



bio·logis

Genetic Information Management

# Genetic Information Management System – GIMS.pharma

Efficient translation of genetic data into clinical practice

# Who we are



# bio·logis

genetic information management

- founded 2013
- located at  
Frankfurt Innovation Center  
Germany
- 25 highly qualified team players  
with background in
  - Software Development
  - Bioinformatics
  - Human Genetics



# bio·logis

genetic information management

## Our goal:

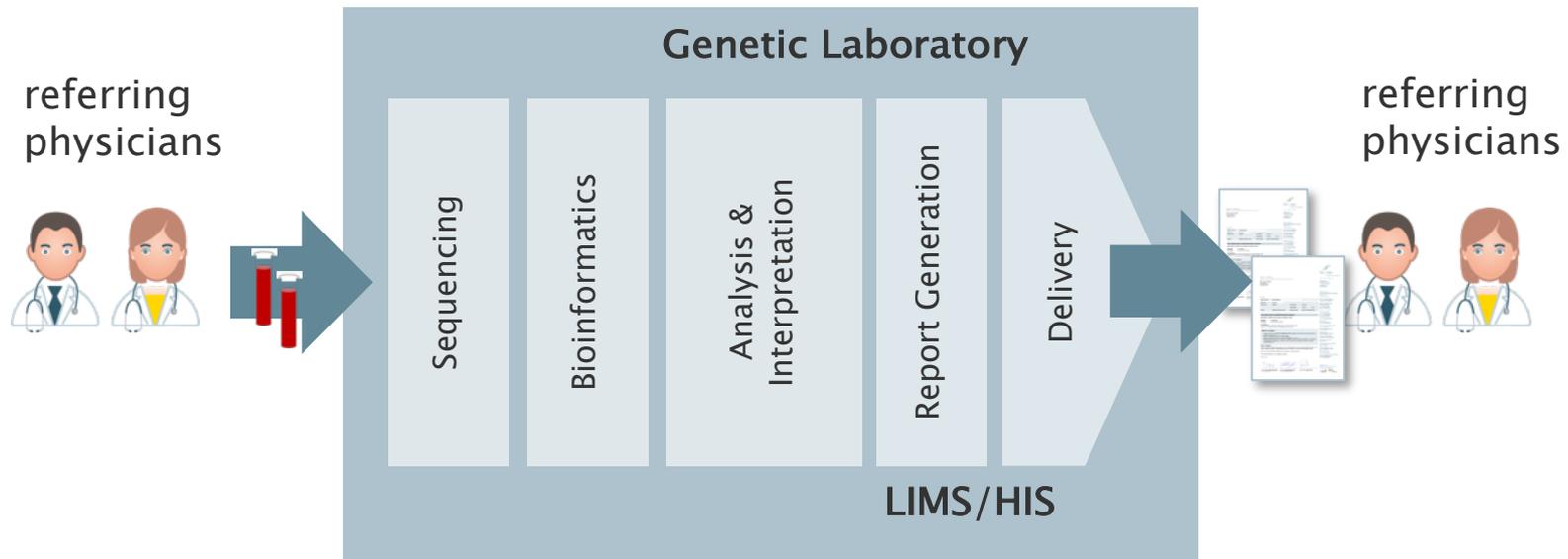
Support implementation of genetic diagnostics  
into clinical practice

## Our offer:

IT-tools and services  
for management and curation of clinically  
relevant interpretations of genetic variants.  
Creation of a virtual space for a human “Interpretome”

**We are bridging the gap  
between genetic knowledge  
and clinical implementation**

# Genetic Diagnostics: Workflow



Problem: highly fragmented manual processes consuming time & money

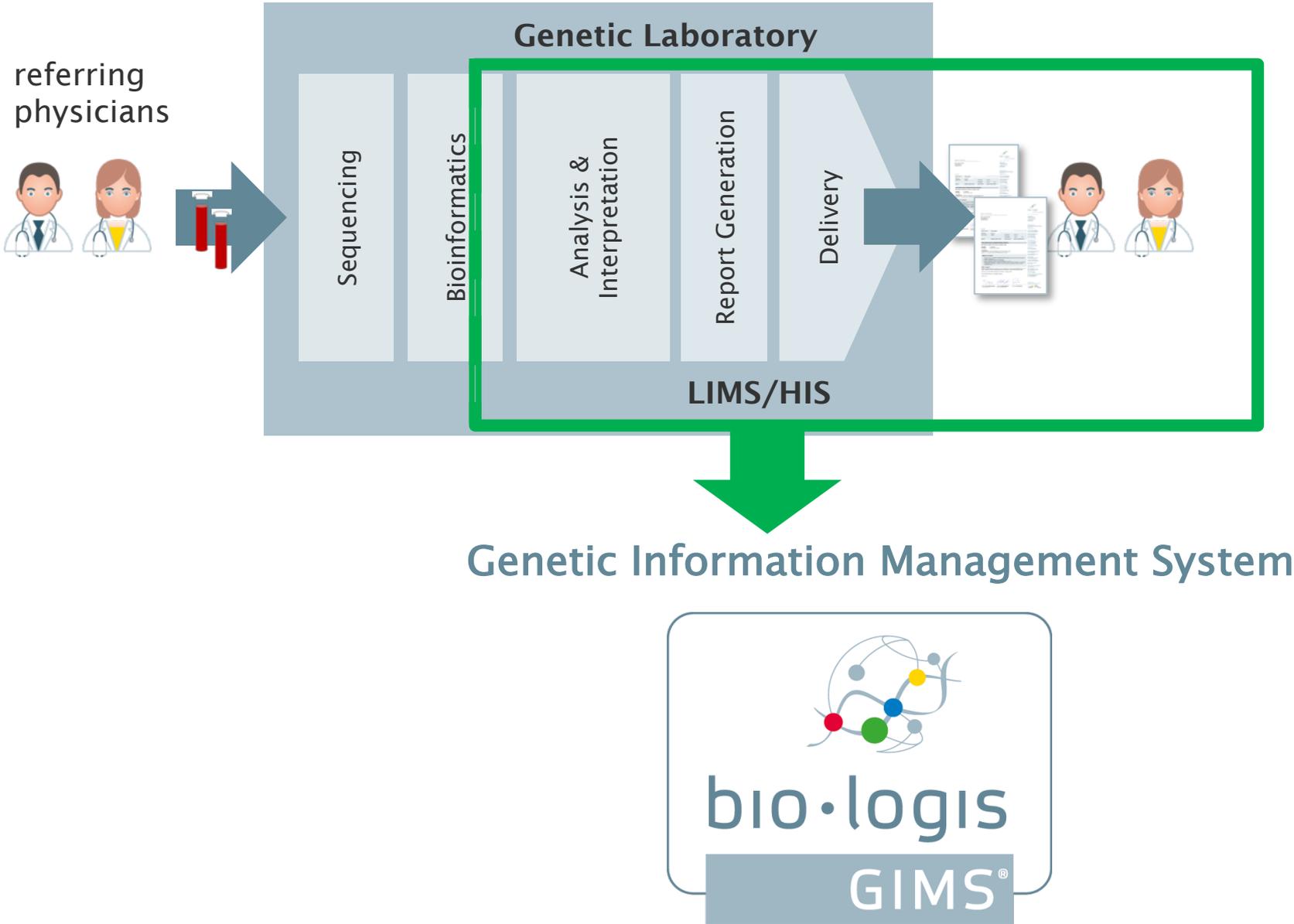


actually NO ONE workflow available for processing of complex genetic diagnostics

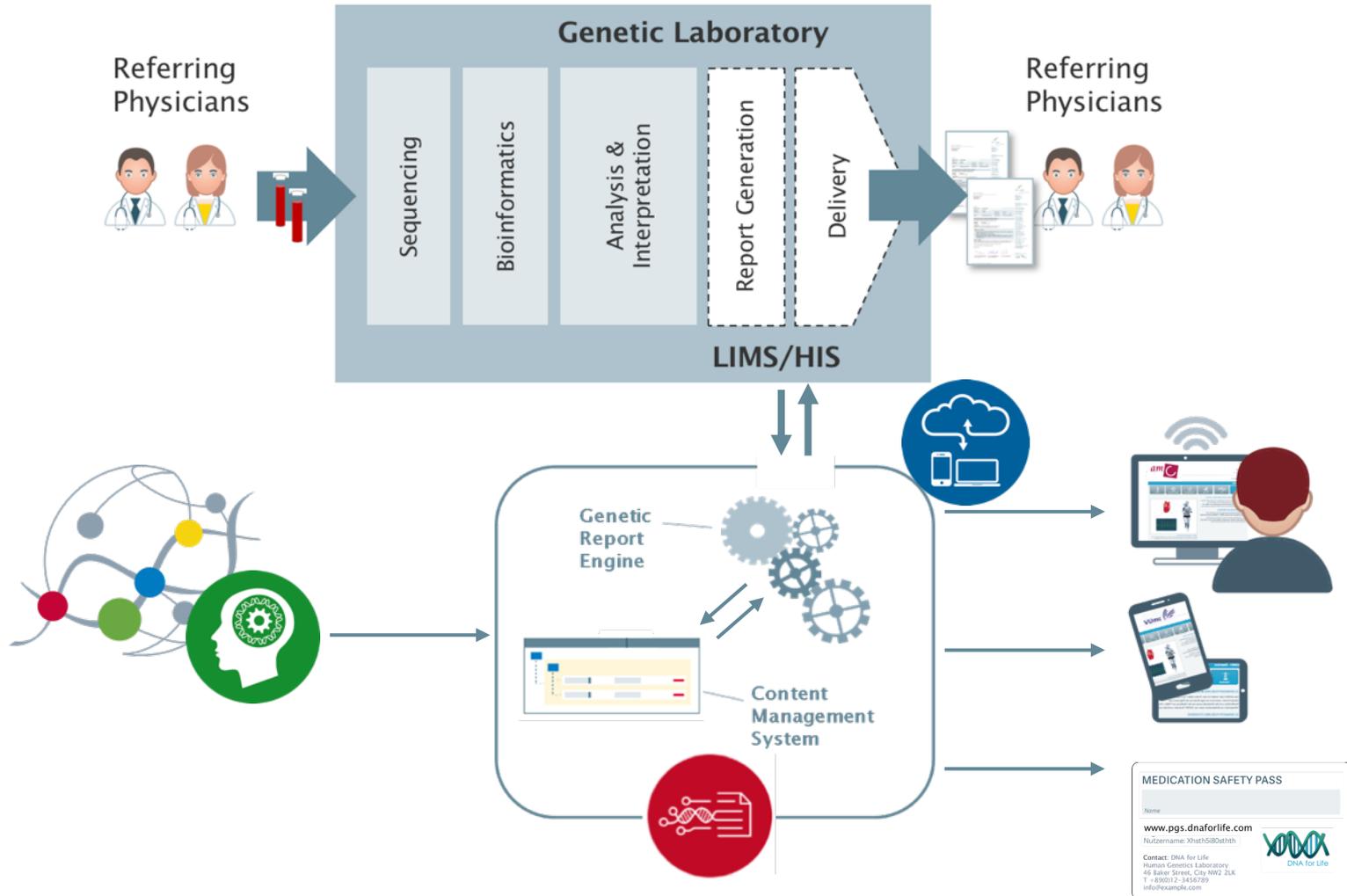


bio.logis delivers the solutions to overcome this situation stepwise

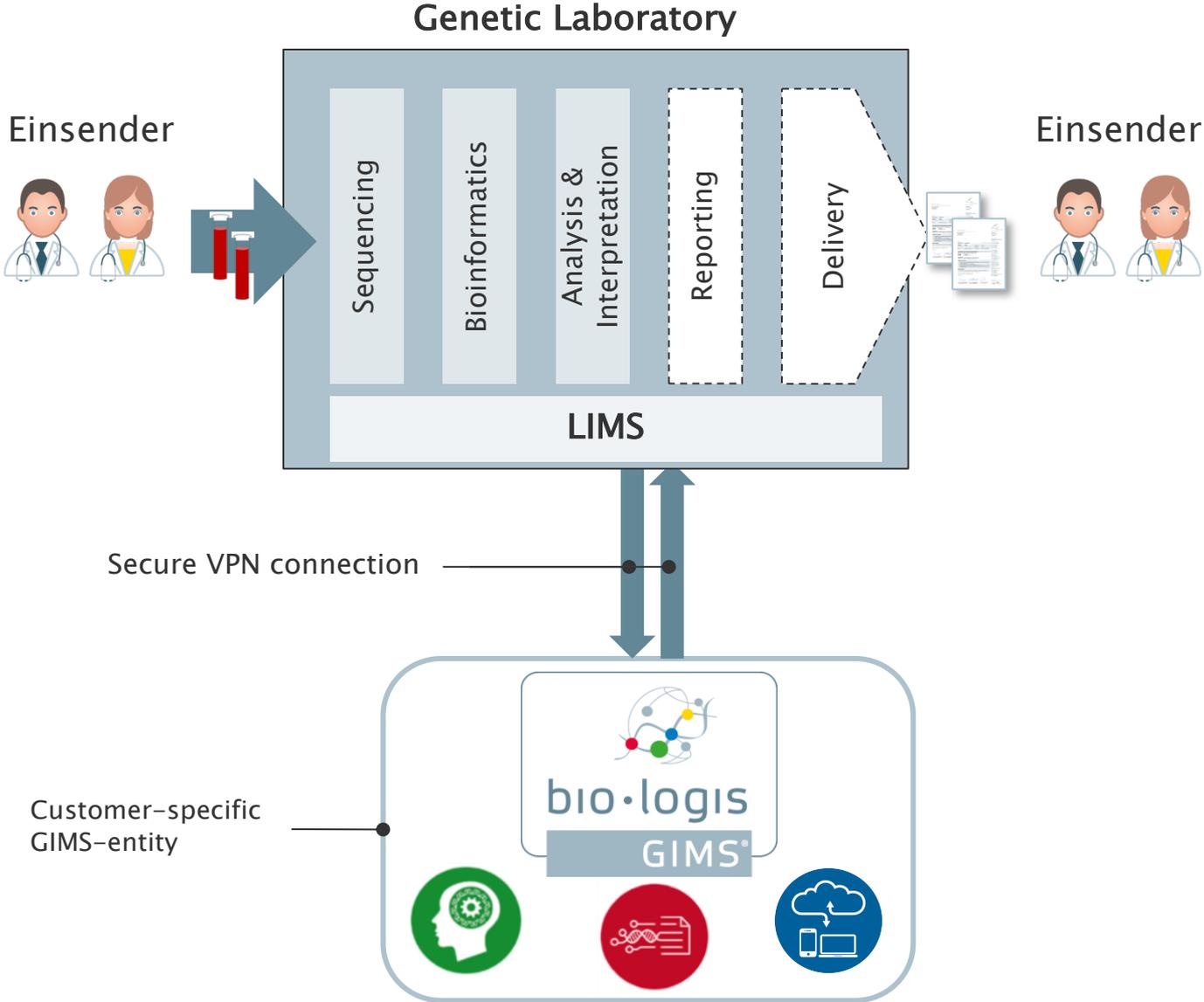
# Genetic Diagnostics: Workflow



# GIMS®: Genetic Information Management System



# GIMS Integration



First focus area:

## Pharmacogenetics

Finding the right drug and dosage for patients based on their individual genetic make-up

**Why is it not used  
in clinical practice?**

# What is needed

Efficient and standardized translation of analysis results into clinical recommendations

Digital decision support at Point of Care

# Use Case



**U-PGx** | Ubiquitous Pharmacogenomics





# U-PGx | Ubiquitous Pharmacogenomics

← → ↻ upgx.eu



U-PGx | Ubiquitous Pharmacogenomics

News / Events Participating organisations Work packages Contact

**WE WANT TO MAKE EFFECTIVE  
TREATMENT OPTIMIZATION  
ACCESSIBLE TO EVERY EUROPEAN  
CITIZEN**

TELL ME MORE

## OUR FOCUS

We want to improve the safety and efficacy of pharmacotherapy for every European patient by enabling clinical pharmacogenomics



### SHARED EUROPEAN GUIDELINES

Maintenance and dissemination of pharmacogenomics guidelines in the European Union



### IMPLEMENTATION AND EVALUATION

Clinical implementation and outcome evaluation of pre-emptive pharmacogenomics in a multitude of European countries



### ENABLING TECHNOLOGIES

Development of powerful and barrier-free clinical decision support systems and novel pharmacogenomics methodologies



### COMMUNICATION AND EDUCATION

Development of a program to reach out to patients, health care professionals, regulatory agencies, politics and health insurance organisations





- EU-funded project within the Horizon 2020 program
- Aiming to support implementation of pharmacogenomics in clinical practice
- bio.logis GIM responsible for implementing GIMS at 7 selected hospitals across Europe



Servicio Andaluz de Salud  
CONSEJERÍA DE SALUD

Univerza v Ljubljani





# U-PGx | Ubiquitous Pharmacogenomics



Dosing recommendations  
 for: 78 drugs  
 based on: ≈50 variants  
 in: 13 genes  
 available: 7 languages

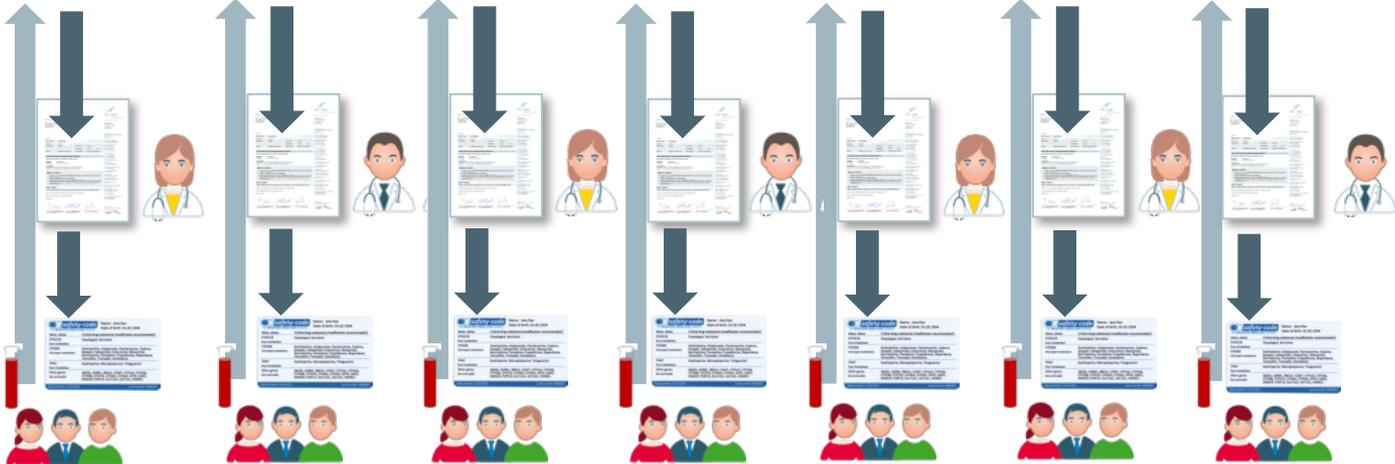


| safety-code                |   | Name: Jane Doe            |
|----------------------------|---|---------------------------|
| The Medication Safety Code |   | Date of birth: 01.02.1934 |
| Gene, status               | Critical drug substances (modification recommended!)  |                           |
| CYP2C19                    | Clopidogrel, Sertraline   |                           |
| Poor metabolizer           |   |                           |
| CYP2D6                     | Amitriptyline, Arripirazole, Clomipramine, Codeine, Doxepin, Haloperidol, Imipramine, Metoprolol, Nortriptyline, Paroxetine, Propafenone, Risperidone, Tamoxifen, Tramadol, Venlafaxine |                           |
| Ultrarapid metabolizer     |   |                           |
| TPMT                       | Azathioprine, Mercaptopurine, Thioguanine   |                           |
| Poor metabolizer           |   |                           |
| Other genes                | ABCB1, ADRB1, BRCA1, COMT, CYP1A2, CYP2A6, CYP2B6, CYP2C3, CYP3A4, CYP3A5, DPYD, G6PD, HMGCR, P2RY12, SULT1A1, UGT1A1, VKORC1   |                           |
| Not actionable             |   |                           |
| Date printed: 15.03.2016   |   |                           |
| Card number: 0000001       |   |                           |

Project Sites



Patients





## Standardized Genotyping

13 Genes ≈ 50 Variants

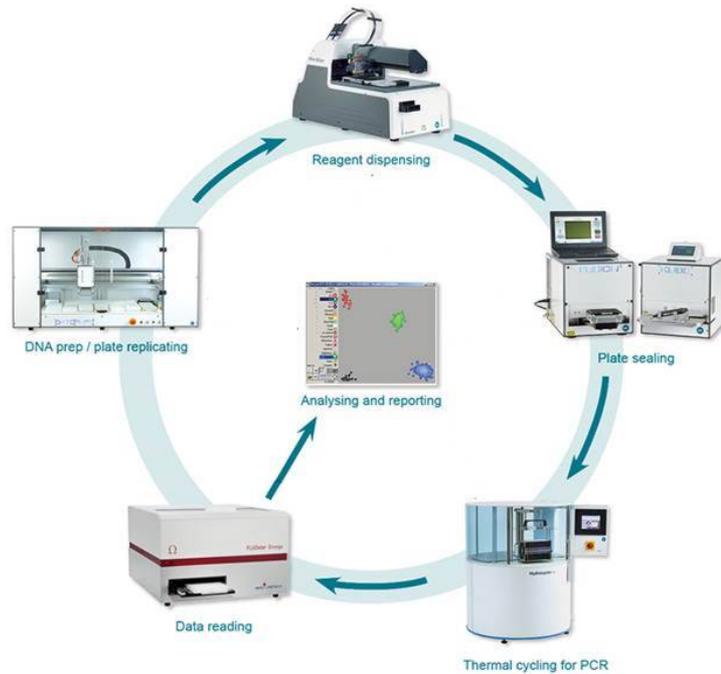


Table 1: Selected pharmacogenes and respective variants (RS number included).

| Genes   | Allele     | Major Nucleotide Variation                     | dbSNP RS ID | Effect on protein | Functional Status     |
|---------|------------|--|-------------|-------------------|-----------------------|
| CYP2B6  | *6/*9      | 516G>T   | rs3745274   | Q172H             | Decreased or Inactive |
| CYP2B6  | *4/*16     | 785A>G   | rs2279343   | K262R             | Decreased or Inactive |
| CYP2B6  | *18        | 983T>C   | rs28399499  | I328T             | Decreased or Inactive |
| CYP2C9  | *2         | 430C>T   | rs1799853   | R144C             | Decreased             |
| CYP2C9  | *3         | 1075A>C  | rs1057910   | I359L             | Decreased             |
| CYP2C9  | *5         | 1080C>G  | rs28371686  | D360E             | Decreased             |
| CYP2C9  | *11        | 1003C>T  | rs28371685  | R335W             | Decreased             |
| CYP2C19 | *2         | 681G>A   | rs4244285   | Splicing defect   | Inactive              |
| CYP2C19 | *3         | 636G>A   | rs4986893   | W212X             | Inactive              |
| CYP2C19 | *4A/B      | 1A>G   | rs28399504  | M1V               | Inactive              |
| CYP2C19 | *5         | 1297C>T  | rs56337013  | R433W             | Inactive              |
| CYP2C19 | *6         | 395G>A   | rs72552267  | R132Q             | Inactive              |
| CYP2C19 | *8         | 358T>C   | rs41291556  | W120R             | Inactive or Decreased |
| CYP2C19 | *9         | 431G>A   | rs17884712  | R144H             | Decreased             |
| CYP2C19 | *10        | 680C>T   | rs6413438   | P227L             | Decreased             |
| CYP2C19 | *17        | -806C>T <sup>3</sup>                           | rs12248560  | X                 | Increased             |
| CYP2D6  | *xN        | Gene duplication or multiplication             | X           | X                 | Increased             |
| CYP2D6  | *3         | 2549delA                                       | rs35742686  | 259Frameshift     | Inactive              |
| CYP2D6  | *4         | 1846G>A  | rs3892097   | Splicing defect   | Inactive              |
| CYP2D6  | *5         | Gene deletion                                  | X           | Gene deletion     | Inactive              |
| CYP2D6  | *6         | 1707delT                                       | rs5030655   | 118Frameshift     | Inactive              |
| CYP2D6  | *8         | 1758G>T  | rs5030865   | G169X             | Inactive              |
| CYP2D6  | *9         | 2615delAAG                                     | rs5030656   | K281 deletion     | Decreased             |
| CYP2D6  | *10        | 100C>T   | rs1065852   | P34S              | Decreased             |
| CYP2D6  | *14A/B     | 1758G>A  | rs5030865   | G169R             | Decreased             |
| CYP2D6  | *17        | 1023C>T  | rs28371706  | T107I             | Decreased             |
| CYP2D6  | *41        | 2988G>A  | rs28371725  | Splicing          | Decreased             |
| CYP3A5  | *3         | 6986A>G  | rs776746    | Splicing defect   | Inactive              |
| CYP3A5  | *6         | 14690G>A                                       | rs10264272  | Splicing defect   | Inactive              |
| CYP3A5  | *7         | 27131_27132insT<br>IVS14 + 1G>A<br>(1905+1G>A) | rs41303343  | 346Frameshift     | Inactive              |
| DPYD    | *2A        | 1691G>A  | rs3918290   | X                 | Inactive              |
| DPYD    | *13        | 1679T>G  | rs55886062  | I560S             | Inactive              |
| DPYD    | X          | 2846A>T  | rs67376798  | D949V             | Decreased             |
| DPYD    | X          | 1236G>A  | rs56038477  | E412E             | Decreased             |
| FS      | X          | 1691G>A  | rs6025      | R506Q             | Decreased             |
| HLA-B   | *5701      | T>G  | rs2395029   |                   | Tagging SNP           |
| SLCO1B1 | *5/*15/*17 | 521T>C   | rs4149056   | V174A             | Decreased             |
| TPMT    | *2         | 238G>C   | rs1800462   | A80P              | Inactive              |
| TPMT    | *3B        | 460G>A   | rs1800460   | A154T             | Inactive              |
| TPMT    | *3C        | 719A>G   | rs1142345   | Y240C             | Inactive              |
| UGT1A1  | *6         | 211(G>A)                                       | rs4148323   | G71R              | Decreased             |
| UGT1A1  | *27        | 686(C>A)                                       | rs35350960  | P229Q             | Decreased             |
| UGT1A1  | *28/*37    | A(TA)6TAA>A(TA)7TAA<br>/A(TA)8TAA              | rs8175347   | X                 | Decreased             |
| VKORC1  | X          | 1173C>T (C6484T)                               | rs9934438   |                   | Increased sensitivity |

<sup>3</sup> Position in genomic DNA sequence is used, since there is no cDNA position for this mutation.



## 78 active ingredients

### Antiarrhythmic drugs:

- Amiodarone
- Disopyramide
- Flecainide
- Kinidine
- Propafenone

### Anticoagulants:

- Acenocoumarol
- Clopidogrel
- Phenprocoumon
- Prasugrel
- Ticagrelor
- Warfarin

### Antidiabetic drugs:

- Glibenclamide
- Gliclazide
- Glimepiride
- Tolbutamide

### Antidepressants:

- Moclobemide

### NARI

- Atomoxetine

### SSRI

- Citalopram
- Duloxetine
- Escitalopram
- Fluoxetine
- Fluvoxamine
- Paroxetine
- Sertraline
- Venlafaxine

### TCA

- Amitriptyline
- Clomipramine
- Doxepin
- Imipramine
- Mirtazapine
- Nortriptyline

### Analgetics:

- Codeine
- Oxycodone
- Tramadol

### beta Blockers:

- Atenolol
- Bisoprolol
- Carvedilol
- Metoprolol
- Soltalol

### HIV therapy:

- Abacavir
- Efavirenz

### Immunotherapy:

- Azathioprine
- Tacrolimus

### Contraceptives:

- Oestrogen containing drugs

### Neuroleptics:

- Aripiprazole
- Clozapine
- Flupentixol
- Fluphenazine
- Haloperidol
- Olanzapine
- Pimozide
- Quetiapine
- Risperidone
- Zuclopenthixol

### PPIs:

- Esomeprazole
- Lansoprazole
- Omeprazole
- Pantoprazole
- Rabeprazole

### Cholesterol-lowering drugs:

- Atorvastatin
- Fluvastatin
- Simvastatin

### Tumor therapy:

- Capecitabine
- Fluorouracil
- Gefitinib
- Irinotecan
- Mercaptopurine
- Tamoxifen
- Tegafur
- Tioguanine

### Others:

- Clonidine
- Dexmethylphenidate
- Eliglustat
- Flucloxacillin
- Methylphenidate
- Phenytoin
- Voriconazole
- Siponimod

# What is needed

Efficient and standardized translation of analysis results into clinical recommendations

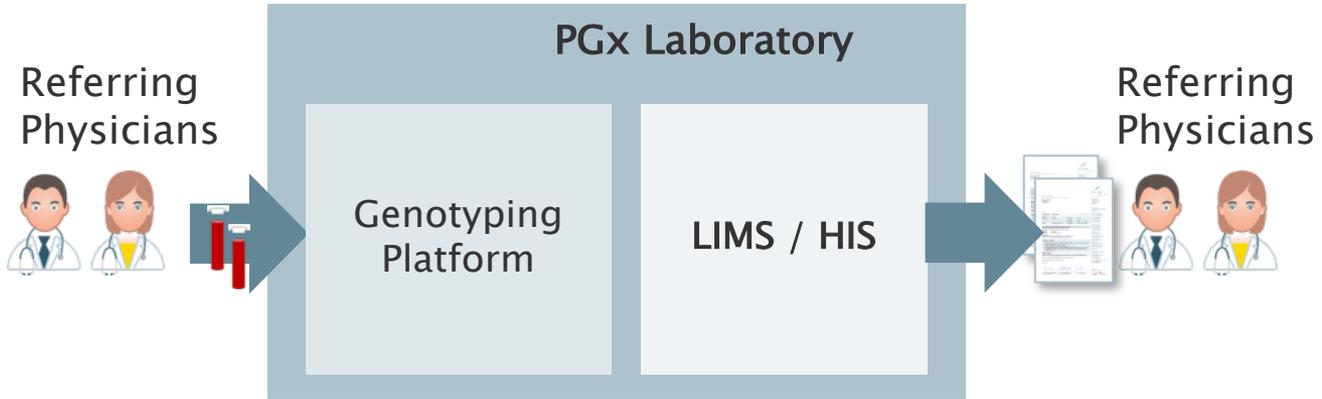
Digital decision support at Point of Care

# The solution

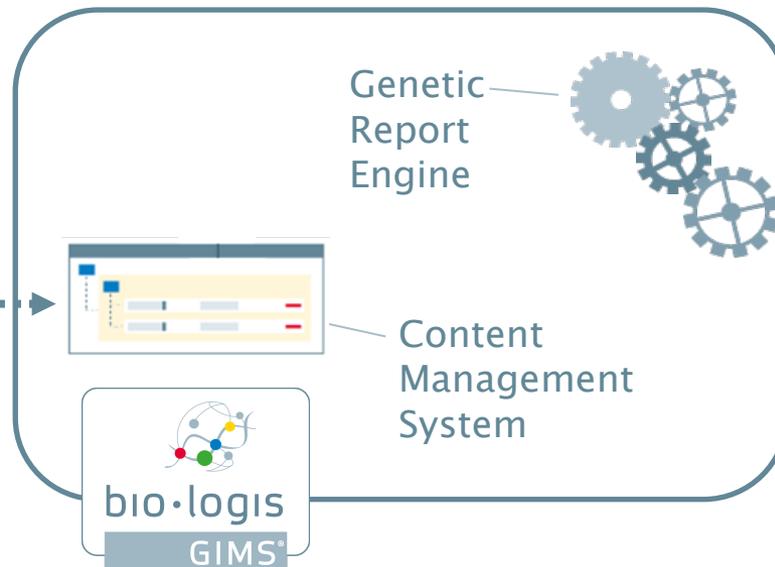




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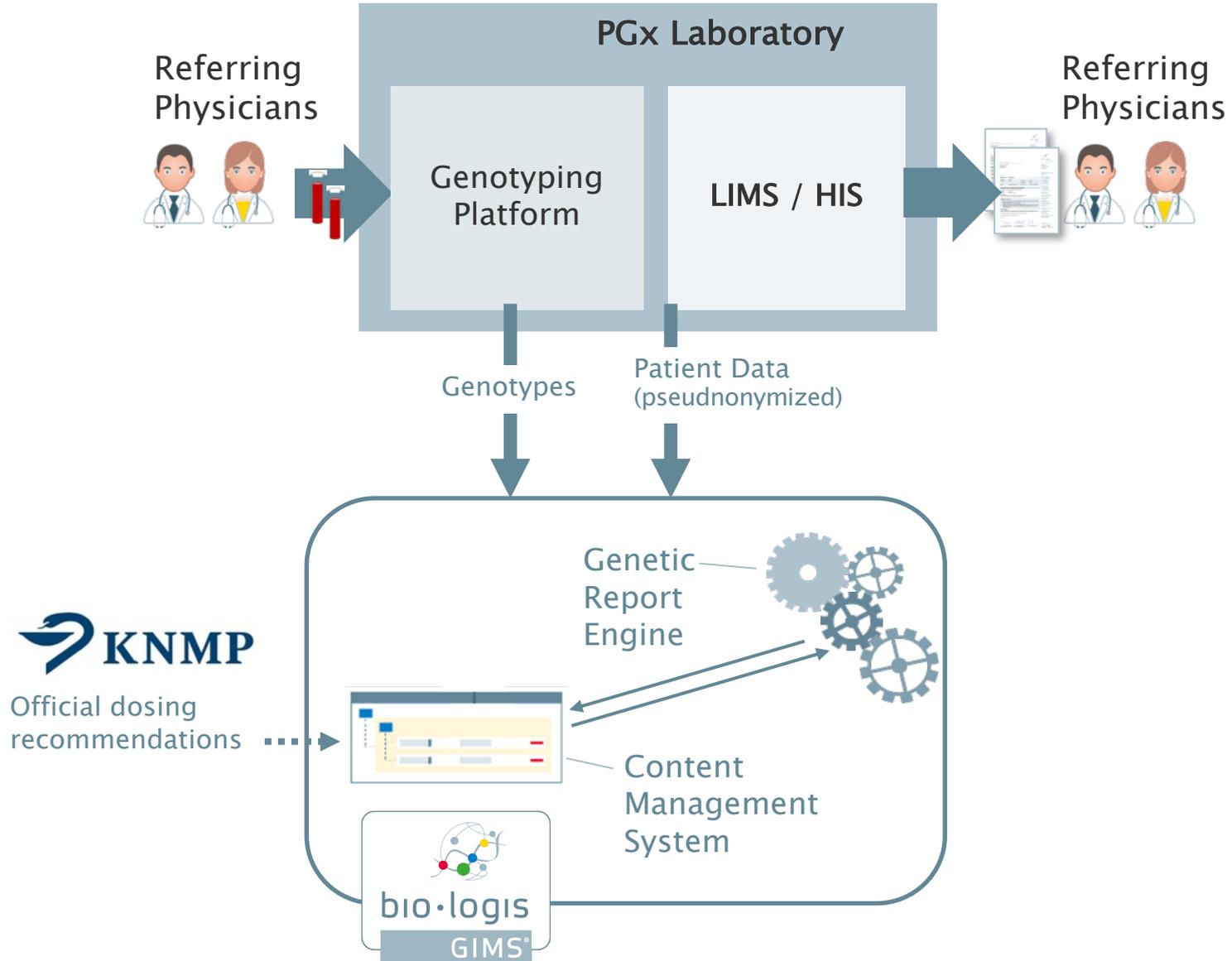


Official dosing recommendations



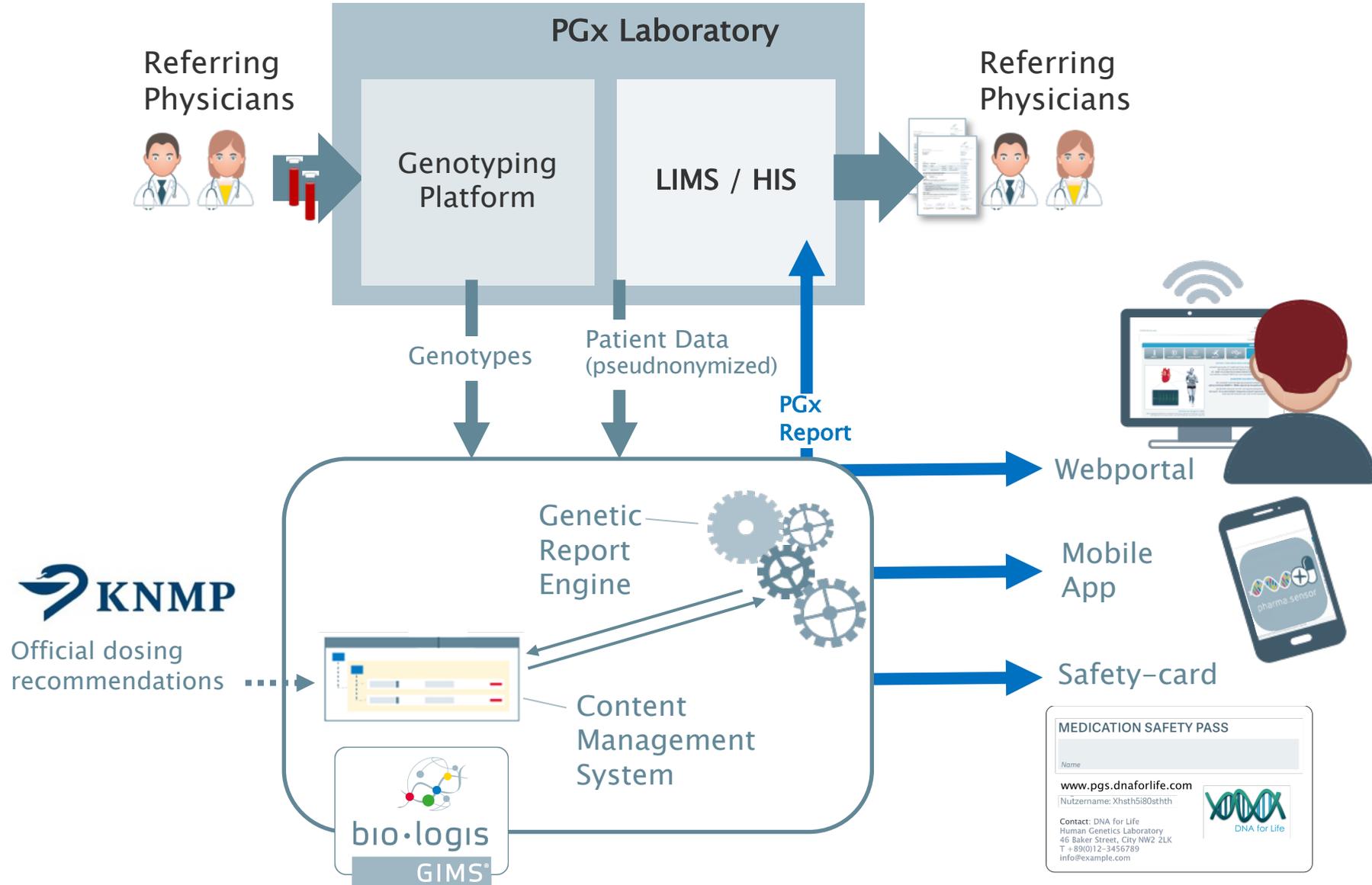


# GIMS.pharma





# GIMS.pharma



# The result: Knowledge for usage at the Point of Care, in real-time

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Wissenschaftliche Kommunikation  
Dr. rer. nat. Stephan Fees  
Dr. rer. nat. Romy Keppeler  
Dr. rer. nat. Tatjana Pabst

akkreditiert durch:  
College of American Pathologists (CAP)



Name / First  
name:

Date of birth: **24 Jan 1963**

Date of receipt: 18 Apr 2011

Sample number: 100000027

Type of sample: Saliva

Indication: Identification of genetic variants

Date of report: **11 May 2016**

## Human genetic report on clinical question: simvastatin intolerance

Analysis of *SLCO1B1* gene

### Genotype:

*SLCO1B1*\*1B/\*15

### Phenotype:

reduced transport capacity

### Interpretation:

Increased simvastatin plasma level possible owing to reduced hepatic uptake.  
Elevated risk for myopathy based on *SLCO1B1* genotype (see table 2).

### Relevance for medication:

- **Maximum daily simvastatin dose of 40 mg\*** (see table 1).
- Monitoring of creatine kinase activity indicated.  
or
- Use alternative medications (e.g. fluvastatin, pravastatin, rosuvastatin) in case of unwanted side effects (see table 1).
- Variants of *CYP2C9* gene are associated with enhanced fluvastatin plasma levels. In case of ADE under fluvastatin therapy genotyping of *CYP2C9* can be considered.

### General information:

Avoid as possible if you are taking

- statins: Co-medication with substances inhibiting *SLCO1B1*
- simvastatin: CYP3A4 inhibitors
- fluvastatin, rosuvastatin: CYP2C9 inhibitors (see table 3)

- Validated and targeted clinical recommendations based on guidelines from expert groups like e. g.
  - Clinical Pharmacogenetic Implementation Consortium (CPIC)
  - Dutch Pharmacogenetics Working Group (DPWG)



# Medication Safety Pass

## Medication Safety Pass

Name

<https://pgx-oms/webapp/>

Username: xWki3S94mFe2



Department of Genetics  
Osbridgeland Medical School  
46 Baker Street, City NW2 2LK  
T: +99 (0)999 9999  
Email: info@example.com

### PLEASE NOTE

DNA variants are often responsible for too high or low efficacy of drugs and adverse events.

For the owner of this Medication Safety Pass DNA variants have been analyzed which may be important to consider for prescribing medication



### Before prescribing:

- Check if personal recommendations are to be considered
- Detailed information available
  - In personal patient account
  - Accessible by using the QR code above



# Genetic Health Record



logged in as  
3bmw-ZQ3LP6Rc

 Log out

-  [Drug check](#)
-  [My DNA analysis](#)
-  [Tutorial pharmacogenetics](#)
-  [Support](#)
-  [Settings](#)

- [Imprint](#)
- [Terms and conditions](#)
- [Privacy policy](#)

search for drugs and active ingredients to get personal recommendations

- Simvastatin**  
active ingredient
- Simvastatin saar 10mg**  
drug
- Simvastatin saar 20mg Filmtabletten**  
drug
- Simvastatin saar 40mg Filmtabletten**  
drug
- Simvastatin ISIS 10mg**  
drug

[→ Simvastatin](#)





# Genetic Health Record



Department  
of Genetics  
Oxbridgeland  
Medical School

logged in as  
**3bmw-ZQ3LP6Rc**

 Log out

-  **Drug check**
-  My DNA analysis
-  Tutorial pharmacogenetics
-  Support
-  Settings

[Imprint](#)

[Terms and conditions](#)

[Privacy policy](#)

Your search returned matches for following active ingredients: ✓ **Simvastatin**

GENE **SLC01B1**

|   |   |  |
|---|---|--|
| ACTIVE INGREDIENT<br><b>Simvastatin</b> |  | MY DNA-VARIANT<br><b>SLC01B1 POOR FUNCTION</b> |
|---|---|--|

[view more](#)

 What's the meaning of the symbols?



# Genetic Health Record



logged in as  
3bmw-ZQ3LP6rc

Log out

Drug check

My DNA analysis

Tutorial pharmacogenetics

Support

Settings

Imprint

Terms and conditions

Privacy policy

🔍 Simvastatin

Your search returned matches for following active ingredients:  Simvastatin

GENE **SLCO1B1**

ACTIVE INGREDIENT  
**Simvastatin**



MY DNA-VARIANT  
**SLCO1B1 POOR FUNCTION**

✕ view less

**recommendation**

gene

scientific background

literature

The genetic polymorphism leads to reduced simvastatin transport to the liver. This increases simvastatin plasma concentrations and therefore the risk of myopathy.

Recommendation:

1. Choose an alternative

Consider any additional risk factors for statin-induced myopathy.

Rosuvastatin and pravastatin are influenced to a lesser extent by SLCO1B1 polymorphisms. They are also not influenced by CYP3A4 inhibitors such as amiodarone, verapamil and diltiazem.

Fluvastatin is not influenced by SLCO1B1 polymorphisms or CYP3A4 inhibitors.



This report does not replace a decision made in cooperation with a physician or genetic counselor.  
To avoid serious health risks, change of medicinal therapy should only be done under medical supervision.





# Mobile App: pharma.sensor



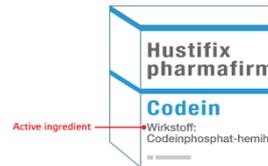
## Do I need to consider DNA variants for the drug I am taking?

Q Drug / Active ingredient

Barcode scan

### Drug Check:

- enter name in search field or
- scan barcode on packaging



### clomipramine

Gene CYP2C19

- for this active ingredient you need to consider DNA variants
- considering DNA variants, there are recommendations for
  - adjusting the dose and/or
  - prescribing an alternative therapy

### If your DNA variants are known, login here:

Login

### Or: Test DNA variants and check whether clomipramine is effective



### Anafranil 10mg Dolorgiet Dragees

Gene CYP2D6

Analysis from

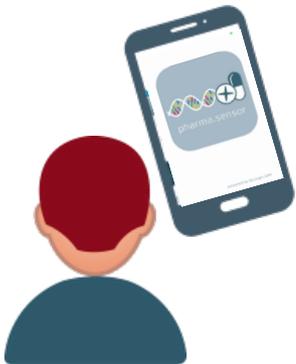


Your result (DNA-variant / phenotype): **CYP2D6 POOR METABOLIZER**

The genetic polymorphism leads to decreased metabolic capacity of CYP2D6. This may cause increased plasma concentrations of clomipramine and the active metabolite and decreased concentrations of the potentially cardiotoxic hydroxy metabolites.

Recommendation:

- Indication DEPRESSION:
  1. decrease the dose to 50% of the standard dose





# Certified as Medical Device & GDPR-proof



Mike Peter  
Sachverständiger für Datenschutz  
BDSF  
MITGLIED



[yourprivacyfirst.de](https://yourprivacyfirst.de)

## Bestätigung / Confirmation

Ich bestätige hiermit / I hereby confirm that

Firma / Company

**bio.logis Genetic Information Management GmbH**

**die Anforderungen der DSGVO /  
the requirements of the GDPR**

umgesetzt hat und sie weiterhin evaluiert /  
has implemented and continues to evaluate them

Landau, 25.04.2019

  
Unterschrift

Mike Peter  
Sachverständiger für Datenschutz  
BDSF  
MITGLIED

# translating DNA into health

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