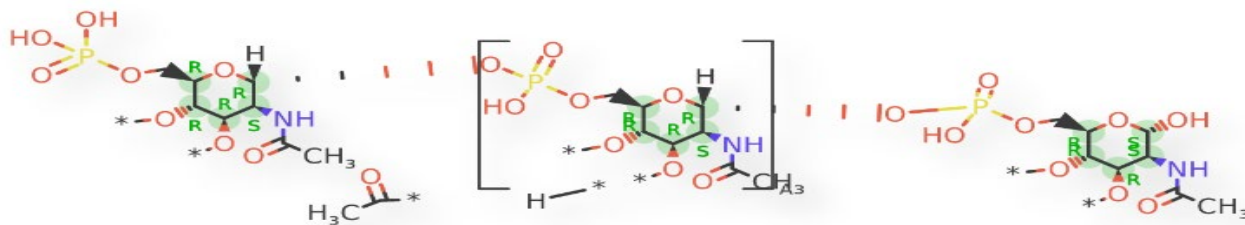
Monomers 3

Structural Units 5

Relationships 1

Material	Material Name	Amount	Type
	2-Acetamide-deoxy-D-Mannose-6-Phosphate		STARTING_MATERIAL
	4-O-acetyl-2-acetamide-2-deoxy-alpha-D-Mannose-6-phosphate		STARTING_MATERIAL
	3-O-Acetyl-2-Acetamide-2-deoxy-alpha-D-Mannose-6-Phosphate		STARTING_MATERIAL

Display of the structure of the Repeat Unit & end groups



CONJUGATED ANTIGEN RECORD 1) STRUCTURALLY DIVERSE DEFINITION

1) STRUCTURALLY DIVERSE DEFINITION

2) PROTEIN DEFINITION OF THE CONJUGATED CRM197 ANTIGEN

Neisseria meningitidis, Serogroup A, Strain C1, Oligosaccharide C.diphtheriae CRM197 Conjugated Antigen-HD-TEST
pending record

Overview

Alternative Definitions 1

Names

Identifiers 4

Specified Substances 1

Relationships 2

Audit Info

References 7

▼ Overview

Substance Class Structurally Diverse

Source Materials

Source Materials **BACTERIUM**
Type

Source Materials INACTIVATED
State

Source Materials
Parent



Neisseria meningitidis, Serogroup A, Whole- HD-TEST

Part	CELL WALL
------	-----------

Fraction Material Type	Conjugated Oligosaccharide
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Fraction Name Oligosaccharide Diphtheria CRM19
7 Conjugated Antigen

ALTERNATIVE
DEFINITION for
[*Neisseria*
meningitidis,
Serogroup A, Strain
C1, Oligosaccharide
C.diphtheriae
CRM197 Conjugated
Antigen-HD-TEST]

alternative record

Overview

Primary Definition

Subunits 1

Disulfide Links 2

▼ Overview

Substance Class	Protein
-----------------	---------

Protein Type PROTEIN

Protein Sub Type CARRIERPROTEIN

Sequence Origin Tox197 gene of corynephage b
eta DNA

Sequence Type COMPLETE

Record UNII alternative record

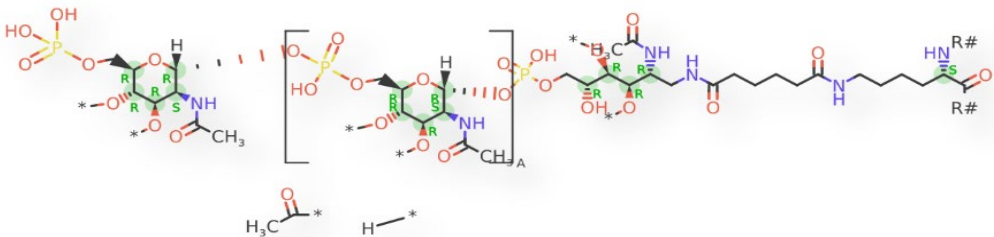
Record Status alternative

Record Version 2 v

Show Definitional
References ▼

Snapshot of partial CRM197 Subunit

Display of Modified Conjugated Oligosaccharide conlugated to Carrier protein



Subunit 1

10	20	30	40	50
GADDVVDSSK	SFVMENFSSY	HGTKPGYVDS	IQKGIQKPKS	GTQGNYYDDDW
60	70	80	90	100
KEFYSTDNKY	DAAGYSVDNE	NPLSGKAGGV	VKVTYPGLTK	VLALKVDNAE
110	120	130	140	150
TIKKELGLSL	TEPLMEQVGT	EEFIKRFGDG	ASRVVLSLPF	AEGSSSVVEYI
160	170	180	190	200
NNWEQAKALS	VELEINFETR	GKRGQDAMYE	YMAQACAGNR	VRRSYVGSLS
210	220	230	240	250

CHALLENGES WITH *CONFIDENTIALITY OF NAMES*

- In order to define the correct names, more information than the SMS names was needed → time consuming
- The explained naming conventions constrain confidential information
 - Strain name
 - In SMS, the preferred term is always public; therefore the preferred term may be in the SMS database, but not as preferred term
 - EU-SRS will be only accessible to EU regulators, so the confidentiality of names is less of an issue
 - SMS and EU-SRS will need to communicate / stay in synch
 - The team is currently investigating the best solution

COLLABORATION



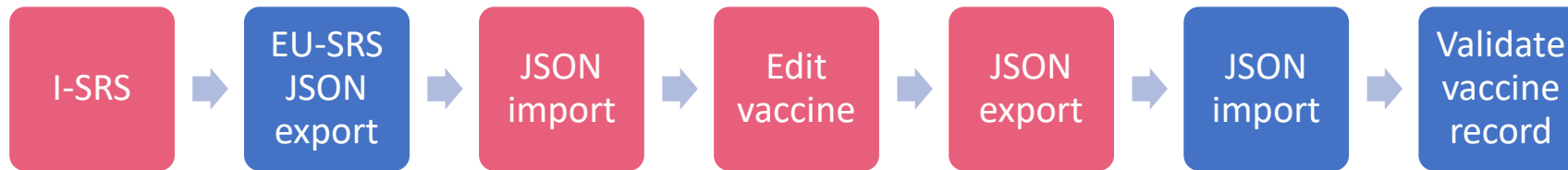
EU-SRS / WP2 IS A *COLLABORATIVE* PROJECT



- SVG members come from a variety of NCA's: Assessors or substance experts (21 members)
- Both human and veterinary experts are involved
- There is a close collaboration with EMA & FDA as well as with WHO-UMC
- **Industry involvement** through Substance Work Group approach

2021: Global Vaccines Initiative

INVOLVING INDUSTRY IN BUILDING VACCINES



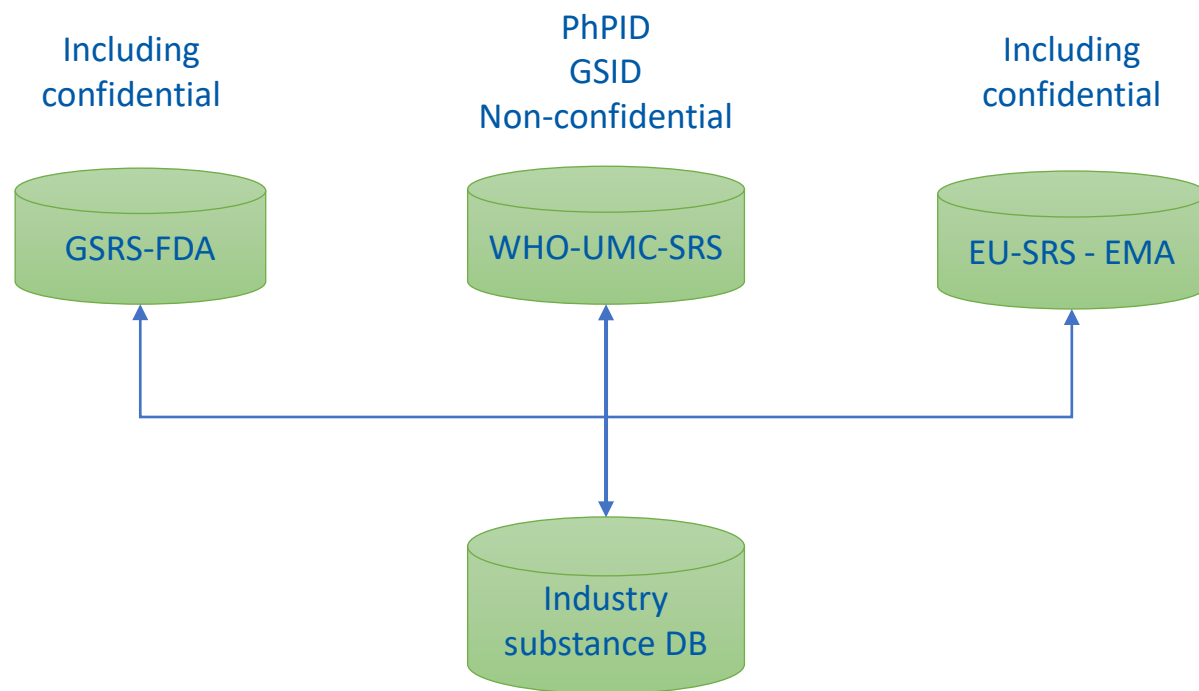
Process details:

- SVG provides guidance on how to build vaccine in the SRS software
- Relevant vaccine is exported in JSON format and shared with industry (Eudralink)
- Industry adds the necessary information based on internal knowledge and exports the updated JSON file to SVG (Eudralink)
- SVG to import the updated substance and validate (peer review) of vaccine in EU-SRS

INDUSTRY INVOLVEMENT IN BUILDING VACCINE RECORDS

- EU-SRS Steering Committee endorsed industry involvement in building vaccines in SRS (legacy/existing vaccines)
- Whether this would be a future model of exchanging data cannot be confirmed at this stage
- 3 pharmaceutical companies have indicated interest to join this activity
- This activity also fits with the discussions in the global vaccines initiative, and it is expected to provide input into GSRS software requirements for data exchange
- Start: meet early February to confirm next steps

THE FUTURE OF SUBSTANCE MANAGEMENT?

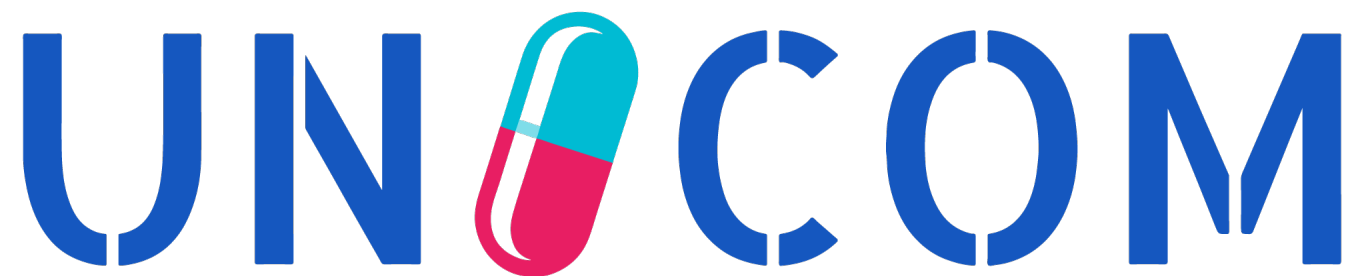


- Substance identifiers (UNII, EUTCT, GSID)
- Global approach, standards
- Same “signature fields” in each SRS
- Data exchange, staging area
- JSON file shared, validated, confirmed
- Share resources (agreed process)
- Scope differences between the instances
 - Early devt substances yes/no
 - Veterinary substances yes/no
 - Food, homeopathics



Questions in the Q & A facility, please





14 January 2022

Vaccine challenges – cleansing, confidentiality and vaccine naming

Industry perspective

Presenter: Jean-Gonzague Fontaine
jean-gonzague.x.fontaine@gsk.com

Product and Substance Master Data Lead (IDMP), GSK R&D, Vaccines BU
ISO TC 215 WG 6 « Pharmacy and Medicines Business » expert



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- ▶ Opinions and positions expressed in these slides are those of Jean-Gonzague Fontaine based on his knowledge and do not represent that of GSK.

- ▶ IDMP, a paradigm shift for pharmaceutical industry
- ▶ Product and substance identification implementation, challenges and benefits
- ▶ Industry contribution to the GSID and PhPID

IDMP, a paradigm shift for pharmaceutical industry

Pharma development and regulatory historically deal with documents
(with a notable exception of clinical – CDISC)

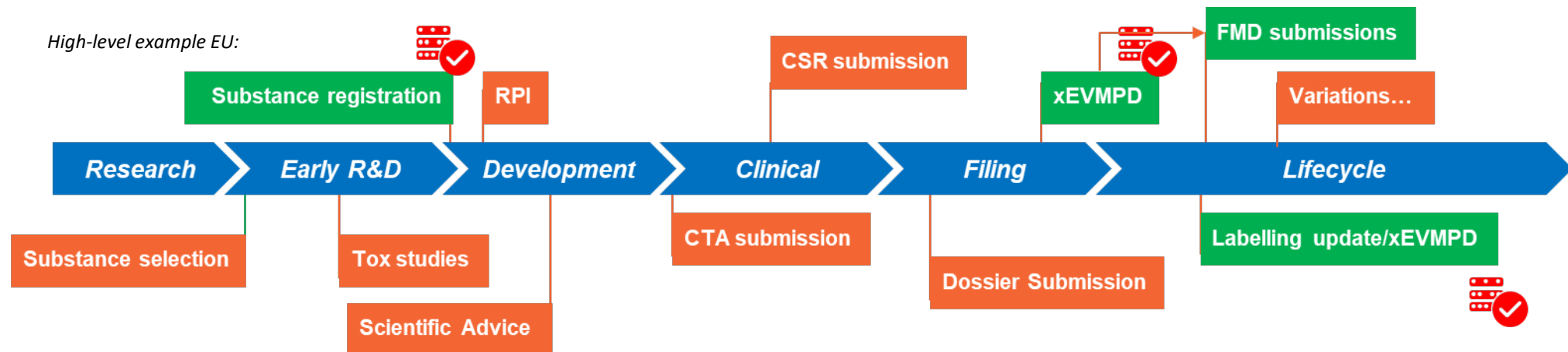
Regulatory process leverages documents such as IMPD or CTD, PSF, etc.,
which contain limited structured data in the form of tables.



Current situation

Current situation: limited structured information managed post new drug or labelling approval.

- ▶ EMA xEVMPD: structured after labelling information has been approved within 15 days from the date of authorization*
- ▶ EMA SMS: early registration, with very basic information (name)
- ▶ FDA SPL: submission of the label information within 14 days of application approval
- ▶ FDA GSRS: registration based on scientific information, for substance identification in the SPL submission



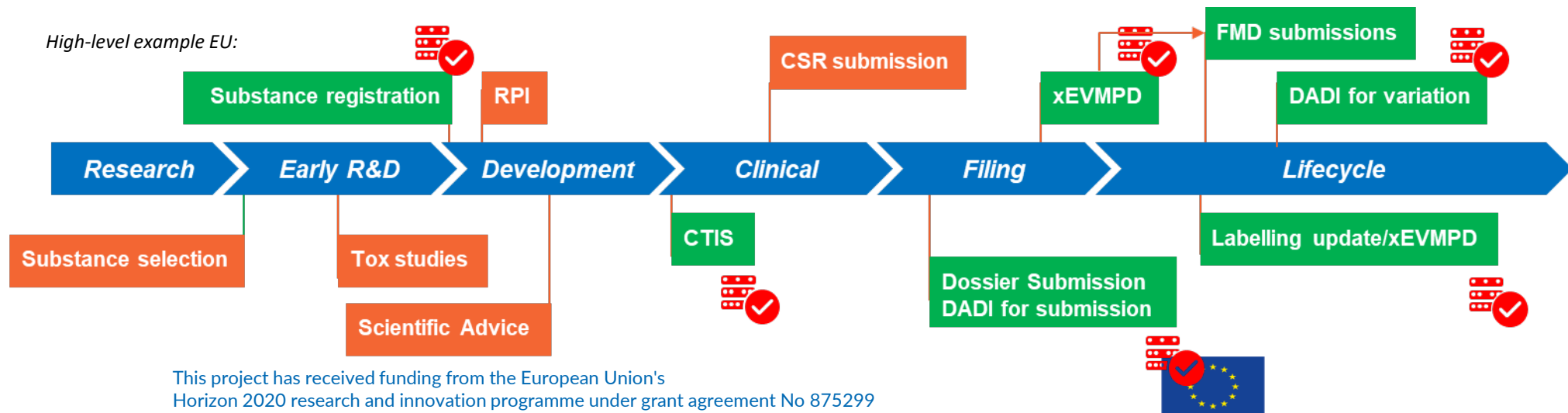
Recent developments in Europe are driving the industry towards in-process data exchanges: DADI and PMS represent a change in the approach to data submission, where data is submitted **in process, aligned/negotiated with authorities**, as opposed to post-authorization reporting.



The consequences of this are the need to manage:

- ▶ The data lifecycle: both submitted **AND** approved data for the same datapoint.
- ▶ Scientifically proven information versus regulatory truth for the same datapoint.
- ▶ Parallel, desynchronized submissions between regulatory authorities.

Implies the capability to manage as data, in system what is managed today in documents.



What are the consequences for regulatory activities (EU focus)?

Additional IDs for the management and identification of the various product dimensions.



Regulatory procedure identification	EMA procedure n°	EMA procedure n°	-
PMS reference	-	PMS ID	***
Authorisation	MA Number	MA number	-
Medicinal Product	PRD code*	MPID (+ PRD code)	***
Package	-	PCID	-
Pharmaceutical Product	-	PhPID	Global PhPID**
Substance	EUTCT, SUB code	EUTCT (+ SUB code)	GSID**

*after labelling approval **managed by WHO ***proposal of a technical report for a global and persistent Product identification globally (ISO TC215 WG6)

► In process data exchange:

- ▷ Product and substance identification critical – data quality
- ▷ Improved transparency
- ▷ Improved lifecycle management and PV on products

Product and substance identification implementation, challenges and benefits



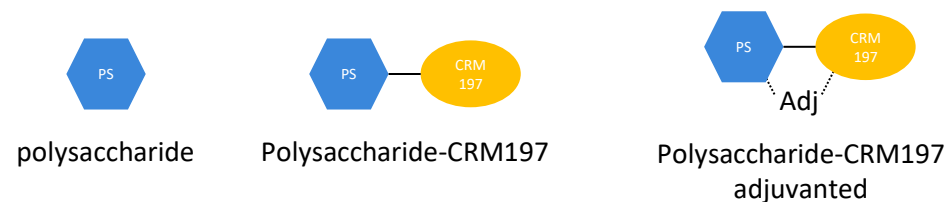
Managing data in regulatory procedures presents added challenges...

- ▶ Additional datapoints, capabilities, controls, etc.

...but also provides benefits:

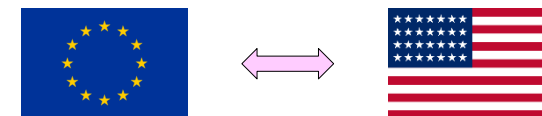
- ▶ Straight forward identification of substances (and so products):

- ▷ Do we mean the polysaccharide (active moiety), the conjugated polysaccharide, or the adjuvanted conjugated polysaccharide (active ingredient)?



- ▷ Do we mean naturally isolated polysaccharide or mechanically sized polysaccharide?

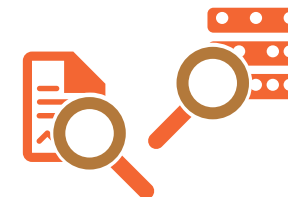
- ▶ Straightforward alignment of product across jurisdictions and reusability of information
Same GSID, same PhPID



- ▶ Regulatory processes' efficiency (Pharma and Regulator)



- ▶ Transparency: information is easily accessible and structured in a more analysable format



Implementing the IDMP substance standard presents challenges for industry:

- ▶ Additional datapoints: defined and understood by business end-users, business rules and lifecycle – risk!
- ▶ Additional capabilities: learning curve/changes for both people and systems
- ▶ Additional controls: data quality checks are a must!
- ▶ Change in mindset – context is based on a mix of system and document based info → burden for scientific functions which support highly precise antigen information
- ▶ Legacy systems, legacy standards
IDMP substance is another representation of antigens in the company
- ▶ Transversal information – first antigen description in early R&D, transferred to development, which will elucidate the complete antigen description as well as its development
- ▶ IDMP substance standard focuses on the scientific description of the antigen (incidental, stable information) where development/commercial manufacturing focus on Material Master articles and batches/lots (transactional information)
- ▶ IDMP substance standard still regulatory driven: however, the scientific knowledge is in the development functions

- ▶ Vaccine Pharmacovigilance will leverage on PhPID (and GSID) for product identification in individual case safety reports and safety databases.
 - ▷ PhPID → Transversal (jurisdiction specific) product tagging
 - ▷ GSID → deduced from the Pharmaceutical Product, enabling cross product analysis (sharing the same substance)
- ▶ Pharmaceutical Product: leverage dose form characteristics for the administrable dose form description. Improved granularity.
- ▶ GSID: Precise substance identification enables answers to transversal research questions.
For example: a regulatory authority may require the adjuvant safety profile regardless of the antigen.

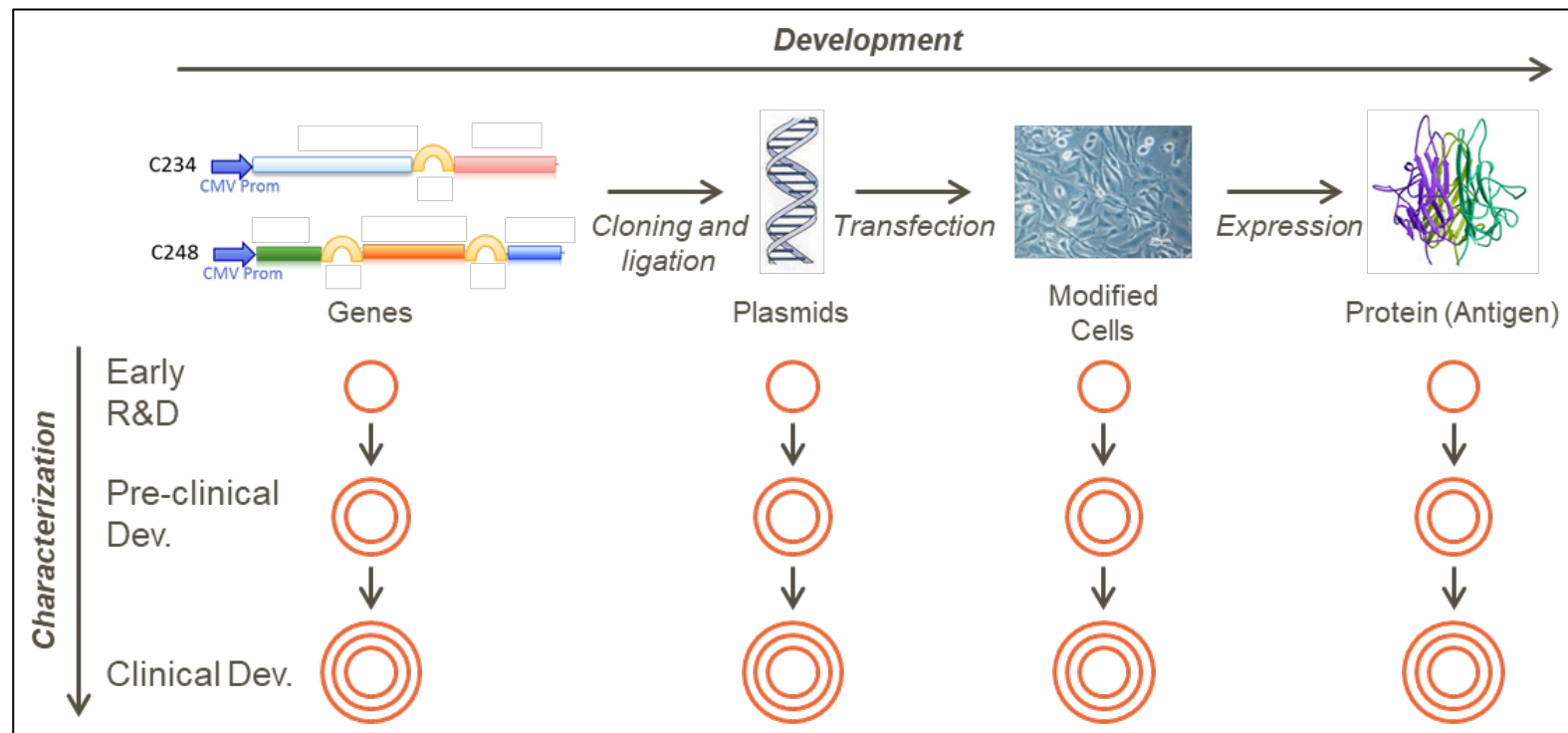
Challenges in Vaccines safety reporting:

- ▶ Reports are often too vague to identify exactly which vaccine was administered:
 - ▷ Missing Tradename (e.g. Hepatitis B vaccine)
 - ▷ No target population (10 or 20 µg? Child or Adult?)
- ▶ PhP (therefore PhPID) does not cover excipients
 - ▷ Impacts identification of side effects from excipients
- ▶ Biological substances are not identical: GSK's hep B antigen is not the same one as the Sanofi hep B antigen → GSID (or PhPID) is critical

Industry contribution to the GSID and PhPID



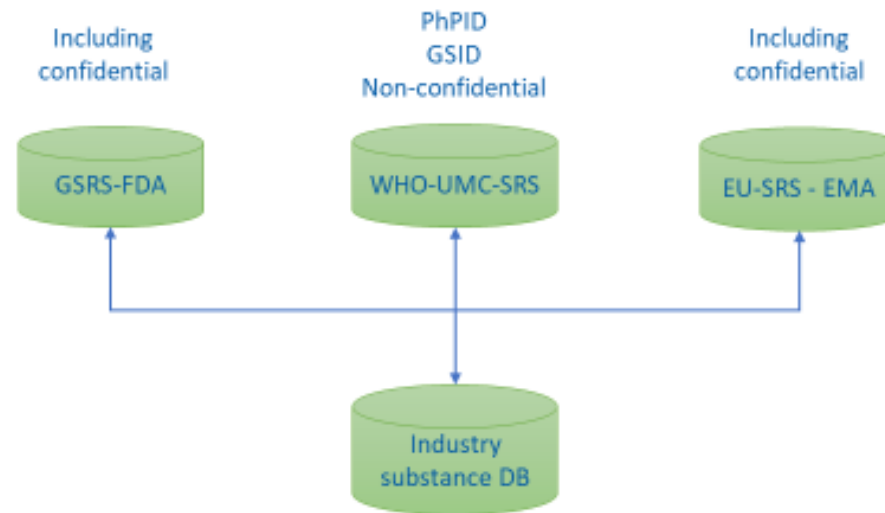
- ▶ Industry is the **originator, owner and expert on substance information and substance-defining data**
- ▶ **Industry involvement in substance creation** and sharing between regulatory authorities
- ▶ For vaccines complex substances, antigen is not always characterised at the beginning of development. **Basic substances description** (Minimum substance fields) **is required** in early phases.
- ▶ **Lifecycle management of a substance:** creation and enrichment, parent-child relationship, etc.



- ▶ Confidential information must not be shared between regulatory authorities, nor with the WHO UMC
- ▶ **Industry may act as a broker** between regulatory authorities, exchanging the same information, holding a public signature.

A PoC is currently been initiated focusing on data exchange and alignment between EU-SRS team and participating companies (voluntary basis)

THE FUTURE OF SUBSTANCE MANAGEMENT?



- Substance identifiers (UNII, EUTCT, GSID)
- Global approach, standards
- Same “signature fields” in each SRS
- Data exchange, staging area
- JSON file shared, validated, confirmed
- Share resources (agreed process)
- Scope differences between the instances
 - Early devt substances yes/no
 - Veterinary substances yes/no
 - Food, homeopathics

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► **System:** leverage GSRS system, MAH instance of SRS and EU-SRS

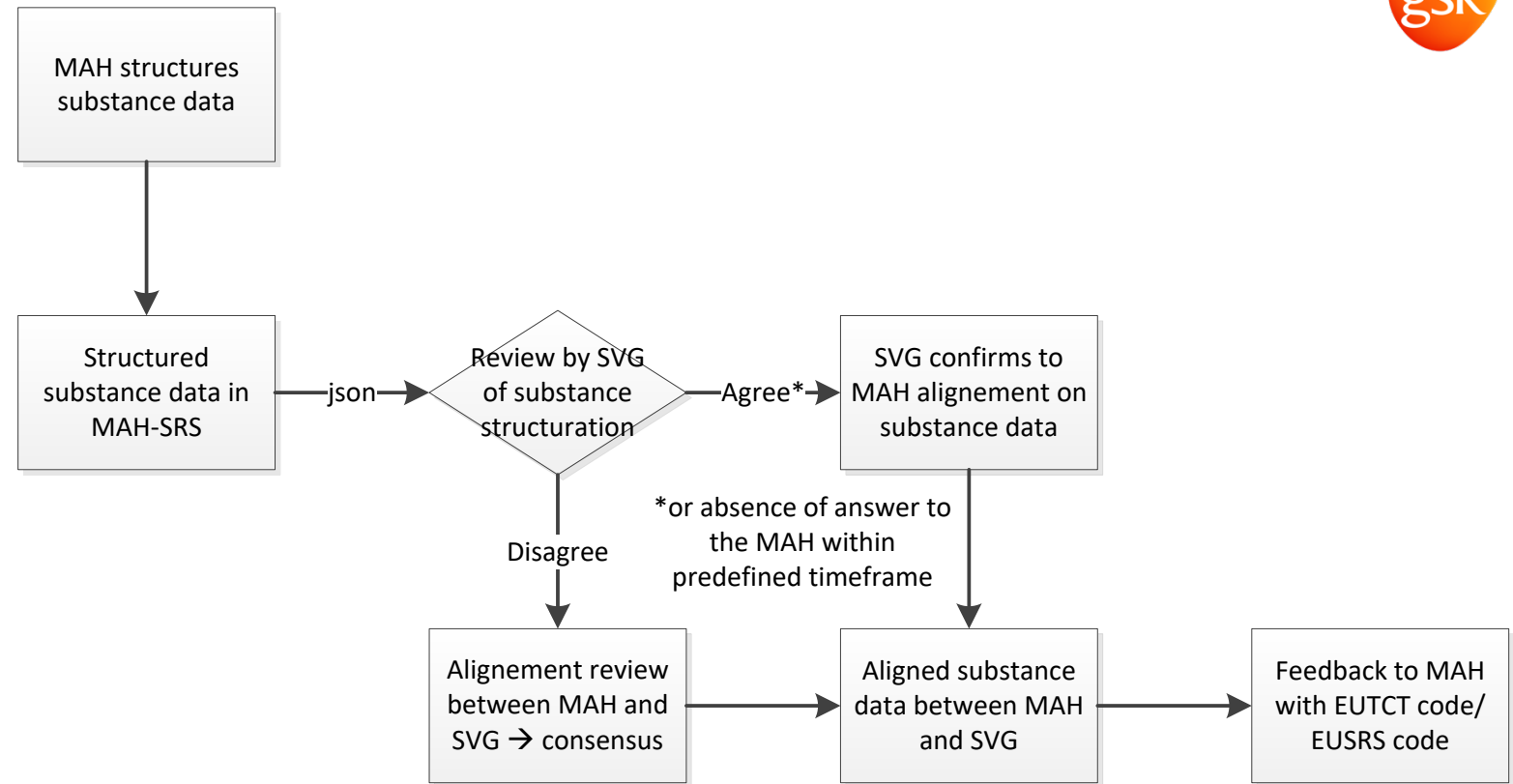
- ▷ Same system on both Regulatory and MAH sides.

► **Process:**

- ▷ Define roles and responsibilities
- ▷ Define the cleansing process →
- ▷ Define & align on business rules
- ▷ Define hash process

► **Data:**

- ▷ **Substance data ownership: MAH.**
Regulatory authority (SVG) can not modify a substance data set.
Enrichment with additional non-definitional data (e.g; references, links, additional codes, etc.)
- ▷ Data is confidential by default, except for the substance ID, substance version and substance hashmap (linked to the version)
Public data, substance ID, version and hashmap are shared with UMC



- ▶ IDMP Implementation of IDMP represents a paradigm shift for industry, with in-process data exchange in addition to documents.
- ▶ There is a need for Pharmacovigilance to define GSID and PhPID for vaccines on a global scale.
- ▶ The definition of GSID and PhPID must include the data owner – the pharmaceutical industry, as they are the expert on their own data.
- ▶ Both regulators and industry will benefit from having an univocal definition of substances and products.

Questions in the Q & A facility, please

For feedback, please go to : <https://forms.gle/SvePmpHJNwDJHFpL8>

Thanks for your time



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