

Quantitative Benefit-risk assessment using MultiCriteria Decision Analysis (MCDA) and its extensions: practical applications

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Preliminary information

- All the programs to reproduce the results of this presentation will be available on the PSI website
- The R code presented here is intended to be simple and understood by all (more efficient programming ways are certainly possible)

Benefit-Risk assessment

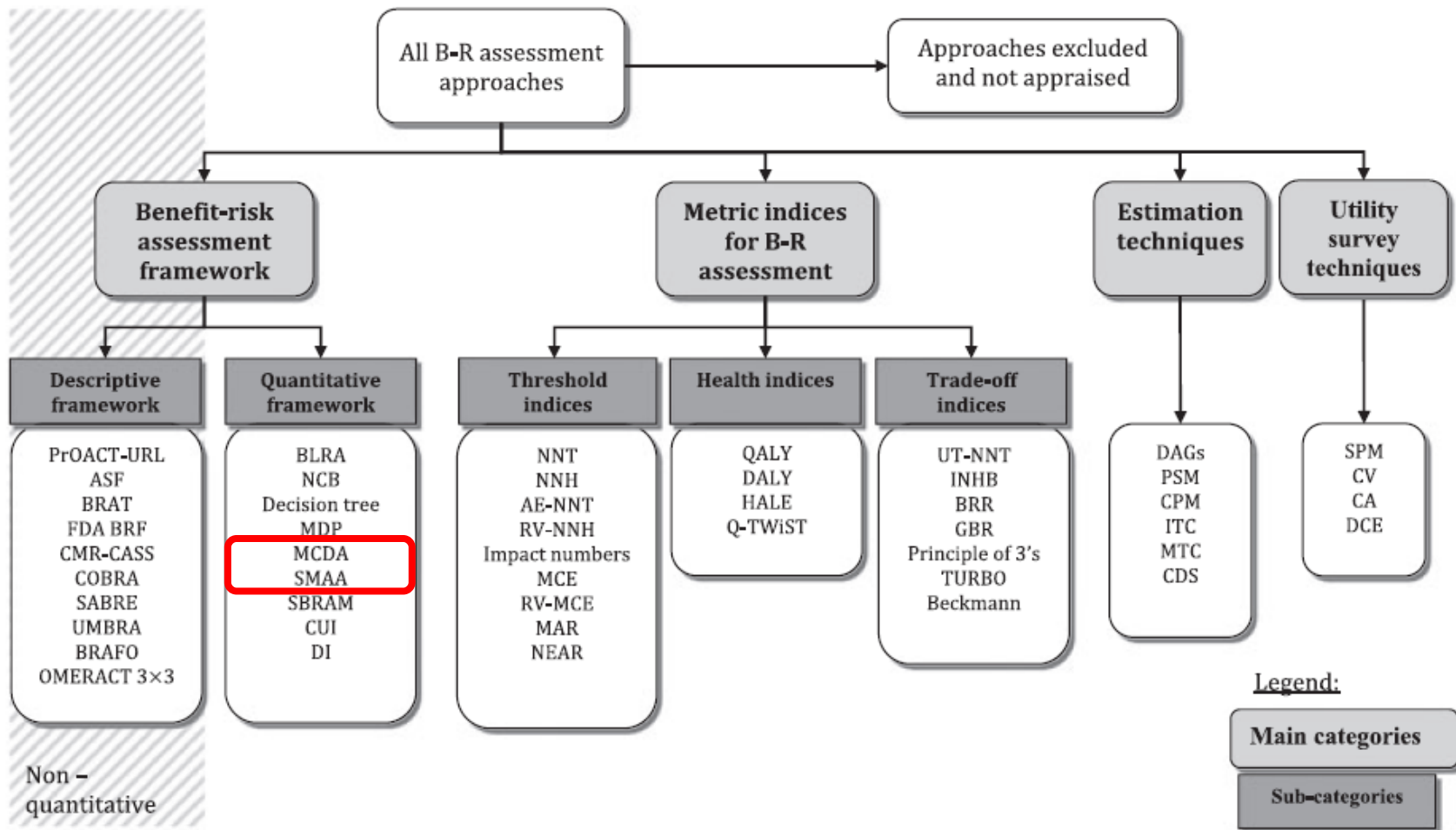
Introduction

- **Benefit-risk assessment:** to compare the benefits and the risks of a treatment
- A medicine should be considered only if it has a favorable benefit-risk balance → **Strong predictor for regulatory approval and long-term viability of a medicine**
- Until 2010, most of the drug benefit-risk assessments were qualitative
- Since then, **structured qualitative frameworks** and **quantitative methods** for benefit-risk assessment were developed
→ **more transparency, consistency and better communication**

Main working groups: PhRMA (Pharmaceutical Research and Manufacturers of America) / EMA (European Medicines Agency) / IMI-PROTECT (Innovative Medicines Initiative - Pharmacoepidemiological Research on Outcomes of Therapeutics by a European Consortium) / EFSPi (European Federation of Statisticians in the Pharmaceutical Industry) / PSI (Statisticians in the Pharmaceutical Industry)

Methodology review

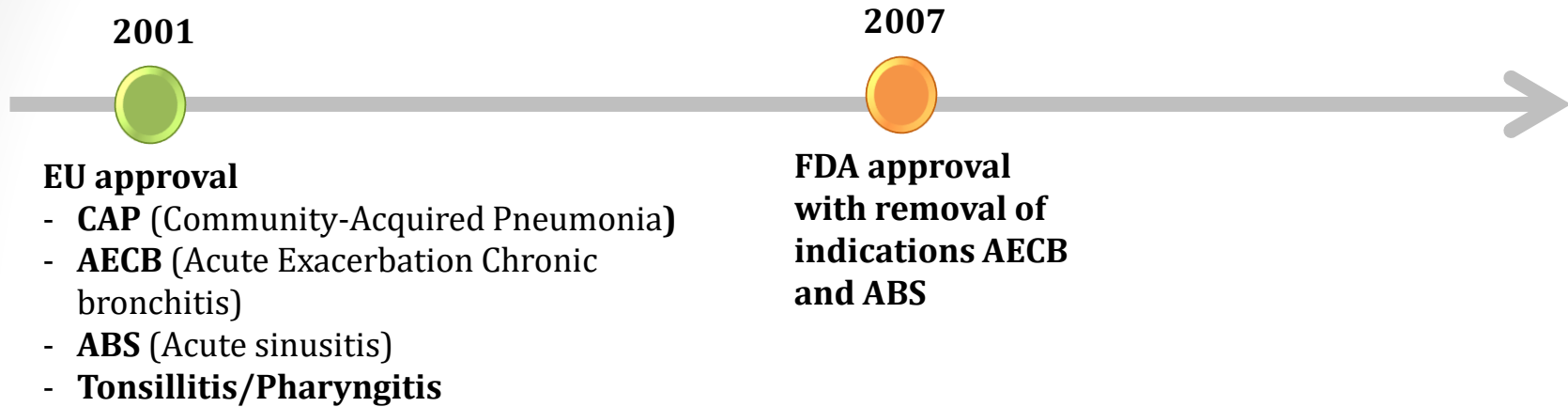
For benefit-risk assessment



Source: Mt-Isa 2014

Motivating example: Telithromycin (Ketek[®])

IMI PROTECT case study

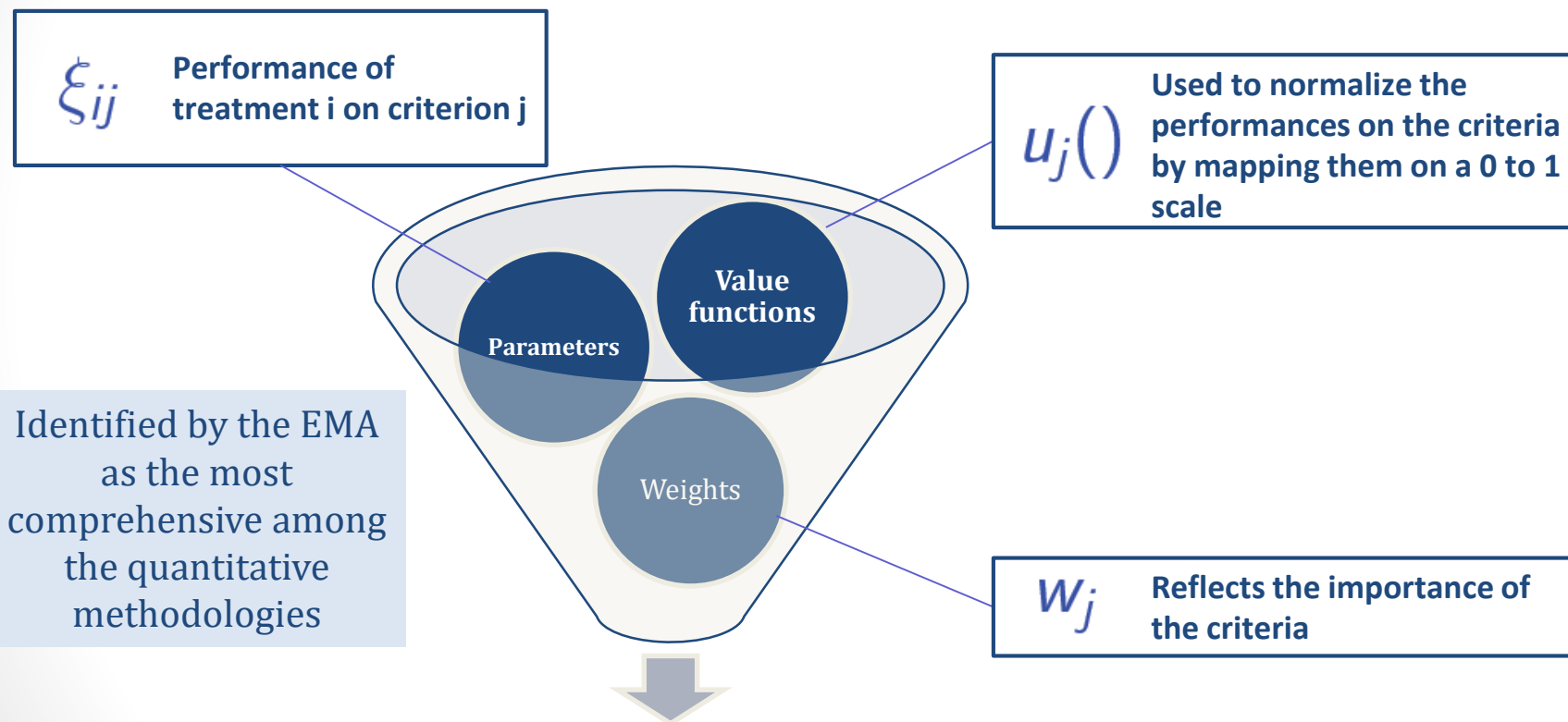


Compared to other macrolides, Telithromycin seems to be associated with a somewhat different risk profile including the following adverse reactions (eye disorders, loss of consciousness, acute liver failure, prolonged QT interval).

We will illustrate the use of quantitative approaches for benefit-risk assessment on telithromycin's CAP indication

Multi-Criteria Decision Analysis (MCDA) and its extensions

Principle



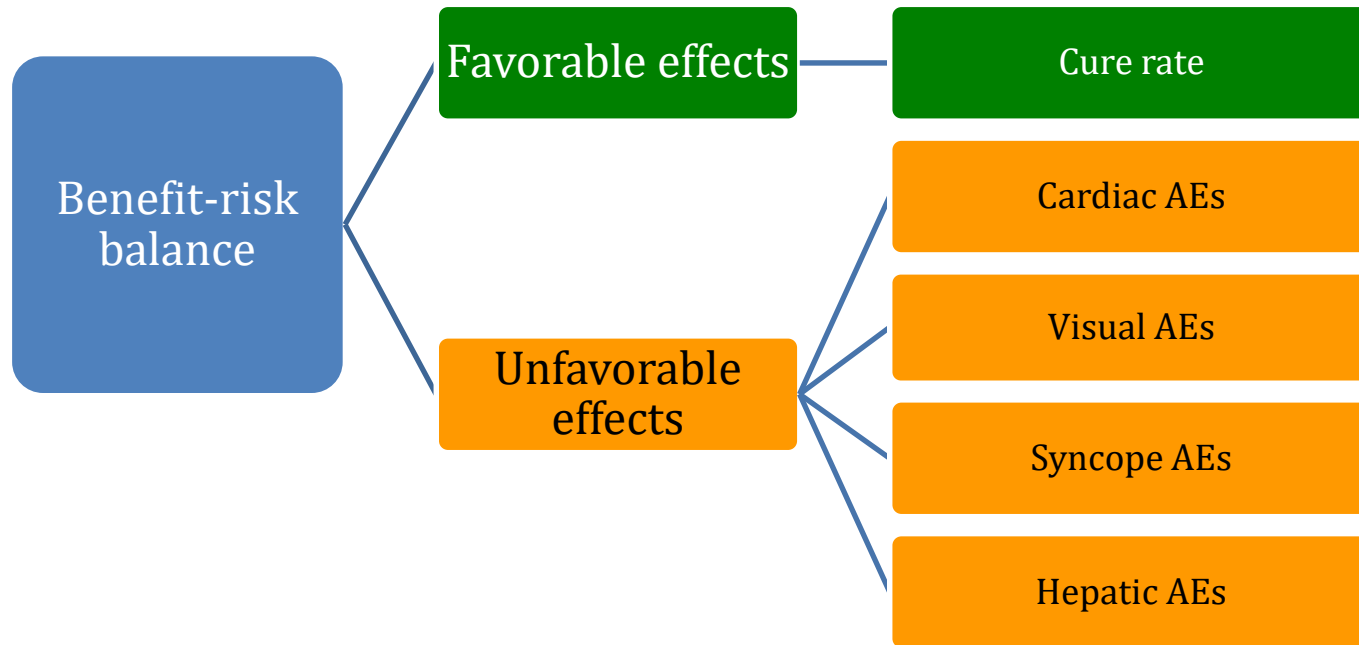
Benefit-Risk utility score

$$u(\xi_i, w) = w_1 u_1(\xi_{i1}) + \dots + w_n u_n(\xi_{in})$$

Motivating example: Telithromycin (Ketek[®])

IMI PROTECT case study – Indication CAP

Value Tree



AE = Adverse Event

Motivating example: Telithromycin (Ketek[®])

IMI PROTECT case study – Indication CAP

- 2 treatments: Ketek & Comparator
- 5 criteria: 1 for favorable effects, 4 for unfavorable effects

Criteria		Ketek [®]		Comparator	
		n/N	ξ_{1j}	n/N	ξ_{2j}
Favorable effects	Cure rate	2185/2417	90%	813/926	87,8%
Unfavorable effects	Hepatic AEs	57/1320	4,3%	46/1121	4,1%
	Cardiac AEs	4/1320	0,3%	3/1121	0,3%
	Visual AEs	14/1320	1,1%	5/1121	0,4%
	Syncope AEs	2/1320	0,2%	3/1121	0,3%

Data coming from the EPAR.

n = number of events ; N = number of patients ; EPAR = European Public Assessment Report

Different models

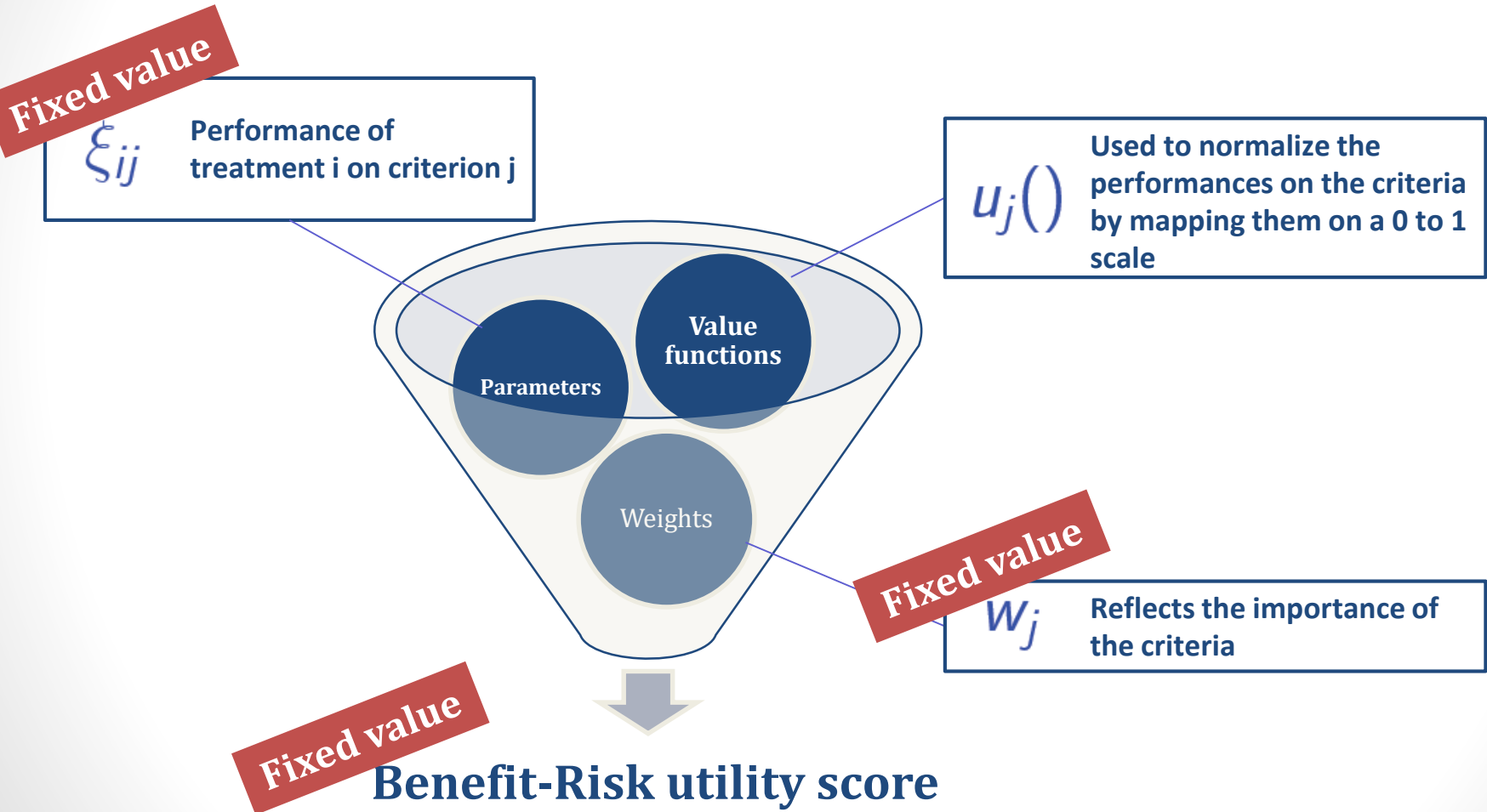
- **Deterministic MCDA (dMCDA)**
- **Probabilistic MCDA (pMCDA)**
- **Stochastic Multicriteria Acceptability Analysis (SMAA)**
- **Dirichlet SMAA**

Deterministic MCDA

Mussen et al. (2007)

Fixed value: uncertainty is ignored

Random variable: uncertainty is taken into account



$$u(\xi_i, w) = w_1 u_1(\xi_{i1}) + \dots + w_n u_n(\xi_{in})$$

Deterministic MCDA

Mussen et al. (2007)

Partial value functions

- Used to **normalize** the performances on the criteria by mapping them on a 0 to 1 scale from **best and worst preferable values of the criteria**
- Linear value functions are often used, but non linear functions can be used

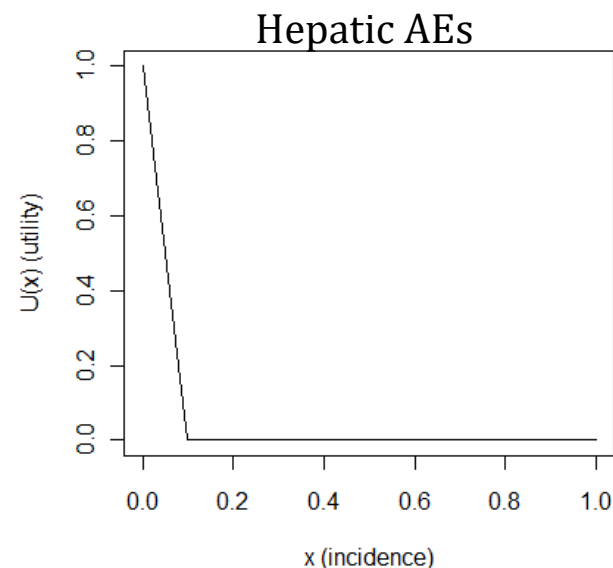
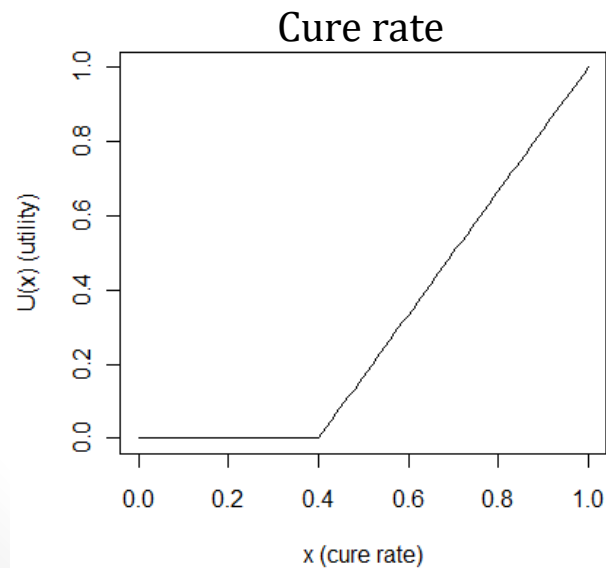
How to define the best and worst preferable values ?

- 95% confidence limits are often used, but it is not advised (data-driven, not reproducible)
- Bounds based on clinical considerations
- Bounds defined on the range of the criteria (e.g. 0-1 for probabilities of event)

Example Telithromycin

Partial value functions

Criteria		Best	Worst	Function
Favorable effects	Cure rate	100%	40%	linear
Unfavorable effects	Hepatic AEs	0%	10%	inverse linear
	Cardiac AEs	0%	10%	inverse linear
	Visual AEs	0%	10%	inverse linear
	Syncope AEs	0%	10%	inverse linear



Example Telithromycin

What do we do?

For each model, the application to the motivation example is presented

R Code

Data

Number of events

```
# Criteria = 1. Cure, 2. Hepatic, 3. Cardiac, 4. Visual, 5. Syncope  
# Treatments = 1. Telithromycin, 2. Comparator
```

```
events = t(matrix(  
  c(2185, 813,  
    57, 46,  
    4, 3,  
    14, 5,  
    2, 3), nrow=5, byrow=T))
```

Number of patients

```
N = t(matrix(  
  c(2417, 926,  
    1320, 1121,  
    1320, 1121,  
    1320, 1121,  
    1320, 1121), nrow=5, byrow=T))
```

dMCDA : example Telithromycin

Criterion parameters	
ξ_{ij} deterministic, proportion of events = # events / (# patients)	xi=events/N
Partial value functions	
$u_j(\xi_{ij}) = \frac{\xi_{ij} - \xi_{ij}''}{\xi_{ij}' - \xi_{ij}''} \text{ (linear)}$ <p> ξ_{ij}' = most preferable value ξ_{ij}'' = least preferable value </p>	<p># Criteria = 1. Cure, 2. Hepatic, 3. Cardiac, 4. Visual, 5. Syncope</p> <p># Most preferable values ; Least preferable values</p> <p>most = c(1, 0, 0, 0, 0)</p> <p>least = c(0.4, 0.1, 0.1, 0.1, 0.1)</p> <p># Partial Value Functions</p> <pre> pvf <- function(x, most, least) { return((x - least) / (most - least)) } values=pvf(xi, most, least) </pre>
Weights	
w_j deterministic, elicited by the clinicians / regulators / patients, with guidance from the statisticians (e.g surveys, swing-weighting, MACBETH tool...)	weights=c(0.30, 0.15, 0.15, 0.15, 0.25)
Utility score	
$u(\xi_{ij}, w) = \sum_{j=1}^n w_j u_j(\xi_{ij})$	<pre> us <- function (v, w) { return (sum(w*v))} us_teli = us(values[,1], weights) us_comp = us(values[,2], weights) </pre>

dMCDA : example Telithromycin

Results: dMCDA

Benefit-risk utility score:	Telithromycin	Comparator
	0.863	0.860

Conclusion from dMCDA: the benefit-risk balance of telithromycin is better than the benefit-risk balance of the comparator

But...

- Small difference
- Ignore uncertainties
- Sensitivity analyses should be conducted (varying the weights, using different criteria...)

dMCDA: conclusion

dMCDA	<ul style="list-style-type: none">✓ Simple summary✗ Deterministic, all sources of uncertainty are ignored
pMCDA	
SMAA	
Dirichlet SMAA	

Probabilistic MCDA

Waddingham et al. (2016)

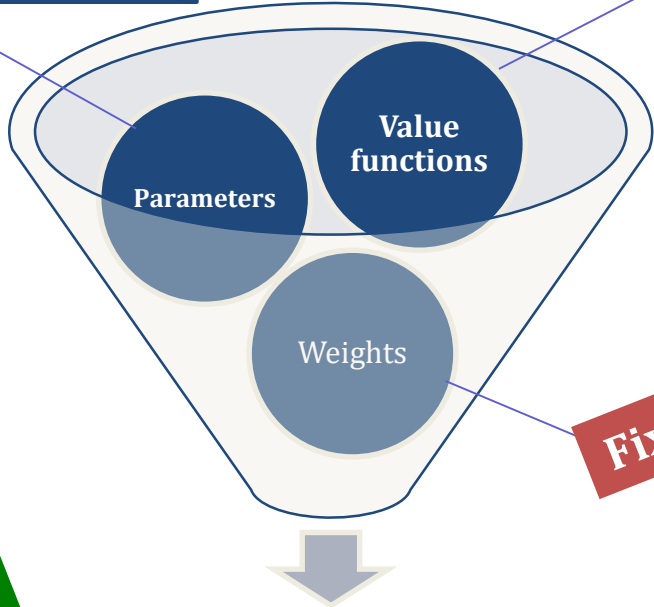
Fixed value: uncertainty is ignored

Random variable: uncertainty is taken into account

Random variable

ξ_{ij} Performance of treatment i on criterion j

$u_j()$ Used to normalize the performances on the criteria by mapping them on a 0 to 1 scale



Fixed value

w_j Reflects the importance of the criteria

Random variable

Benefit-Risk utility score

$$u(\xi_i, w) = w_1 u_1(\xi_{i1}) + \dots + w_n u_n(\xi_{in})$$

pMCDA : example Telithromycin (1/3)

Criterion parameters

$\xi_{ij} \sim \text{Beta}(a, b)$

a = # events +1

b = # non-events +1

random variables instead
of single summary values

```
Ntrt=2 ; Nendpt=5;
```

```
nsim=100000 # nb of simulations to obtain the posterior  
distributions
```

```
# Criteria = 1. Cure, 2. Hepatic, 3. Cardiac, 4. Visual, 5. Syncope
```

```
# Parameters of the posterior beta distribution
```

```
a=events+1
```

```
b=N-events+1
```

```
xi = array(0, c(nsim, Ntrt, Nendpt))
```

```
for (i in 1:Ntrt) {
```

```
  for(j in 1:Nendpt) {
```

```
    xi[,i,j]=rbeta(nsim, a[i,j], b[i,j])
```

```
  }
```

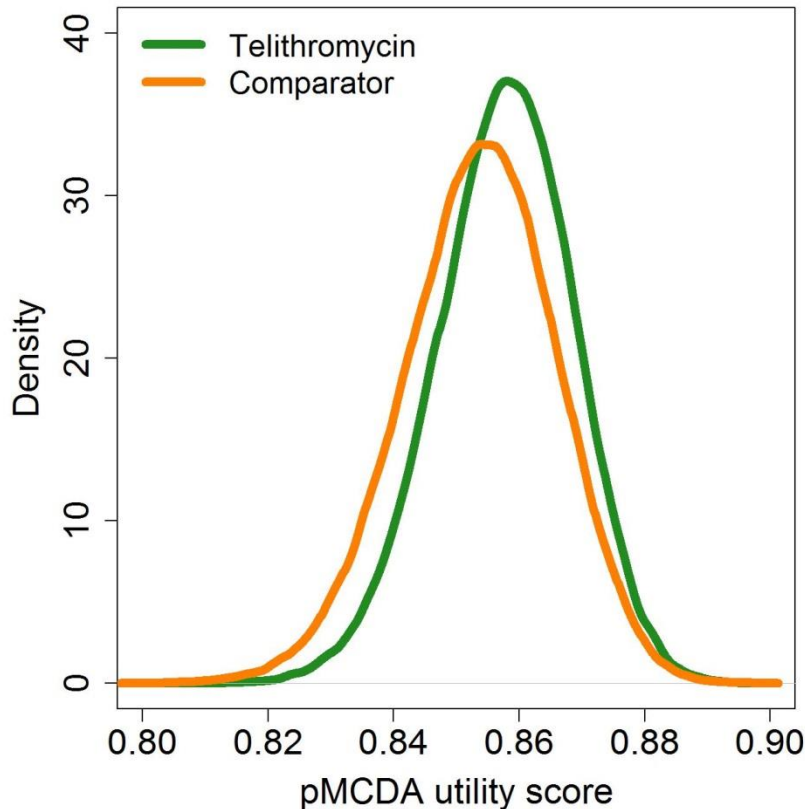
pMCDA : example Telithromycin (2/3)

Partial value functions	
$u_j(\xi_{ij}) = \frac{\xi_{ij} - \xi_{ij}''}{\xi_{ij}' - \xi_{ij}''} \text{ (linear)}$ <p> ξ_{ij}' = most preferable value ξ_{ij}'' = least preferable value </p>	<p># Most preferable values most = c(1, 0, 0, 0, 0)</p> <p># Least preferable values least = c(0.4, 0.1, 0.1, 0.1, 0.1)</p> <pre> pvf <- function(x, most, least) { return((x - least) / (most - least))} values = array(0, c(nsim, Nendpt, Ntrt)) for (i in 1:nsim) { values[i,]=pvf(t(xi[i,]), most, least) } </pre>
Weights	
w_j remain deterministic	weights=c(0.30, 0.15, 0.15, 0.15, 0.25)
Utility score	
$u(\xi_{ij}, w) = \sum_{j=1}^n w_j u_j(\xi_{ij})$ <p>random variables</p>	<pre> us <- function (v, w) { return (sum(w*v))} us_teli=us_comp=diff=vector(length=nsim) for (i in 1:nsim) { us_teli[i] = us(values[i,1], weights) us_comp[i] = us(values[i,2], weights) diff[i]=us_teli[i]-us_comp[i] } </pre>

pMCDA : example Telithromycin (3/3)

Results: pMCDA

- Distribution of the B-R utility scores
- Statistics on the B-R utility scores



Treatment	Median (95% CrI)
Telithromycin	0.858 (0.836;0.875)
Comparator	0.854 (0.829;0.873)
Difference	0.004 (-0.028;0.032)

- Probability to be better than the comparator

**Probability
Telithromycin > Comparator**

60%

pMCDA: conclusion

dMCDA	<ul style="list-style-type: none">✓ Simple summary✗ Deterministic, all sources of uncertainty are ignored
pMCDA	<ul style="list-style-type: none">✓ Takes into account uncertainty in treatment effects on the criteria✗ Preferences of decision-makers (weights) are explicitly required
SMAA	
Dirichlet SMAA	

SMAA

Tervonen et al. (2011)

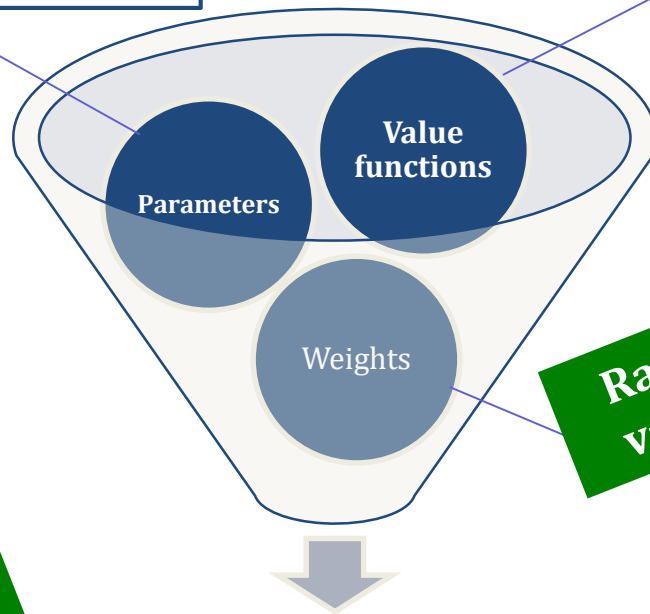
Fixed value: uncertainty is ignored

Random variable: uncertainty is taken into account

Random variable

ξ_{ij} Performance of treatment i on criterion j

$u_j()$ Used to normalize the performances on the criteria by mapping them on a 0 to 1 scale



Random variable

w_j Reflects the importance of the criteria

Random variable

Benefit-Risk utility score

$$u(\xi_i, w) = w_1 u_1(\xi_{i1}) + \dots + w_n u_n(\xi_{in})$$

SMAA

Tervonen et al. (2011)

Assumption: the weights have a uniform distribution on a space of weights, that needs to be defined

- No information, i.e. no preference between the criteria

$$W = \{w \in \mathbb{R}^n, w > 0, \sum_{j=1}^n w_j = 1\}$$

- Restrictions of the space of the weights
 - Upper and/or lower bounds
 - Complete ranking of the criteria
 - Equality of weights between benefits and risks

Example for 3 criteria

w_1, w_2, w_3

Figure: Full space

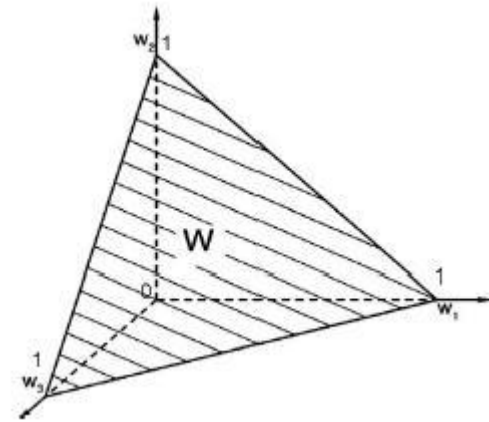
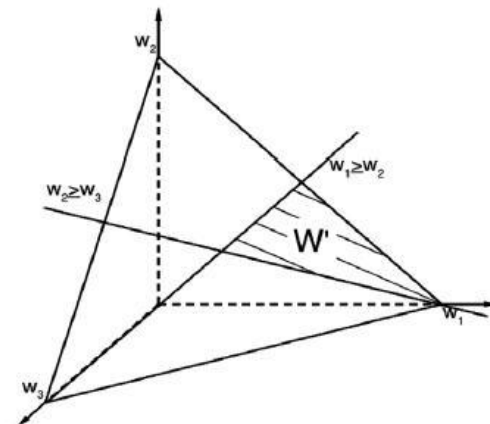


Figure: Ranking: $w_1 \geq w_2 \geq w_3$



SMAA: example Telithromycin (1/3)

Criterion parameters	
$\xi_{ij} \sim \text{Beta}(a, b)$ a = # events + 1 b = # non-events + 1 random variables instead of single summary values	<pre> Ntrt=2 ; Nendpt=5; nsim=100000 # nb of simulations to obtain the posterior distributions # Criteria = 1. Cure, 2. Hepatic, 3. Cardiac, 4. Visual, 5. Syncope # Parameters of the posterior beta distribution a=events+1 b=N-events+1 xi = array(0, c(nsim, Ntrt, Nendpt)) for (i in 1:Ntrt) { for(j in 1:Nendpt) { xi[,i,j]=rbeta(nsim, a[i,j], b[i,j]) } } </pre>
Partial value functions	
$u_j(\xi_{ij}) = \frac{\xi_{ij} - \xi_{ij}''}{\xi_{ij}' - \xi_{ij}''} \text{ (linear)}$ $\xi_{ij}' = \text{most preferable value}$ $\xi_{ij}'' = \text{least preferable value}$	<pre> # Most preferable values most = c(1, 0, 0, 0, 0) # Least preferable values least = c(0.4, 0.1, 0.1, 0.1, 0.1) pvf <- function(x, most, least) { return((x - least) / (most - least))} values = array(0, c(nsim, Nendpt, Ntrt)) for (i in 1:nsim) { values[i,,]=pvf(t(xi[i,,]), most, least)} </pre>

SMAA: example Telithromycin (2/3)

Weights

w_j **random variables** with a joint uniform distribution on a weight space W

Example: use simplex.sample from package hitandrun to generate uniform unit simplexes

```
library(hitandrun)  
weights=simplex.sample(Nendpt, nsim, sort=FALSE)$samples
```

Utility score

$$u(\xi_{ij}, w) = \sum_{j=1}^n w_j u_j(\xi_{ij})$$

random variables

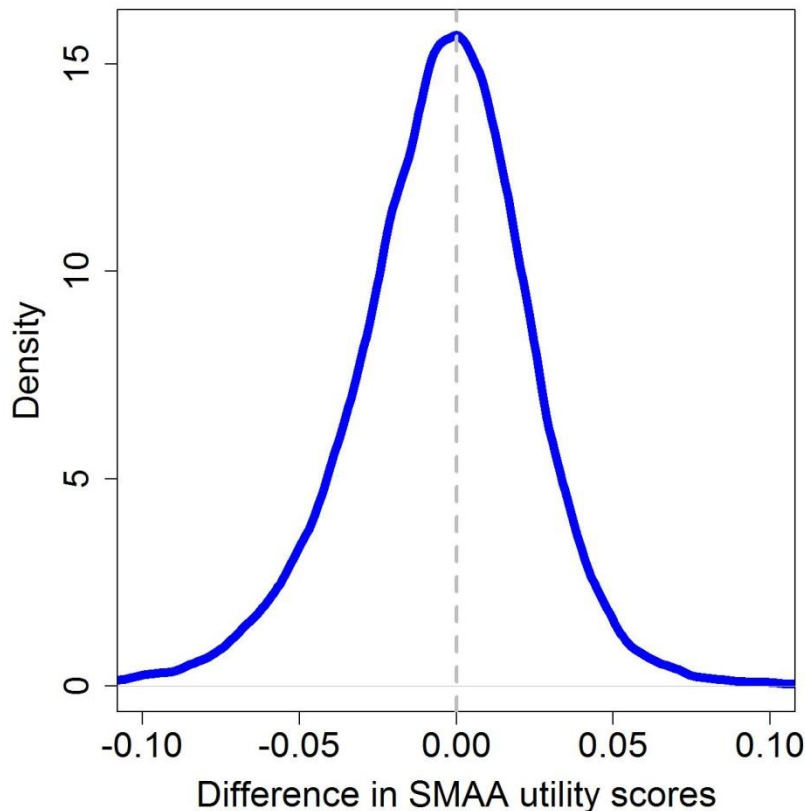
```
us <- function (v, w) { return (sum(w*v))}  
us_teli=us_comp=diff=vector(length=nsim)  
for (i in 1:nsim) {  
  us_teli[i] = us(values[i,,1], weights[i,])  
  us_comp[i] = us(values[i,,2], weights[i,])  
  diff[i]=us_teli[i]-us_comp[i]  
}
```

SMAA: example Telithromycin (3/3)

Results: SMAA

Without weight elicitation

- Distribution of the difference in B-R utility scores
- Statistics on the difference in B-R utility scores



Treatment	Median (95% CrI)
Difference	-0.005 (-0.067;0.04)

- Probability to be better than the comparator

Probability Telithromycin > Comparator
45%

SMAA: conclusion

dMCDA	<ul style="list-style-type: none">✓ Simple summary✗ Deterministic, all sources of uncertainty are ignored
pMCDA	<ul style="list-style-type: none">✓ Takes into account uncertainty in treatment effects on the criteria✗ Preferences of decision-makers (weights) are explicitly required
SMAA	<ul style="list-style-type: none">✓ Takes into account uncertainty in treatment effects on the criteria✓ Does not require the elicitation of preferences to weigh the criteria✗ Interpretation less straightforward✗ High degree of uncertainty in the results
Dirichlet SMAA	

Dirichlet SMAA

Saint-Hilary et al. (2017)

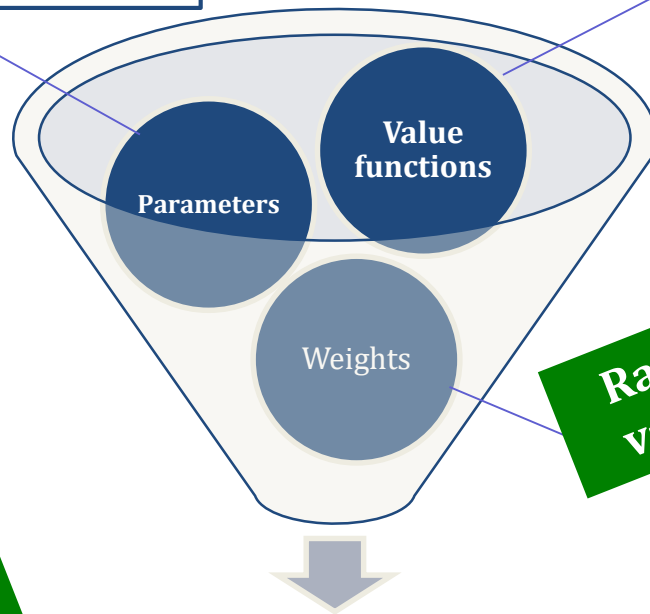
Fixed value: uncertainty is ignored

Random variable: uncertainty is taken into account

Random variable

ξ_{ij} Performance of treatment i on criterion j

$u_j()$ Used to normalize the performances on the criteria by mapping them on a 0 to 1 scale



Random variable

w_j Reflects the importance of the criteria

Random variable

Benefit-Risk utility score

$$u(\xi_i, w) = w_1 u_1(\xi_{i1}) + \dots + w_n u_n(\xi_{in})$$

Dirichlet SMAA

Saint-Hilary et al. (2017)

- w_j : weights are **random variables**, following a **Dirichlet distribution**

$$(w_1, \dots, w_n) \sim \text{Dirichlet}(\alpha_1, \dots, \alpha_n)$$

Property:

the means of all w_i stay the same if all α_i are scaled with the same multiplicative constant, with variances getting smaller as the parameters α_i grow.

- We rewrite the Dirichlet distribution as follows:

$$(w_1, \dots, w_n) \sim \text{Dirichlet}(c \cdot (w_1^0, \dots, w_n^0))$$

With: (i) $0 \leq w_1^0, \dots, w_n^0 \leq 1$ with $\sum_{j=1}^n w_j^0 = 1$

(ii) c , a scaling constant, that can vary from 0 to $+\infty$

Dirichlet SMAA

Saint-Hilary et al. (2017)

- The variances of w_j are inversely proportional to c .
- They equal to infinity when $c = 0$ and to zero when $c = +\infty$.
- **Dirichlet SMAA** corresponds to:
 - pMCDA, when $c = +\infty$, as weights are deterministic ($w_j = w_j^0, j = 1, \dots, n$)
 - SMAA, without weight elicitation when $w_1^0 = \dots = w_n^0 = 1/n$ and $c = n$
- c : **confidence level** of the decision-makers in the elicitation of their preferences, which impact on the results can be assessed using different values of c .

Dirichlet SMAA: example Telithromycin (1/3)

Criterion parameters	
$\xi_{ij} \sim \text{Beta}(a, b)$ a = # events + 1 b = # non-events + 1 random variables instead of single summary values	<pre> Ntrt=2 ; Nendpt=5; nsim=100000 # nb of simulations to obtain the posterior distributions # Criteria = 1. Cure, 2. Hepatic, 3. Cardiac, 4. Visual, 5. Syncope # Parameters of the posterior beta distribution a=events+1 b=N-events+1 xi = array(0, c(nsim, Ntrt, Nendpt)) for (i in 1:Ntrt) { for(j in 1:Nendpt) { xi[,i,j]=rbeta(nsim, a[i,j], b[i,j]) } } </pre>
Partial value functions	
$u_j(\xi_{ij}) = \frac{\xi_{ij} - \xi_{ij}''}{\xi_{ij}' - \xi_{ij}''} \text{ (linear)}$ $\xi_{ij}' = \text{most preferable value}$ $\xi_{ij}'' = \text{least preferable value}$	<pre> # Most preferable values most = c(1, 0, 0, 0, 0) # Least preferable values least = c(0.4, 0.1, 0.1, 0.1, 0.1) pvf <- function(x, most, least) { return((x - least) / (most - least))} values = array(0, c(nsim, Nendpt, Ntrt)) for (i in 1:nsim) { values[i,,]=pvf(t(xi[i,,]), most, least)} </pre>

Dirichlet SMAA: example Telithromycin (2/3)

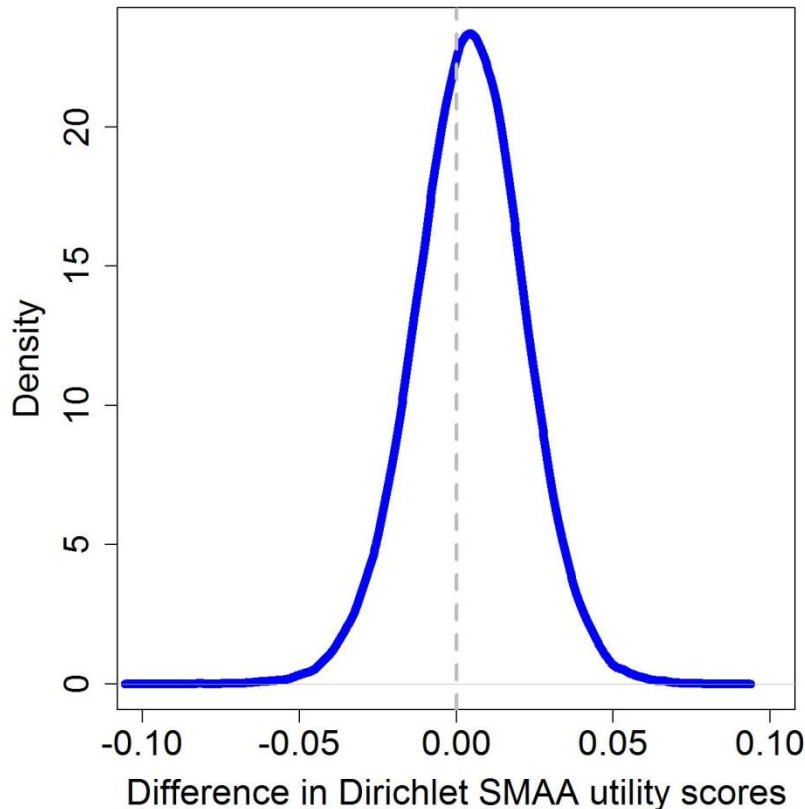
Weights	
<p>w_j random variables with a Dirichlet distribution</p> <p>c = confidence factor, level of confidence of the decision-makers in their weight elicitation</p>	<pre>library(gtools) # Example for c=50 c=50 weights=rdirichlet(nsim,c(0.30, 0.15, 0.15, 0.15, 0.25)*c)</pre>
Utility score	
$u(\xi_{ij}, w) = \sum_{j=1}^n w_j u_j(\xi_{ij})$ <p>random variables</p>	<pre>us <- function (v, w) { return (sum(w*v))} us_teli=us_comp=diff=vector(length=nsim) for (i in 1:nsim) { us_teli[i] = us(values[i,,1], weights[i,]) us_comp[i] = us(values[i,,2], weights[i,]) diff[i]=us_teli[i]-us_comp[i] }</pre>

Dirichlet SMAA: example Telithromycin (3/3)

Results: Dirichlet SMAA

For a given confidence factor (here, $c=50$)

- Distribution of the difference in B-R utility scores
- Statistics on the difference in B-R utility scores



Treatment	Median (95% CrI)
Difference	0.004 (-0.031;0.033)

- Probability to be better than the comparator

Probability Telithromycin > Comparator
60%

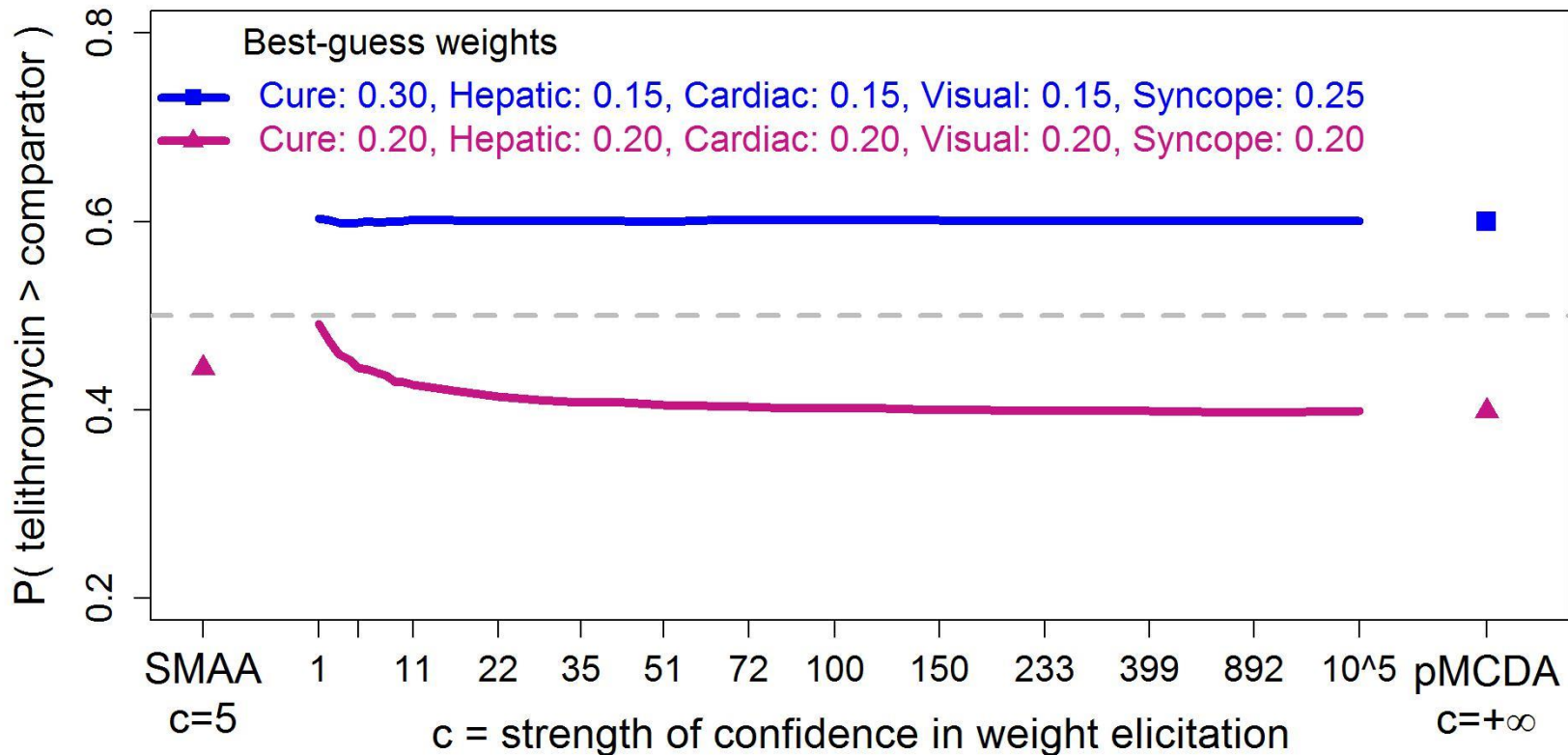
Dirichlet SMAA: example Telithromycin (3/3)

Results: Dirichlet SMAA

Varying confidence factor

- Probability to be better than the comparator

*Taking into account the **uncertainty** of the decision-makers in their weight elicitation*



Dirichlet SMAA: conclusion

dMCDA	<ul style="list-style-type: none">✓ Simple summary✗ Deterministic, all sources of uncertainty are ignored
pMCDA	<ul style="list-style-type: none">✓ Takes into account uncertainty in treatment effects on the criteria✗ Preferences of decision-makers (weights) are explicitly required
SMAA	<ul style="list-style-type: none">✓ Takes into account uncertainty in treatment effects on the criteria✓ Does not require the elicitation of preferences to weigh the criteria✗ Interpretation less straightforward✗ High degree of uncertainty in the results
Dirichlet SMAA	<ul style="list-style-type: none">✓ Takes into account uncertainty in treatment effects on the criteria✓ Takes into account uncertainty in weight elicitation, and allows flexibility by making the variance of the weights vary✓ Permits to account for a new source of uncertainty: the level of confidence of the decision-makers in their weight elicitation✓ All parameters have a natural interpretation: treatment effects, decision-makers' preferences and their strength of confidence

Other examples in backup slides

- dMCDA: Gardasil[®] vaccine for preventing anal cancer in males
- pMCDA: Natalizumab for the treatment of Relapsing Remitting Multiple Sclerosis (RRMS)
- Dirichlet SMAA: fictive case-study in depression (inspired by a real case)

MCDA and its extensions

Conclusion

- Powerful quantitative **decision-making** tools
 - Recognized by the EMA
- **Subjectivity**: input from clinical/regulatory/patients needed to determine the criteria and their relative importance, as well as the range of preferences
 - Sensitivity analyses should be performed
 - Need to consider the various sources of uncertainty
- Relative **complexity**
 - Collecting and summarizing the data on multiple criteria, possibly from different sources
- Usually used late in the development → could be applied in **Early development** using biomarkers

Main references (1/2)

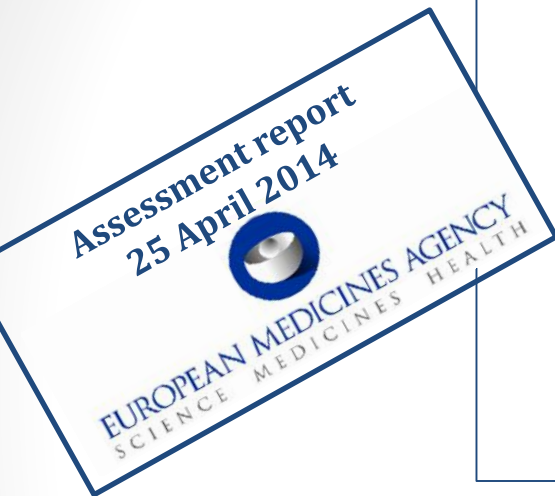
- EMA (2010). Benefit–risk methodology project. Work package reports: applicability of current tools and processes for regulatory benefit–risk assessment. Available at <http://www.ema.europa.eu/>
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- EFSPi-SIG Benefit-risk <http://www.benefit-risk-assessment.com/welcome-to-the-benefit-risk-blog-of-the-efspi-benefit-risk-sig/>
- Mt-Isa, S., Ouwens, M., Robert, V., Gebel, M., Schacht, A., and Hirsch, I. (2015). Structured benefit–risk assessment: a review of key publications and initiatives on frameworks and methodologies. *Pharmaceutical Statistics* **15**, 324–332. doi: 10.1002/pst.1690
- Mt-Isa, S. *et al.* (2014). Balancing benefit and risk of medicines: a systematic review and classification of available methodologies. *Pharmacoepidemiology and Drug Safety* **23**, 667–678. doi: 10.1002/pds.3636
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Main references (2/2)

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Back-up slides

Example 1: Gardasil® vaccine for preventing anal cancer in males



(...) the MAH has used the ‘problem, objectives, alternatives, consequences, trade-offs, uncertainty, risk attitude, linked decisions’ (PrOACT-URL) and the multi criteria decision analysis’ (MCDA) approaches, which are two similar and well-structured approaches to estimate the overall benefit-risk balance, both on a **qualitative (PrOACT and MCDA) and a quantitative (MCDA)** point of view. **These two approaches allow taking into consideration all the potential benefits and all the potential risks within a single evaluation.**

Quantitative benefit-risk assessment by MCDA of the quadrivalent HPV vaccine for preventing anal cancer in males

Lydie Marcelon, Thomas Verstraeten, Geraldine Dominiak-Felden & François Simondon

To cite this article: Lydie Marcelon, Thomas Verstraeten, Geraldine Dominiak-Felden & François Simondon (2016) Quantitative benefit-risk assessment by MCDA of the quadrivalent HPV vaccine for preventing anal cancer in males, *Expert Review of Vaccines*, 15:1, 139-148, DOI: [10.1586/14760584.2016.1107480](https://doi.org/10.1586/14760584.2016.1107480)



Example 1: Gardasil[®] vaccine for preventing anal cancer in males

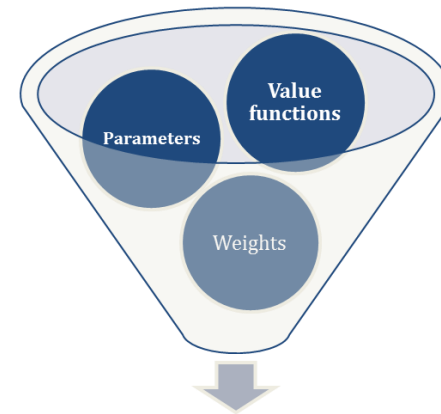
PrOACT-URL

Generic **qualitative** framework to structure decision problems



dMCDA

Quantitative approach to synthesize the results



Benefit-Risk utility score

$$u(\xi_i, w) = w_1 u_1(\xi_{i1}) + \dots + w_n u_n(\xi_{in})$$

Example 1: Gardasil[®] vaccine for preventing anal cancer in males

Data sources

Identification of key benefits and risks	Weights	Treatment performances on the criteria
1) Sanofi Pasteur MSD clinicians and epidemiologists, with working experience on the qHPV vaccine 2) Panel of six external experts		<ul style="list-style-type: none"> Merck/Sanofi Pasteur MSD-sponsored clinical trials Post-authorization study reports

Treatment groups

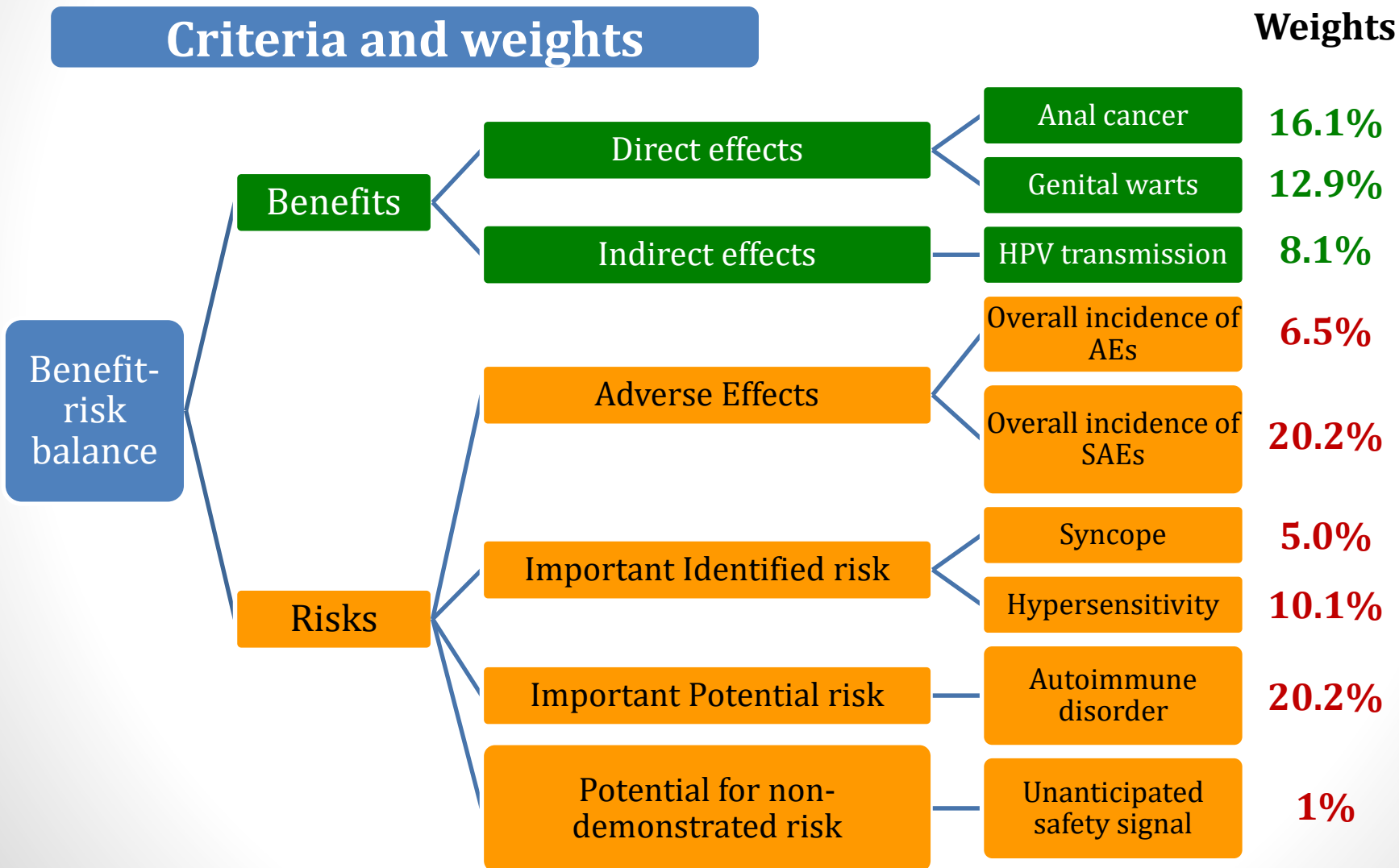
- Gardasil[®]
- No vaccination

Example 1: Gardasil[®] vaccine for preventing anal cancer in males

For submission

Criteria and weights

Weights



Example 1: Gardasil[®] vaccine for preventing anal cancer in males

Results: dMCDA

Benefit-risk utility score:	Gardasil [®]	No vaccination
	66	46

Sensitivity analyses: results are robust to changes in

- the **weight** assigned to the individual criteria or nodes
- the **model parameters** (e.g. inclusion of data less favorable to the vaccine or excluding all beneficial effects other than anal cancer prevention)

Assessment report
25 April 2014

EUROPEAN MEDICINES AG
SCIENCE MEDICINES H

"MCDA is a method considered to be **useful as a complementary and supportive tool**. Through a number of steps the purpose is to **bring together evaluations of options on both benefits and risks into one overall evaluation** taking into account what is considered best current evidence."

Example 1: Gardasil[®] vaccine for preventing anal cancer in males

Conclusion

Assessment report
25 April 2014

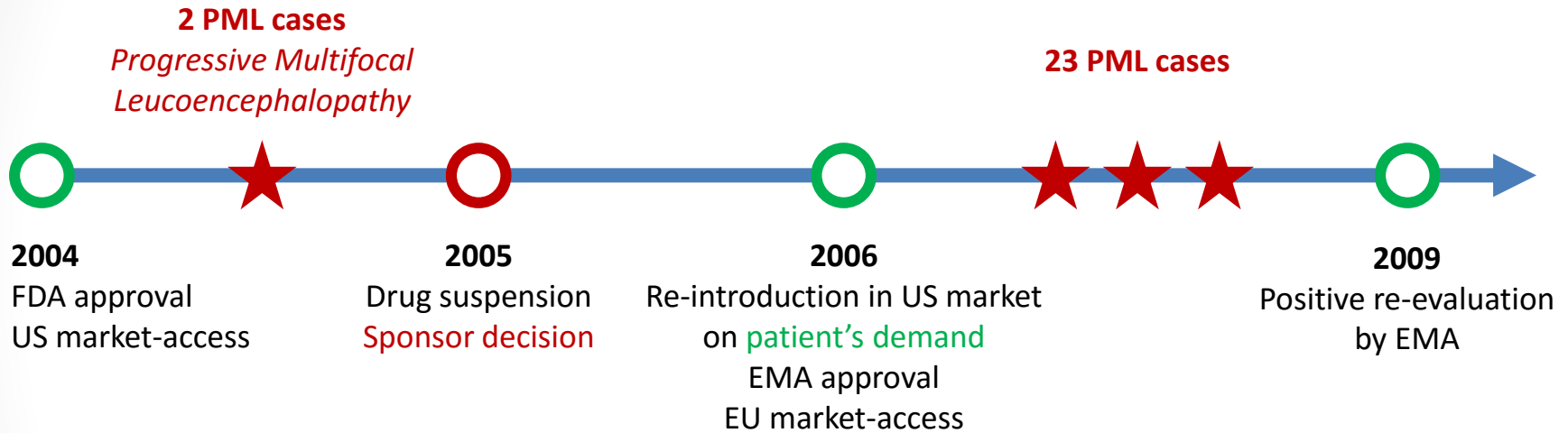


EUROPEAN MEDICINES AGENCY
SCIENCE MEDICINES HEALTH

“The benefit-risk balance [of Gardasil[®]] is considered positive.”



Example 2: Natalizumab for the treatment of Relapsing Remitting Multiple Sclerosis (RRMS)



→ Rare serious side effect in an effective treatment for a serious disease

Was the decision right to keep natalizumab on the market given that increased episodes of PML were observed?

Example 2: Natalizumab for the treatment of Relapsing Remitting Multiple Sclerosis (RRMS)

Data sources

Identification of key benefits and risks	Weights	Treatment performances on the criteria
Individual experts (based largely on data from the SPC and the EPAR for natalizumab)	Patient representatives (Decision conference held on 23 Sept 2011)	Mainly: <ul style="list-style-type: none"> • EPARs • Literature search

EPAR: European public assessment reports

SPC: Summary of Product Characteristics

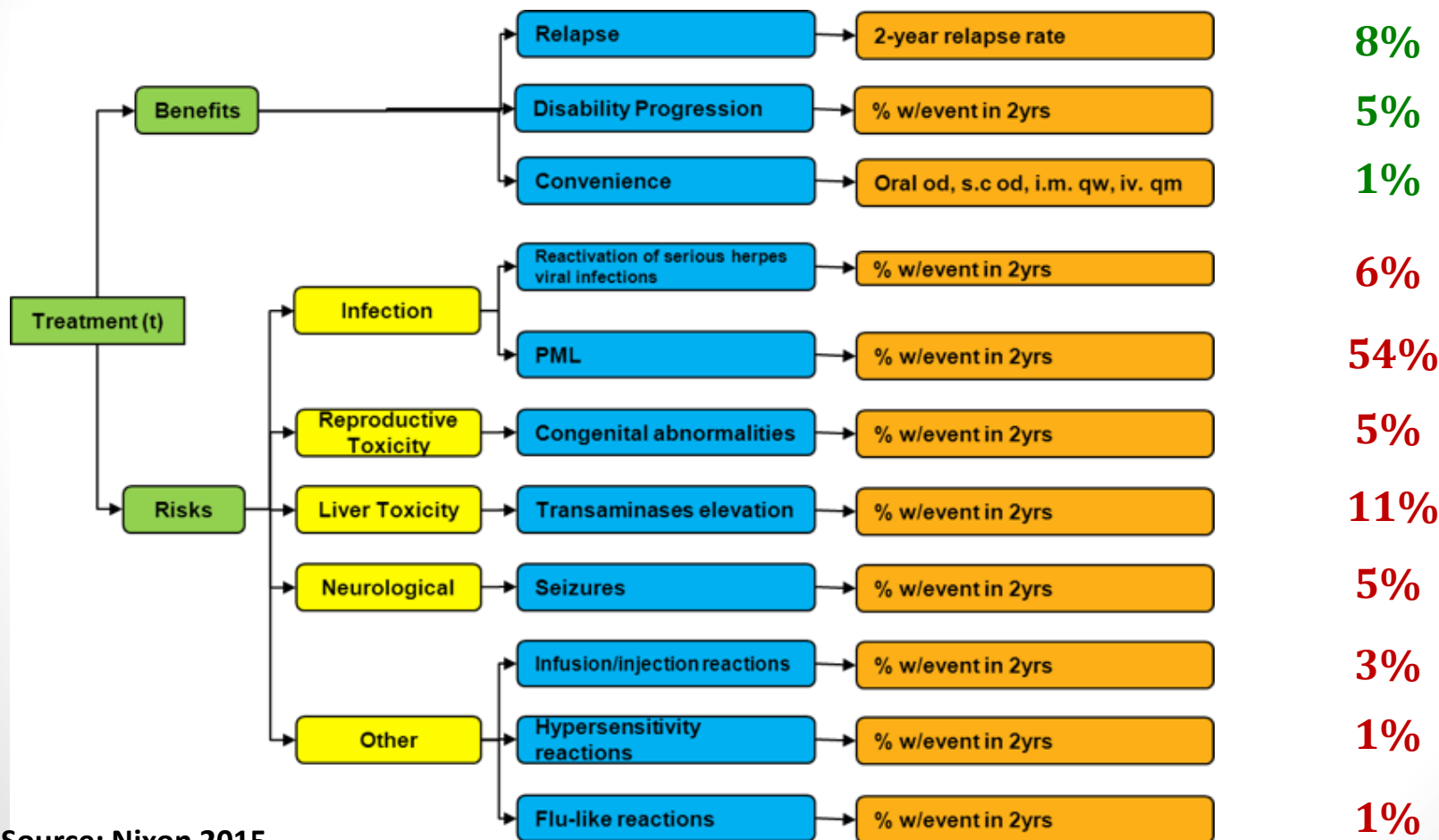
Treatment groups

- Natalizumab
- Placebo
- Glatiramer Acetate
- Beta-interferon

Example 2: Natalizumab for the treatment of Relapsing Remitting Multiple Sclerosis (RRMS)

Criteria and weights

Weights

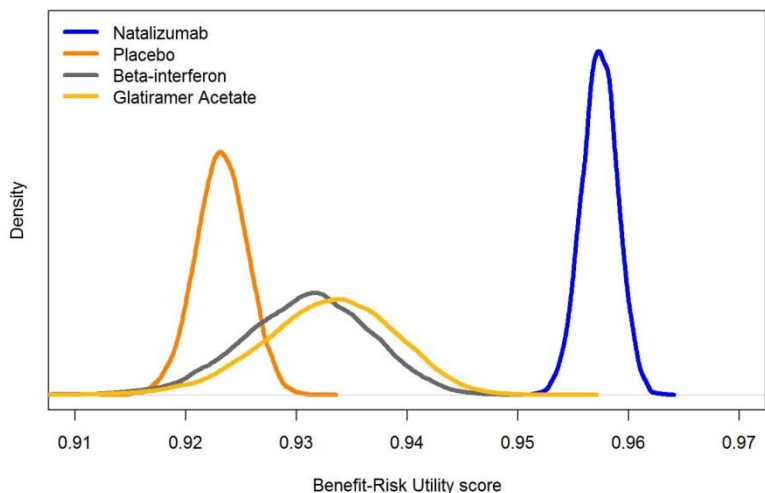


Source: Nixon 2015

Example 2: Natalizumab for the treatment of Relapsing Remitting Multiple Sclerosis (RRMS)

Results: pMCDA

- Distribution of the B-R utility scores
- Statistics on the B-R utility scores



Treatment	Median (95% CrI)
Placebo	0.92 (0.92;0.93)
Natalizumab	0.96 (0.95;0.96)
Beta-interferon	0.93 (0.92;0.94)
Glatiramer Acetate	0.93 (0.92;0.94)

- Probability to be better than the control

Treatment	Probability Treatment > Placebo
Natalizumab	100%
Beta-interferon	80%
Glatiramer Acetate	79%

Source: Waddingham 2016

Example 2: Natalizumab for the treatment of Relapsing Remitting Multiple Sclerosis (RRMS)

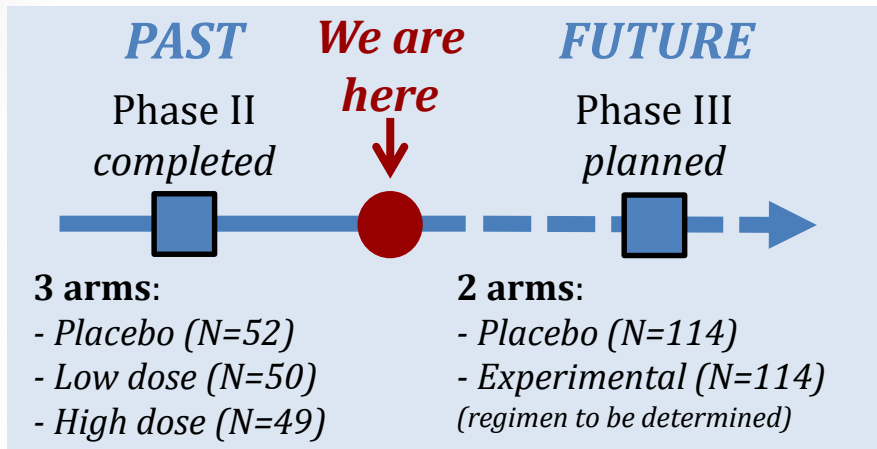
Conclusion

- Despite the incidence of the serious rare adverse events PML, Natalizumab has the best benefit-risk profile

Example 3: Fictive case-study in depression

(inspired by a real case)

Context: Go/No-Go Ph III



Results from Ph II

- Effective treatment
- **Dose-response relationship** for efficacy and safety
- Hypokalemia may be a **serious adverse effect**

Considered strategies for Ph III

- Low dose
- High dose
- Low dose with possible dose-increase
- High dose with potassium supplementation

Which dose/regimen has the best chance to have a positive Benefit-Risk balance versus Placebo in Phase III?

Example 3: Fictive case-study in depression

(inspired by a real case)

Data sources

Identification of key benefits and risks	Weights	Treatment performances on the criteria
Sponsor multidisciplinary team (clinical, pharmacovigilance, regulatory, statistics etc.)		Phase II trial

Treatment groups

- Low dose
- High dose
- Low dose with possible dose-increase
- High dose with potassium supplementation

Note: details about the combination of new hypotheses with the data (dose-increase, potassium supplementation) and about predictions of the next study are not included in this presentation

Example 3: Fictive case-study in depression

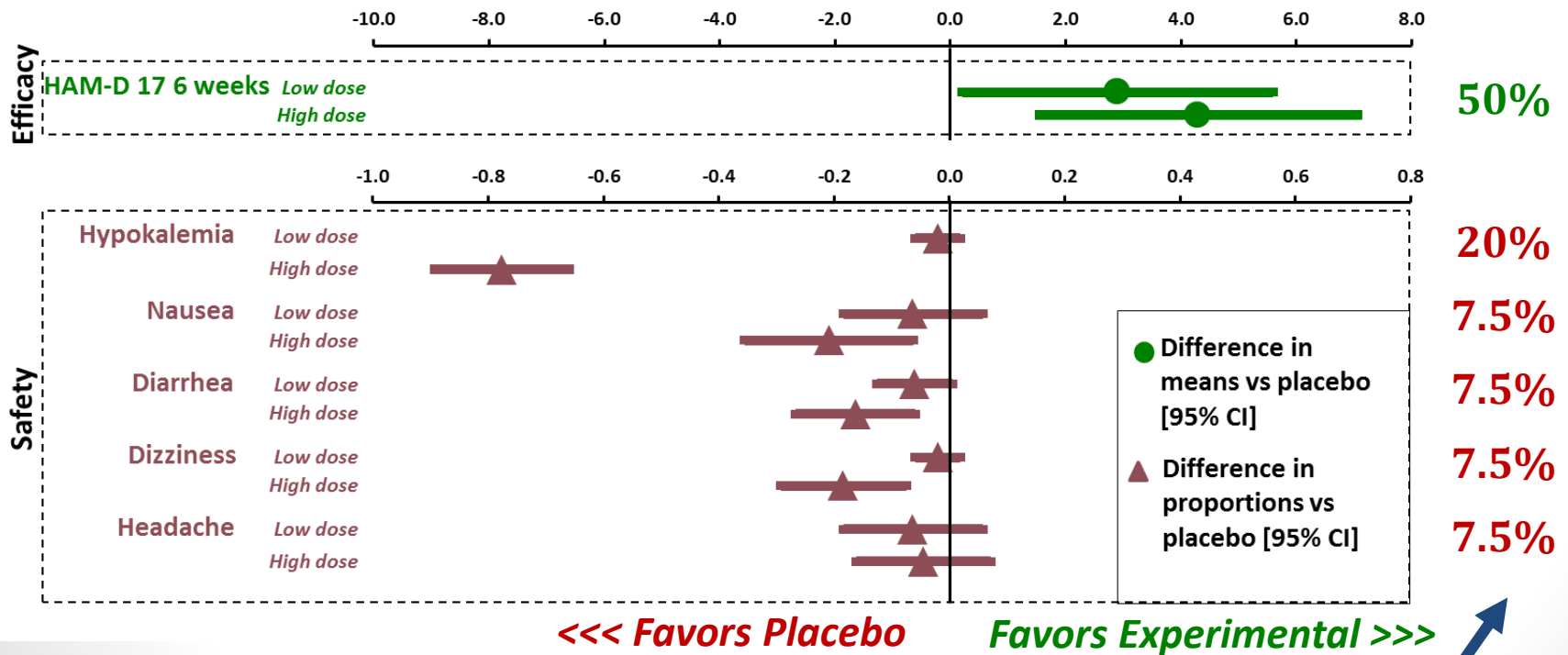
(inspired by a real case)

Criteria and weights

Results from Phase II

Primary efficacy criterion and 5 more frequent adverse events

Weights



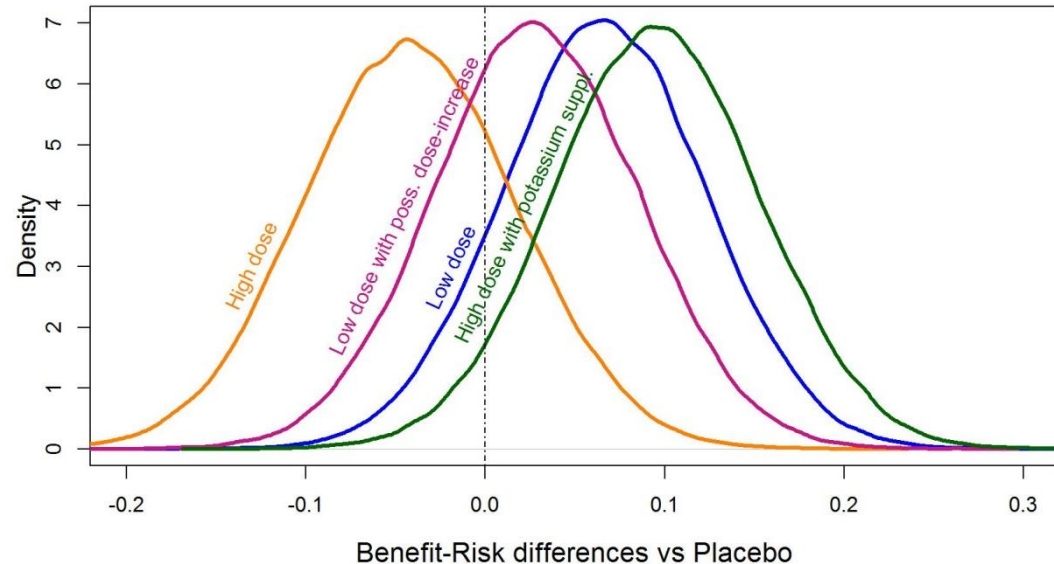
But decision-makers are not quite confident in their weight elicitation...

Example 3: Fictive case-study in depression

(inspired by a real case)

Results: Dirichlet SMAA

- Predictive distribution of the differences in B-R utility scores vs Placebo



- Statistics on differences in B-R utility scores vs Placebo

Treatment	Diff vs Placebo Median (95% CrI)
Low dose	0.07 (-0.04;0.18)
High dose	-0.04 (-0.16;0.08)
Dose-increase	0.03 (-0.08;0.14)
High dose suppl	0.10 (-0.02;0.21)

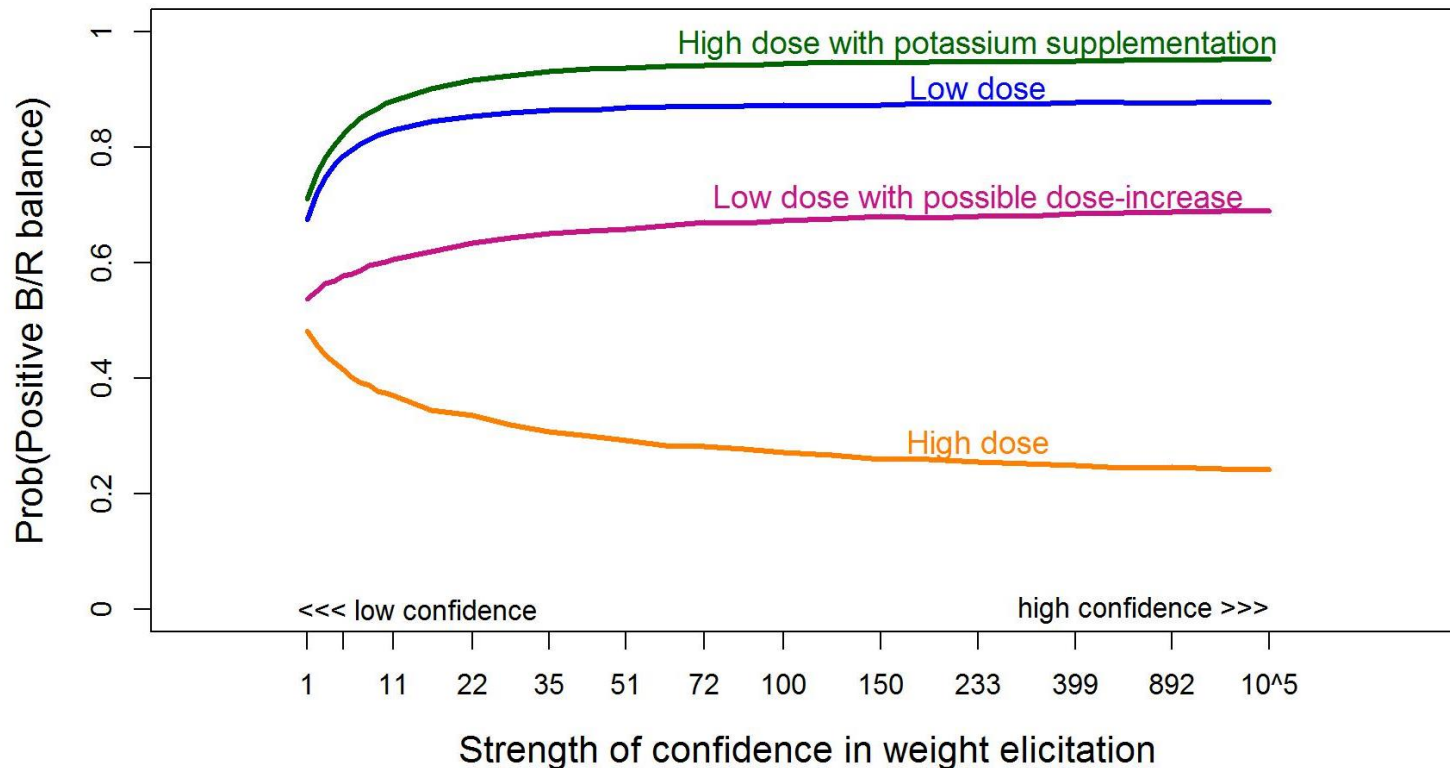
Example 3: Fictive case-study in depression

(inspired by a real case)

Results: Dirichlet SMAA

- Probability to be better than placebo in the next Ph III

*Taking into account the **uncertainty** of the decision-makers in their elicitation of preferences (weights)*



Example 3: Fictive case-study in depression

(inspired by a real case)

Conclusion

- High dose with potassium supplementation seems to be the regimen with the best benefit-risk balance vs placebo