

# **Bayesian Markov models for the cost-effectiveness analysis of HPV vaccination**

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*June 12th, 2014*

- ① Aim of the research
- ② Literature review
- ③ General introduction
  - Bayesian Markov models
- ④ Model assumptions
  - Reference population and follow-up
  - Cervical screening and HPV vaccination
  - Model structure
  - The process of sexual mixing
    - Herd immunity
  - Distributional assumptions and sources of prior information
  - Transition probabilities
- ⑤ Preliminary results
  - Convergence and autocorrelation
  - Cost-effectiveness analysis

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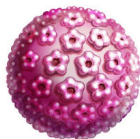
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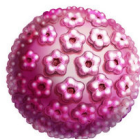
## Identifying the most cost-effective vaccination strategy against human papillomavirus (HPV)

- 1 Incorporating the effects of herd immunity into the Bayesian Markov model
- 2 Including boys in a quadrivalent HPV vaccination scheme
- 3 Considering a great variety of HPV-induced diseases



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- Mainly sexually transmitted virus
- Infects both mucous membrane and skin
- In rare cases transmission through:
  - Shared towels
  - Public saunas
  - Digital-genital contact
- Around 40 identified genotypes, including 13 high-risk types
  - HPV 16 and 18: 79.1% of all cervical cancers
  - HPV 6 and 11: anogenital warts and recurrent respiratory papillomatosis (RRP)
  - HPV 1 and 2: benign skin warts
- Contributory cause of anal, vaginal, vulvar, penile and head/neck cancers

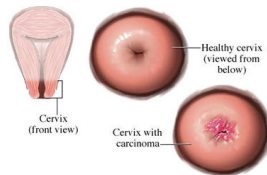
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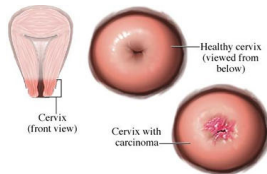


## Cervical cancer

- Yearly 2,890 new cervical cancer diagnoses
- Cervical cancer 11th most frequent cancer in females

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## Genital warts prevalence

- 4.7% in females
- 2.2% in males



## Yearly costs borne by the NHS

- £17 million for genital warts treatment
- £157 million for cervical cancer treatment



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## European public health insurance systems

- National Health Service (NHS) in the UK
- Servizio Sanitario Nazionale (SSN) in Italy
- Couverture Maladie Universelle (CMU) in France



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## Country-specific information

- Data on costs and utilities specific for Italy
- Cost-effectiveness analysis in an Italian context

- 5 databases searched with variety of search word combinations
- Altogether 116 publications reviewed and summarized
- Hybrid models for HPV vaccination
  - ① Simulate the process of sexual mixing
  - ② Calculate age-and gender-specific HPV prevalence by means of
    - Difference equations
    - ODEs
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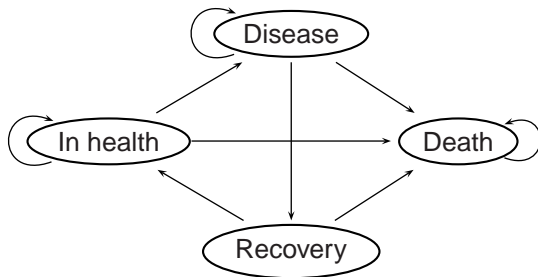
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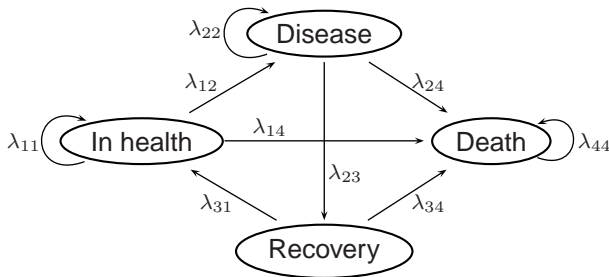
To the best of our knowledge, our methodology of including dynamic interactions between individuals directly into a static Bayesian Markov model is unique in the field of HPV transmission and disease progression modelling.

## 1. Define a structure



- Exhaustive and mutually exclusive health states

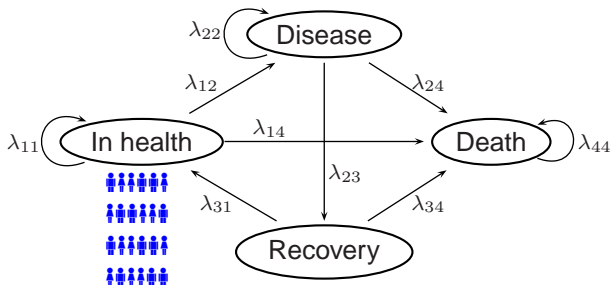
## 2. Estimate the transition probabilities



- Transition probabilities  $\lambda$  are functions of  $\theta$
- Assigning flat and informative distributions to parameters  $\theta$

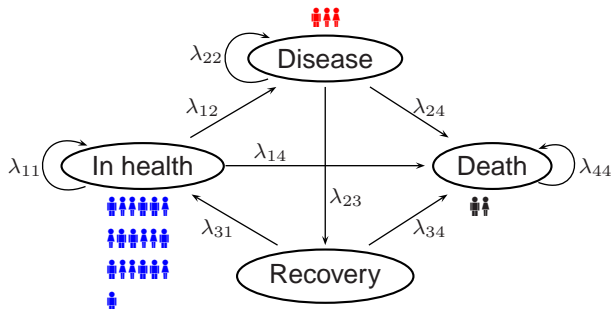


## 3. Run the simulation: $t = 0$



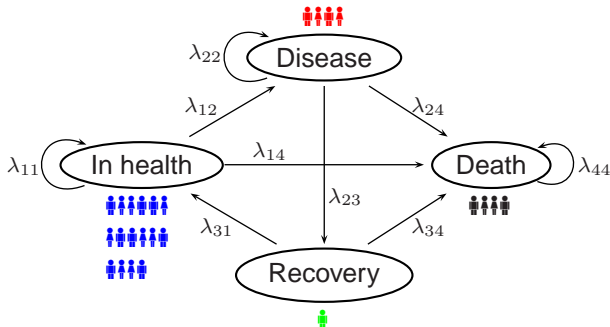
- Each health state is assigned a value of utility
- Ranging between 0 (death) and 1 (perfect health)

## 3. Run the simulation: $t = 1$

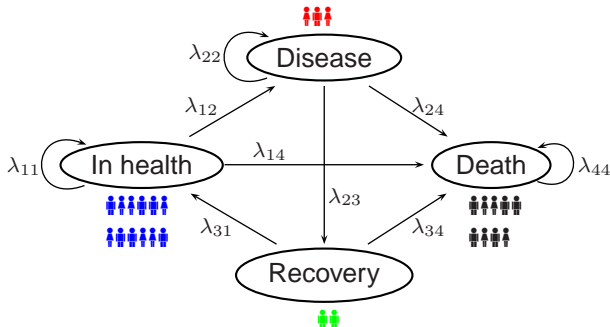


- Markov cycle length of 1 year

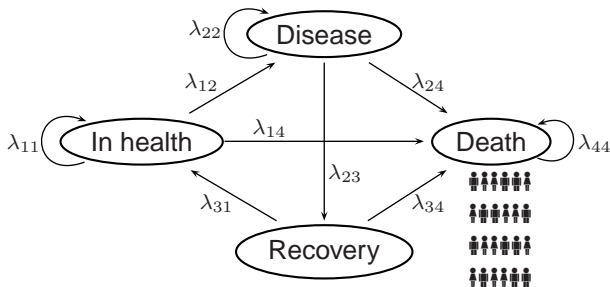
## 3. Run the simulation: $t = 2$



3. Run the simulation:  $t = 3$



3. Run the simulation:  $t = T$



- Health economic analysis of multi cohort vaccination strategy

## Reference population and follow-up

- 24 cohorts of females and males aged 12-35 years
- Follow-up period of 55 years
- Population dynamics: Entering of healthy 12 year old individuals during first 10 years of observation

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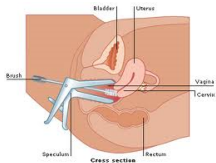
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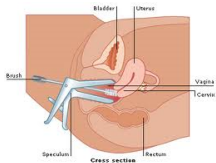
## Interventions $i$

- ① *Screening-only*: Screening in females, no intervention in males
- ② *Female-only vaccination*: Screening and vaccination in 12 year old females, no intervention in males
- ③ *Universal vaccination*: Screening and vaccination in 12 year old females, vaccination in 12 year old males
  - Sensitivity analyses to male vaccination age
- ④ *Catch-up vaccination*: Screening and vaccination in 12 year old females with a catch-up at 15 years and no intervention for males
  - Sensitivity analysis to catch-up coverage rate



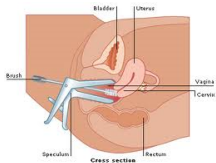
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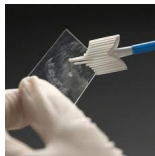
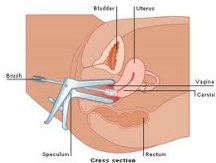
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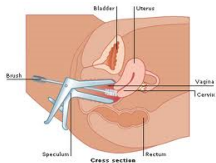
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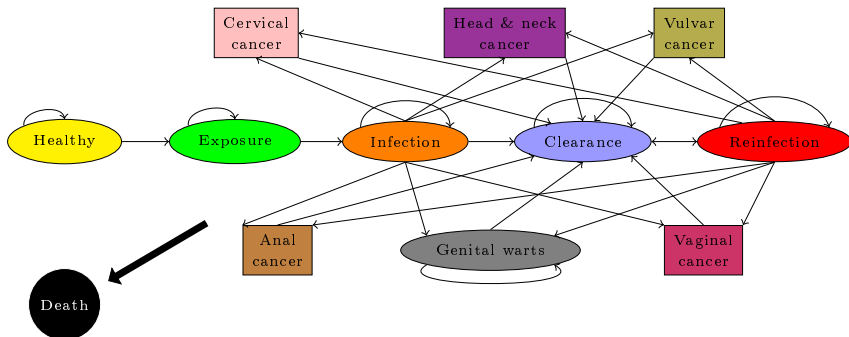


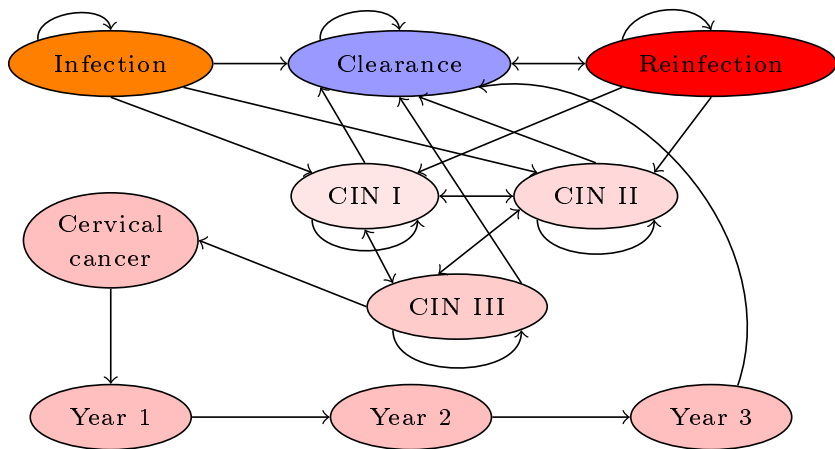
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$S_f = 36$  health states







- Most important aspect of our research
- Transforms the Bayesian MM into a hybrid model
- Accounting for herd immunity
  - unvaccinated individuals are indirectly protected
  - females and males benefit from male HPV vaccination by
    - 1 decrease in prevalence of HPV and induced diseases
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  - vaccine benefits are no longer underestimated in CE-analyses





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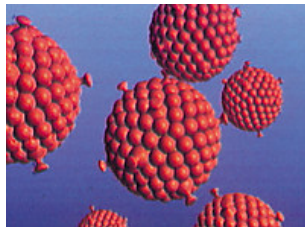
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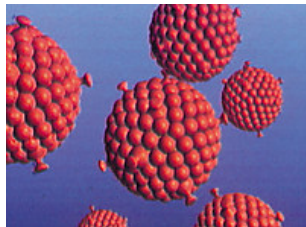
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- 2-10 lifetime sex partners



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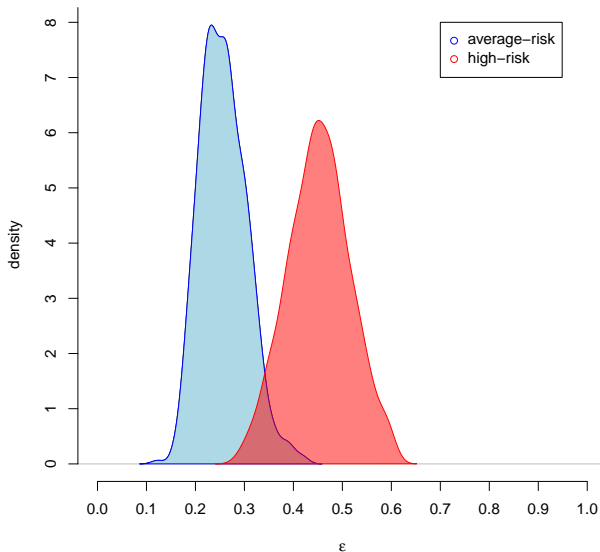
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## High-risk sexual activity

- 20% of the population
- 11 or more lifetime sex partners
- Promiscuity correlates with
  - smoking
  - a low education level
  - early first sexual intercourse before the age of 18

Distributions of HPV transmission probabilities  $\epsilon$



Sexual partnership matrix for female (average-risk group)

Age	12	15	20	25	30	35	40	45	50	55	60	65
	-14	-19	-24	-29	-34	-39	-44	-49	-54	-59	-64	-80
12-19	1%	26%	58%	15%	1%	0%	0%	0%	0%	0%	0%	0%
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Annual max, average and mean partner acquisition rate for females

	Females			Males		
Age	Min	Mean	Max	Min	Mean	Max
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  - $m_{g,s',a,a'} = 36\% \times 1.38 = 0.4968$ , for  $a' = 20-24$ ;
  - $m_{g,s',a,a'} = 49\% \times 1.38 = 0.6762$ , for  $a' = 25-29$ ;
  - $m_{g,s',a,a'} = 12\% \times 1.38 = 0.1656$ , for  $a' = 30-34$ ;

Sexual partnership matrix for female (average-risk group)

Age	12	15	20	25	30	35	40	45	50	55	60	65
	-14	-19	-24	-29	-34	-39	-44	-49	-54	-59	-64	-80
12-19	1%	26%	58%	15%	1%	0%	0%	0%	0%	0%	0%	0%
20-24	0%	0%	36%	49%	12%	2%	0%	0%	0%	0%	0%	0%
...	...	...	...	...	...	...	...	...	...	...	...	...
65-80	0%	0%	0%	0%	0%	0%	0%	0%	0%	0%	14%	86%

Annual max, average and mean partner acquisition rate for females

	Females			Males		
Age	Min	Mean	Max	Min	Mean	Max
12-19	0.74	1.26	1.78	0.90	1.92	2.94
20-24	0.54	0.96	1.38	0.68	1.38	2.09
...	...	...	...	...	...	...
60-	0.05	0.10	0.15	0.04	0.11	0.18

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Sexual partnership matrix for female (average-risk group)

Age	12 -14	15 -19	20 -24	25 -29	30 -34	35 -39	40 -44	45 -49	50 -54	55 -59	60 -64	65 -80
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## Equation introduced by Korostil et al.

$$\kappa_{g,s,a} = \epsilon \sum_{s',a'} m_{g,s,s',a,a'} \left( \frac{I_{g',s',a'}}{N_{g',s',a'}} \right)$$

- $\epsilon$  represents the HPV transmission probability per partnership
- $m_{g,s,s',a,a'}$  represents the sexual mixing matrix
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At each time point  $t$ , the probability of HPV infection depends on the pool of opposite sex partners

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- Cooper et al.'s formula is based on the assumption of constant transition probabilities over the whole observation period
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- These are directly integrated into the health state allocation algorithm of the MM
- → Health economic analysis by means of output of MM



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- 5 Calculating overall costs and utilities, resulting in the cost-effectiveness analysis



- **Just Another Gibbs Sampler (JAGS)**
- Integrated into R by means of package R2jags
- 2 parallel chains ( $n_{chains} = 2$ )
- $n_{iter} = 40,000$  simulations
- burn-in of  $n_{burn} = 4,000$
- thinning step of  $n_{thin} = 360$

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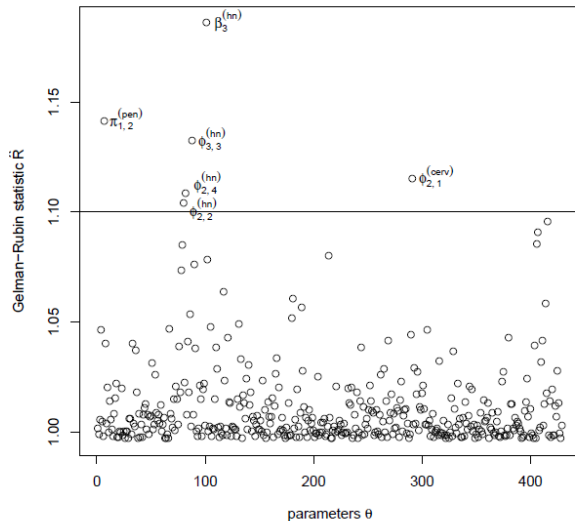
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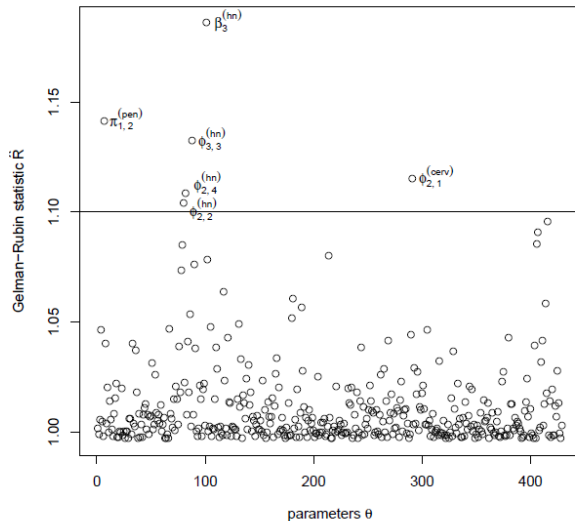
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Convergence evaluation with  $\hat{R}$  for all model parameters  $\theta$





Convergence evaluation with  $\hat{R}$  for all model parameters  $\theta$



$$\hat{R} = \sqrt{\frac{\widehat{Var}(\theta_k|y)}{W(\theta_k)}}$$

## The PVC

- Sum of overall costs in intervention  $i$  for time  $t = 1$  to  $t = 55$
- Commonly discounted by  $\nu_c = 0.03$

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## Comparison of universal to female-only vaccination

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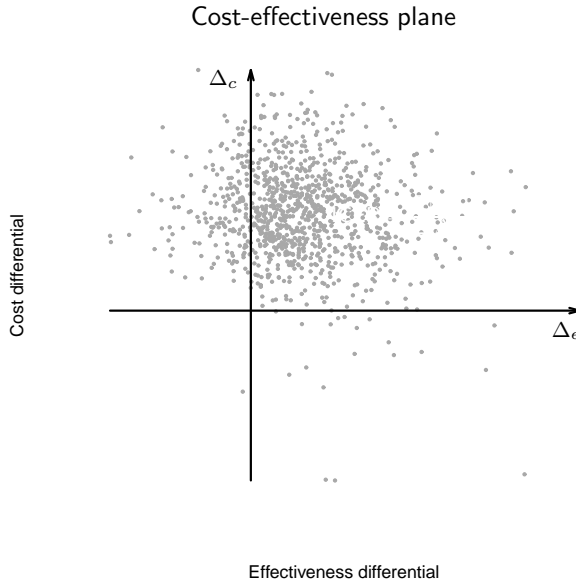
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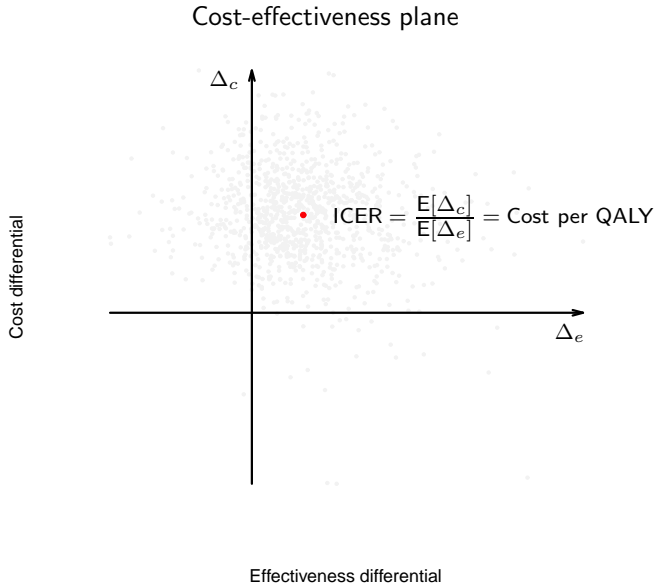
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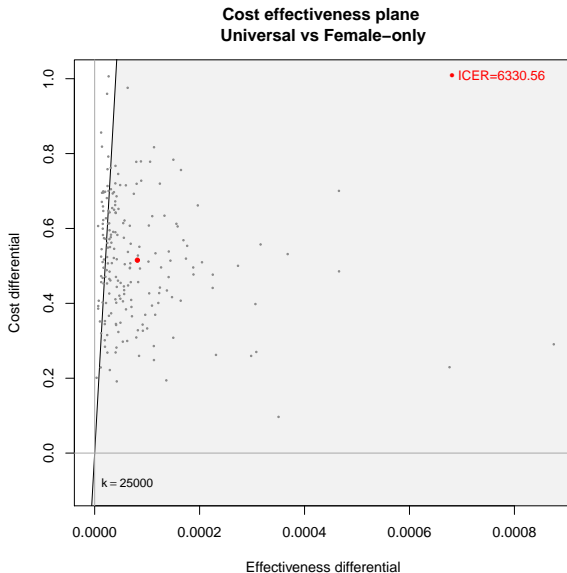
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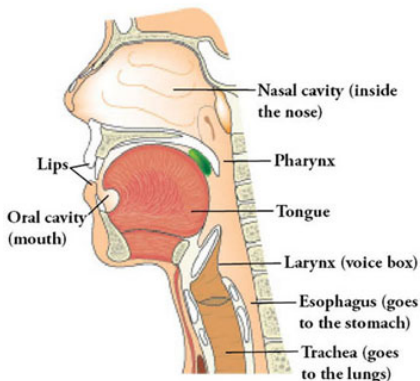
# Thank you very much for your attention.



# Appendix

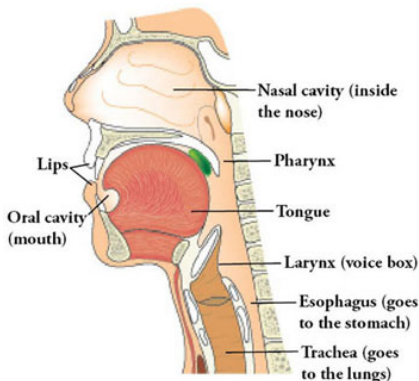
## Anal, head/neck, vaginal, vulvar, and penile cancer

- Multifactorial diseases
- HPV-induced:
  - more than 90% of anal cancers
  - more than 50% of vaginal, vulvar and penile cancers
  - 60–70% of oropharyngeal cancers
- Other head/neck cancers mainly attributed to tobacco and alcohol



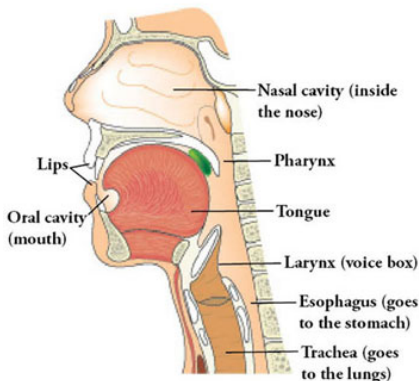
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## Databases



- Scopus
- Pubmed
- Cochrane Library
- Web of Science
- Centre for Review and Dissemination (CRD)
  - Database of Abstracts of Reviews of Effects (DARE)
  - NHS Economic Evaluation Database (EED)
  - Health Technology Assessment (HTA)



## Search word combinations

((cost-effectiveness) OR (cost-utility) OR (cost-benefit))  
AND ((HPV vaccine) OR (human papillomavirus vaccine) OR  
HPV or (human papillomavirus))

## For universal vaccination extended by

AND (boys OR male)



## Methodology

- Static vs. dynamic
- Deterministic vs. stochastic
- Ordinary differential equation (ODE) vs. Markov model vs. hybrid model
- Population-based vs. individual-based vs. microsimulation model

## Model assumptions

- Country of investigation
- HPV types involved
- HPV-induced diseases
- Vaccine coverage rate
- Vaccine efficacy
- Vaccination age
- Male vaccination
- Duration of immunity
- Application of booster
- Levels of sexual activity
- Sexual mixing strategy
- Cervical screening strategy
- Duration of follow-up
- Time step of follow-up

## Research outcome

- Cost-effectiveness analysis
- HPV-prevalence reduction

## Universal HPV vaccination: 26 publications

- 8 reuse methodology
- 8 ordinary differential equation (ODE) models
- 1 static Markov model
- 2 network models
- 3 difference equation models
- 3 hybrid models
- 1 prevalence-based model

## Female-only HPV vaccination: 90 publications

- $\approx 50\%$  reuse methodology
- 25 static Markov models
- 4 microsimulation models
- 3 cohort models
- 1 prevalence-based model
- 2 difference equation models
- 1 network model
- 10 ODE models
- 8 hybrid models

## Universal vaccination

- 8 publications: cost-effective results
- 7 publications: non-cost-effective results
- 11 publications: only HPV prevalence reductions

## Female-only vaccination

- 75 publications: cost-effective results
- 1 publication: non-cost-effective results
- 10 publications: only HPV prevalence reductions
- 4 publications: no research outcomes

## Universal vaccination

- Taira et al.
  - Difference equation model for HPV transmission
  - Static Markov disease progression model
- Kim et al.
  - ODE model for HPV transmission
  - Microsimulation disease progression model
- Horn et al.
  - ODE model for HPV transmission
  - Static Markov disease progression model

- Probabilistic nature
- Exhaustive and mutually exclusive health states
- Moving between health states according to specified transition probabilities
  - Assigning flat and informative distributions with suitable ranges
  - Prior information out of the literature or from expert opinion
  - Updating posterior distributions with available data
  - Propagating parameter uncertainty by Markov Chain Monte Carlo estimations (MCMC)
- Model calibration with age- and gender-specific data on prevalence of HPV infection and induced diseases
- Each health state is assigned a value of utility
  - Ranging between 0 and 1
  - 0 represents death, 1 perfect health
  - Specified with Time Trade-Off (TTO) method
- Health economic analysis of multi cohort HPV vaccination strategy

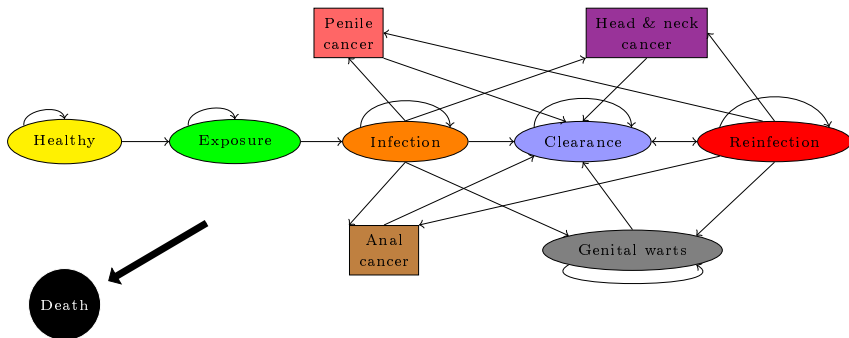
- **Combination of**
  - natural history of disease infection and progression models
  - dynamic sexual disease transmission models
- **Age- and gender-specific HPV prevalence can be calculated beforehand**
  - by means of discrete or continuous time models
  - these probabilities inform the disease progression model afterwards
- **Alternative: the process of sexual mixing can be integrated directly into the static disease progression model**

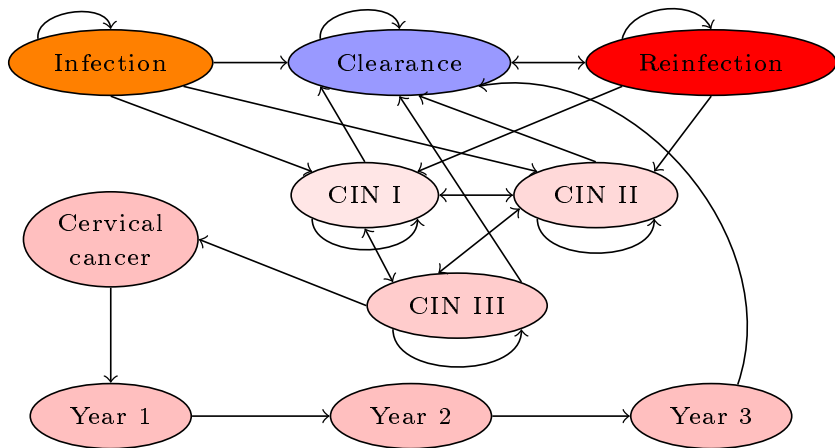
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- **Alternative: the process of sexual mixing can be integrated directly into the static disease progression model**

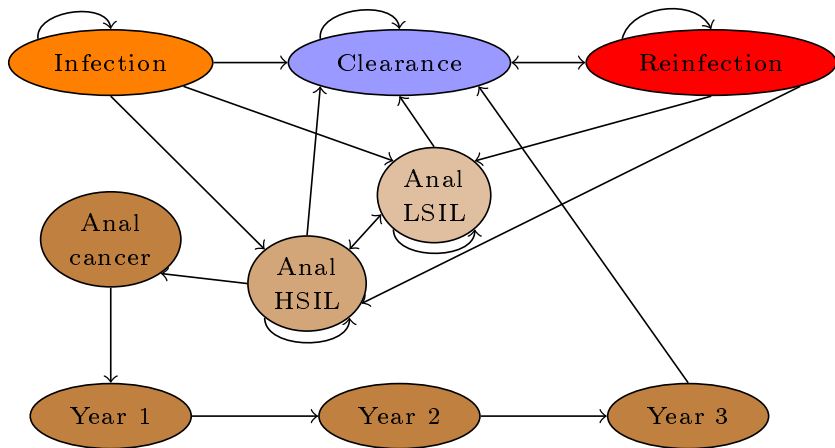


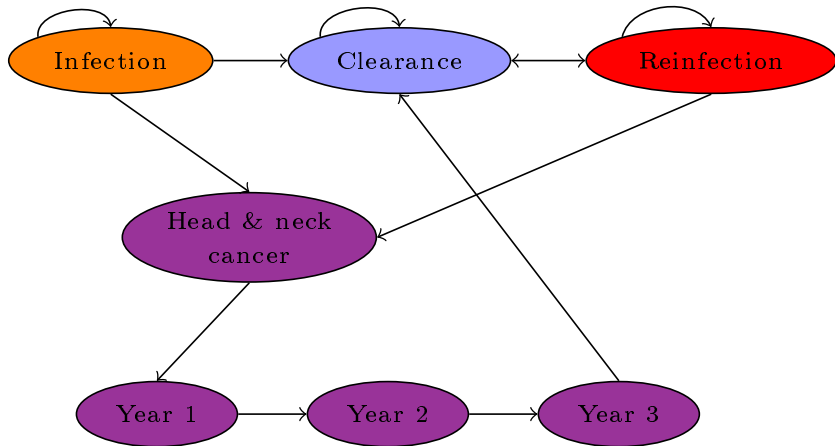
- **Combination of**
  - natural history of disease infection and progression models
  - dynamic sexual disease transmission models
- **Age- and gender-specific HPV prevalence can be calculated beforehand**
  - by means of discrete or continuous time models
  - these probabilities inform the disease progression model afterwards
- **Alternative: the process of sexual mixing can be integrated directly into the static disease progression model**

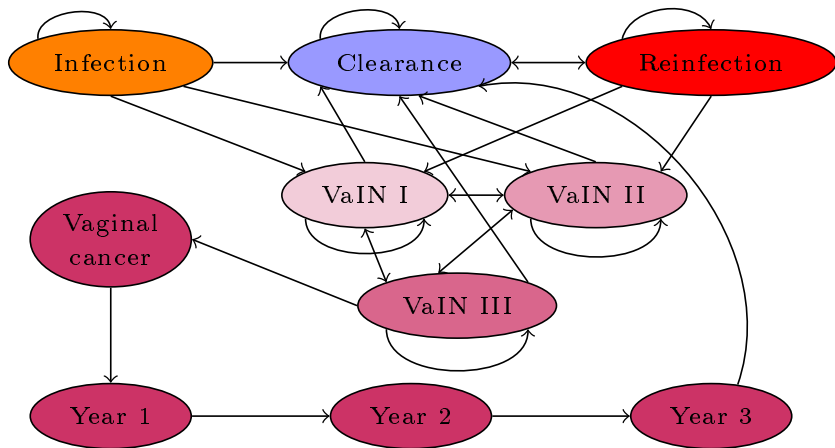
$S_m = 22$  health states

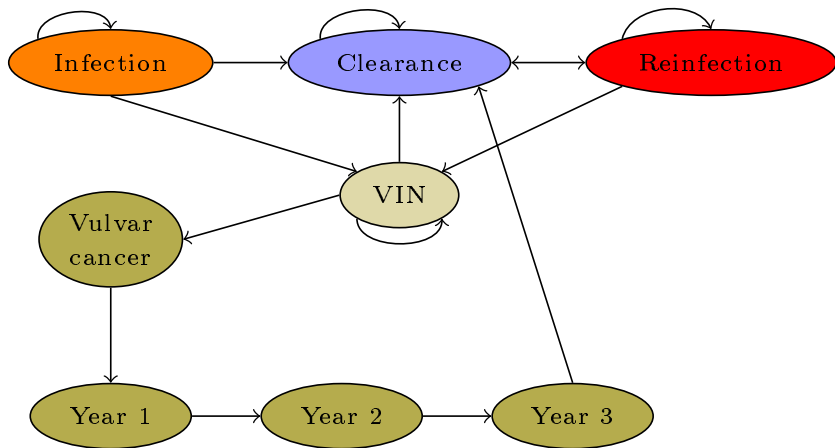


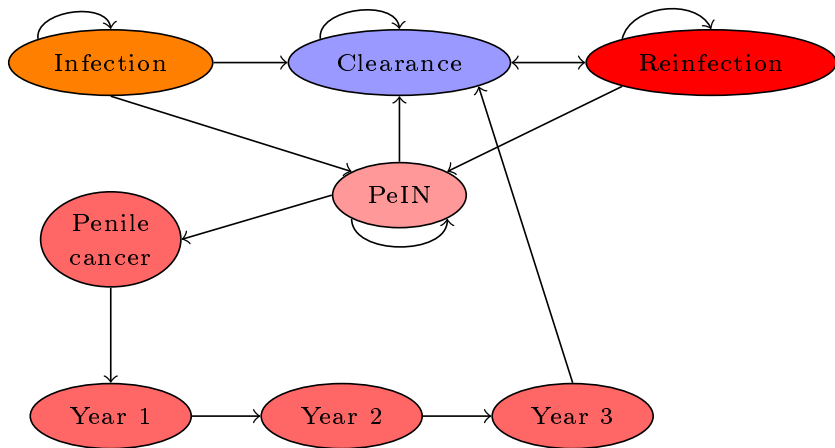














- Individuals move across health states according to  $p_{i,a,j,h}$ , where
  - $i$  indexes the respective health intervention;
  - $a$  indexes the individual's age;
  - $j$  indexes the original health state;
  - $h$  indexes the target health state.
- All transitions from one health state have to sum up to 1 (constraint of probabilities)
- Transitions to the set of health states  $\mathcal{H}$  are possible
- Transitions to health states outside of  $\mathcal{H}$  are set to 0
- Remaining in respective state is induced by  $1 - \sum_{p_{i,a,j,h}} \forall h \in \mathcal{H}$
- Different transition probabilities for females and males as a consequence of different numbers of health states and gender-specific parameters
- Gender-specific parameters with the index  $g = 0$  represent females

- Individuals can have sex (indicated by  $s_a$ ) and move to  $h = 2$
- Individuals can die (indicated by  $d_{a,0}$ ) and move to  $h = 9$
- Individuals can remain in perfect health ( $h = 1$ )

$$p_{i,a,1,h} = 0 \forall h \notin \{1, 2, 9\}$$

$$p_{i,a,1,2} = s_a$$

$$p_{i,a,1,9} = d_{a,0}$$

$$p_{i,a,1,1} = 1 - \sum_{h \neq 1} p_{i,a,1,h}$$

- Individuals in  $j = 2$  can have acquire HPV infection (indicated by  $\lambda_{0,s,a}$ ) and move to  $h = 3$
- Individuals in  $j = 2$  can die (indicated by  $d_{a,0}$ ) and move to  $h = 9$
- Individuals in  $j = 2$  can remain in exposure ( $h = 2$ )

$$p_{1,a,2,h} = 0 \forall h \notin \{2, 3, 9\}$$

$$p_{1,a,2,3} = \lambda_{0,s,a}$$

$$p_{1,a,2,9} = d_{a,0}$$

$$p_{1,a,2,2} = 1 - \sum_{h \neq 2} p_{1,a,2,h}$$

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- Individuals in  $j = 2$  can remain in exposure ( $h = 2$ )

$$p_{2,a,2,h} = 0 \forall h \notin \{2, 3, 9\}$$

$$p_{2,a,2,3} = \alpha_1[\omega_3(1 - \gamma_1)\lambda_{0,s,a} + (1 - \omega_3)(1 - \zeta\gamma_1)\lambda_{0,s,a}] + (1 - \alpha_1)\lambda_{0,s,a}$$

$$p_{2,a,2,9} = d_{a,0}$$

$$p_{2,a,2,2} = 1 - \sum_{h \neq 2} p_{2,a,2,h}$$

- $\alpha_1$  represents the vaccine coverage in female-only vaccination
- $\gamma_1$  represents the vaccine efficacy
- $\omega_3$  represents the vaccine compliance
- $\zeta$  represents the reduction in effectiveness due to noncompliance

Cohort	Time of follow-up												
	0	1	2	3	4	5	6	7	8	9	10	...	55
1	25	26	27	28	29	30	31	32	33	34	35	...	80
2	24	25	26	27	28	29	30	31	32	33	34		79
3	23	24	25	26	27	28	29	30	31	32	33		78
4	22	23	24	25	26	27	28	29	30	31	32		77
5	21	22	23	24	25	26	27	28	29	30	31		76
6	20	21	22	23	24	25	26	27	28	29	30		75
7	19	20	21	22	23	24	25	26	27	28	29		74
8	18	19	20	21	22	23	24	25	26	27	28		73
9	17	18	19	20	21	22	23	24	25	26	27		72
10	16	17	18	19	20	21	22	23	24	25	26		71
11	15	16	17	18	19	20	21	22	23	24	25		70
12*	14	15*	16	17	18	19	20	21	22	23	24	...	69
13*	13	14	15*	16	17	18	19	20	21	22	23		68
14*	12	13	14	15*	16	17	18	19	20	21	22		67
15		12	13	14	15	16	17	18	19	20	21		66
16			12	13	14	15	16	17	18	19	20		65
17				12	13	14	15	16	17	18	19		64
18					12	13	14	15	16	17	18		63
19						12	13	14	15	16	17		62
20							12	13	14	15	16		61
21								12	13	14	15		60
22									12	13	14		59
23										12	13		58
24											12	...	57

- Running 2 chains in parallel to calculate posterior distributions of parameters  $\theta = (\theta_1, \dots, \theta_k)$
- Choosing two different starting points with larger variance compared to the underlying data
- Comparing within-chain variance  $W(\theta_k)$  to between-chain variance  $B(\theta_k)$
- $n_{sims}$  represents the length of the MCMC sample

$$\widehat{Var}(\theta_k|y) = \frac{n_{sims}-1}{n_{sims}}W(\theta_k) + \frac{1}{n_{sims}}B(\theta_k)$$

Convergence is monitored by assessing the *potential scale reduction*

$$\hat{R} = \sqrt{\frac{\widehat{Var}(\theta_k|y)}{W(\theta_k)}}$$

- $\hat{R}$  is the factor by which the scale of the posterior distribution of  $\theta_k$  can be further reduced
- A longer MCMC run will possibly improve convergence
- $\hat{R} \leq 1.1$  represents sufficient convergence

- MCMC iterations are by definition correlated
- Current observation depends on previous one
- The higher the autocorrelation, the lower the equivalence between MCMC output and a proper *iid* sample

$$n_{eff} = \frac{n_{sims}}{1 + 2 \sum_{t=1}^{\infty} corr_t}$$

- $corr_t$  is the *lag t* autocorrelation
- $n_{eff} \approx n_{sims}$  indicates negligible autocorrelation
- In case of high autocorrelation
  - convergence can still be reached
  - extreme quantiles of the posterior distribution are typically estimated without precision



## **Presentation of preliminary results under baseline assumptions**

- vaccination of 12 year old females and males
- high vaccine coverage rate in the catch-up vaccination

## **Detailed explanation of calculation process including**

- overall costs and utilities
- present values of cost (PVC) and utility (PVU)
- *Incremental Cost-Effectiveness Ratio* (ICER)
- cost-effectiveness plane

## Costs include

- diagnostic procedures of health states
  - 2 pap smears and 2 colposcopies in females with CIN I-III
  - 1 HPV DNA test in females with CIN III and cervical cancer
  - anoscopy, biopsy, cytology in individuals with anal LSIL and HSIL
  - diagnostic costs of other HPV-induced diseases already included in treatment costs
- treatment of HPV-induced precancerous lesions and cancers
- vaccine administration and product costs in female-only, universal and catch-up interventions

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**Unit** costs and utilities are multiplied by the number of individuals in intervention  $i$  at time  $t$  in the respective health state, and by the probabilities of diagnosis, to result in **overall** measures.

$$\begin{aligned}
 C_{i,t} = & C_{i,t}^{scr} + C_i^{vac} + C_{i,t}^{gw} + C_{1,i,t}^{cin} + C_{2,i,t}^{cin} + C_{3,i,t}^{cin} + C_{i,t}^{cerv} + C_{i,t}^{lsil} \\
 & + C_{i,t}^{hsil} + C_{i,t}^{an} + C_{i,t}^{hn} + C_{i,t}^{vin} + C_{i,t}^{vulv} + C_{1,i,t}^{vain} + C_{2,i,t}^{vain} + C_{3,i,t}^{vain} \\
 & + C_{i,t}^{pein} + C_{i,t}^{pen}
 \end{aligned}$$

$$\begin{aligned}
 U_{i,t} = & U_{i,t}^{health} + U_{i,t}^{inf} + U_{i,t}^{gw} + U_{1,i,t}^{cin} + U_{2,i,t}^{cin} + U_{3,i,t}^{cin} + U_{r,i,t}^{cerv} + U_{i,t}^{lsil} \\
 & + U_{i,t}^{hsil} + U_{r,i,t}^{an} + U_{r,i,t}^{hn} + U_{i,t}^{vin} + U_{r,i,t}^{vulv} + U_{1,i,t}^{vain} + U_{2,i,t}^{vain} + U_{3,i,t}^{vain} \\
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 \end{aligned}$$



- Standard measure in cost-effectiveness analyses
- Incremental cost per QALY gained
  - **Quality-Adjusted Life Year**
  - Utility of health state is multiplied with amount of time spent within
- All model parameters in vector  $\theta = (\theta^3, \theta^2)$ 
  - $\theta^3$  representing parameters in  $i = 3$  (universal vaccination)
  - $\theta^2$  representing parameters in  $i = 2$  (female-only vaccination)
- Ratio of expectations of cost- and effectiveness-differentials
  - $\Delta_c = \text{PVC}_3 - \text{PVC}_2$
  - $\Delta_e = \text{PVU}_3 - \text{PVU}_2$

$$ICER = \frac{E[\text{PVC}|\theta^3] - E[\text{PVC}|\theta^2]}{E[\text{PVU}|\theta^3] - E[\text{PVU}|\theta^2]} = \frac{E[\Delta_c]}{E[\Delta_e]}$$

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## Positive algebraic sign

- Universal vaccination both higher costs and effects than female-only vaccination
- Universal vaccination both lower costs and effects than female-only vaccination

## Negative algebraic sign

- Universal vaccination higher costs and lower effects than female-only vaccination
- Universal vaccination lower costs and higher effects than female-only vaccination → cost-saving ICER

ICER values between €30,000 and €45,000 are deemed to be cost-effective according to the Italian Health Economics Association (AEIS). In contrast, the NHS in the UK define ICERs under £25,000 to be cost-effective.

## Description of the graph

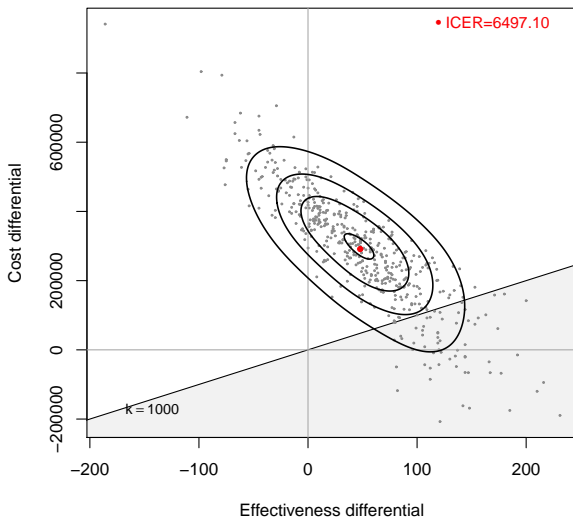
- The x-axis is the effectiveness differential  $\Delta_e$
- The y-axis is the cost differential  $\Delta_c$
- Each point represents a possible future in terms of the expected measures of differential cost and benefit
- The spread of the distribution of points accounts for uncertainty
- The shaded part of the plane indicates the sustainability area (ICERs below the threshold of cost-effectiveness)
- The ICER is displayed as a red dot with its corresponding value

## Interpretation of the graph

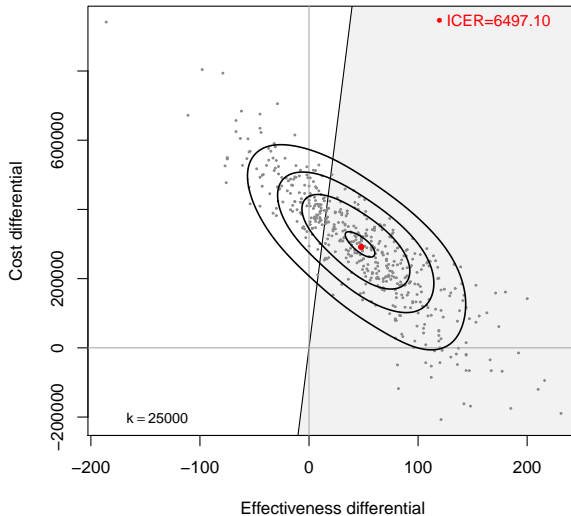
- Points lying in the north-eastern quadrant (i.e. when  $\Delta_e > 0$  and  $\Delta_c > 0$ ) suggest that universal vaccination proves more effective as well as more expensive than female-only vaccination.
- Points lying in the north-western quadrant (i.e. when  $\Delta_e < 0$  and  $\Delta_c > 0$ ) suggest that universal vaccination proves less effective and more expensive than female-only vaccination.
- Points lying in the south-western quadrant (i.e. when  $\Delta_e < 0$  and  $\Delta_c < 0$ ) suggest that universal vaccination proves less effective as well as less expensive than the reference intervention.
- Finally, points lying in the south-eastern quadrant (i.e. when  $\Delta_e > 0$  and  $\Delta_c < 0$ ) suggest that universal vaccination proves more effective and less expensive than female-only vaccination.



## Cost effectiveness plane New Chemotherapy vs Old Chemotherapy



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- **Programming tasks**

- Including layers of uncertainty in the deterministic age- and gender-specific mixing matrices and partner acquisition rates
- Implying the necessity of a booster application
- Generalizing the R code to enable an easier calculation of scenarios next to the baseline
- Conducting a full cost-effectiveness analysis

- **Reading literature** on standard methodology in infectious disease transmission modelling

- **Writing tasks**

- Publishing the cost-effectiveness analysis results, focusing especially on the finding of staggered male vaccination age
- Publishing the methodology of the hybrid Bayesian Markov model
- Writing up the final PhD thesis

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