Data

Bayesian contributions to radiation dose estimation in biological retrospective dosimetry.

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Context				

- Accidents leading to unplanned exposure of humans to ionizing radiation (IR) have occurred many times
 - overexposure in radiotherapy services or occupational settings
 - large-scale nuclear accidents
- Unclear radiation exposure scenarios and/or inconsistent findings
 - workers at risk of exposure may not wear their obligatory personal dosimeter
 - workers at risk of exposure may not store it correctly after use.
- Estimation of the **absorbed radiation dose** received by an exposed or suspected exposed individual may be crucial to:
 - Optimize patient-centered care
 - Predict the derived health consequences for both early and late effects
 - Perform rapid triage of exposed versus non-exposed persons
 - Clarify unclear radiation exposure scenarios
 - Appease the "worried well" persons

Dose assessment \Rightarrow Proof of exposure by court and professional associations

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Biological retrospective dosimetry

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- It offers the only possibility to estimate the individual absorbed dose
 - even weeks or months after a potential exposure (Kulka et al. (2018)).
 - when a direct measurement of IR exposure is not or no longer possible

Main goal

Estimation of the individual absorbed radiation dose from microscope counting of radiation-related chromosomal anomalies

- Radiation exposure causes chromosomal DeoxyriboNucleic Acid (DNA) lesions like double-stand breaks
- $\bullet~$ The broken fragments may repair incorrectly $\Rightarrow~$ Chromosome aberrations



Bayesian contributions

Conclusion & Perspectives

The dicentric chromosome assay (DCA)

- Dicentrics have a low naturally occurring background frequency
- Frequencies of dicentrics increase with the absorbed dose
 ⇒ Well-established and highly specific biological marker of radiation exposure
- Scoring dicentrics in peripheral human blood lymphocytes : "gold standard" biological method for retrospective dose estimation (IAEAb (2011)).



Photo: Olivier Seignette/Mikaël Lafontan/Médiathèque IRSN

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Main que	stions			

Given the number of dicentrics per cell observed in blood lymphocytes:

Question Q1

Can it be stated that a strictly positive radiation dose has been received by :

- In all of the analyzed cells (whole-body irradiation)?
- **3** only a fraction of the analyzed cells (partial irradiation)?
- **o** none of the analyzed cells ? (Relevant for unclear exposure scenarios)

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Main que	stions			

Given the number of dicentrics per cell observed in blood lymphocytes:

Question Q1

Can it be stated that a strictly positive radiation dose has been received by :

- In all of the analyzed cells (whole-body irradiation)?
- **2** only a fraction of the analyzed cells (partial irradiation)?
- **③** none of the analyzed cells ? (Relevant for unclear exposure scenarios)

Question Q2

What is the **estimated absorbed dose** and the **uncertainty associated** to this estimation?



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Standard approaches

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4 real radiation accident victims (2006-2013) In-vivo data provided by IRSN/LRAcc

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Number of dicentrics



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Id	Circum	stances of accident	Clinical signs	Physical dosimetry	Conventional cvtogenetics	
06-11	Exposure to y-r	ays	Vomiting (4h30), nausea, hair loss, Lymphocytes: 0.8 × 10 ⁻³	No	⇒ 👖	no: Number of peripheral blood lymphocytes
11-08	Medical contex to a γ-source (0	t; 10 minutes located next To 60)	Hematopoetic syndrom 7 days after exposure	No		analysed
08-03	Put the γ-sourc his pocket (10 r -> Hand burn	e (lr) in his hand then in ninutes to 1 hour)	lymphocytes: 1.05 × 10 ⁻³	0.25 Sv		Ro: Number
05-03	Exposure head Shoulders 5cms Neck 20cms aw	and chest : 15-30 seconds away from the X source ay from the X source	Erythema (collarbone) Lymphocytes: 2.39 × 10 ^{.3}	0.045 Sv	· ()	chromosomes observed in each cell
s -	06-11	2 -	08-03	2 - T	05-03	
80 -		8 -	8 -	80 -		
7roaddfees 0.4 0.6		Protections	Probabilies 0.4 0.6	Protobiliters 0.6		,

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0.2 0.4 0.6 0.8

0.4 0.6 0.8



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8 suspected exposed individuals (2006-2013) In-vivo data provided by IRSN/LRAcc



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4 real <u>suspected individuals</u> (2006-2013) <u>From</u> IRSN/LRAcc

Id	Circumstances of accident	Clinical signs	Physical dosimetry
06-63	Exposure to γ-rays (10-15 minutes)	No	No
06-70	Spent the night 25 centimeters away from a γ-source	No	No
06-13	Colleague of 06-11	No	No
06-15	Colleague of 06-11	No	No



For some of them, no dicentric was observed ...



In-vitro irradiation of blood samples - various healthy donors - different doses

Number of analyzed cells	Dose (Gray)	Number of dicentrics
19194	0	21
1676	0.05	3
1552	0.10	6
481	0.15	3
1057	0.24	11
1768	0.30	38
1187	0.33	18
2919	0.50	83
1538	0.80	100
869	1	90
1525	1.6	269
1844	2	545
352	2.31	122
784	3	482
534	4	521
341	4.70	381
94	5.77	143



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Dose-response model \mathcal{M}_A for in-vivo data Exposed and suspected exposed individuals

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Let's consider a given individual with n_0 analyzed cells:

- D₀ : Unknown absorbed dose (in Gray) received by each cell
- R_k : Number of dicentrics observed in each cell k ($k = 1, ..., n_0$)

In case of LOW-LET radiation and homogeneous irradiation

$$(\mathcal{M}_A) \qquad R_k \sim^{i.i.d} \mathsf{Poisson}(\lambda_0) \\ \lambda_0 = A + \alpha D_0 + \beta D_0^2$$

Bayesian contributions

- $\theta = (A, \alpha, \beta)$: unknown parameters with A > 0, $\beta > 0$, $\alpha > -2\sqrt{A\beta}$
- A: background expected number of dicentrics per cell at dose $D_0 = 0$

•
$$Y_0 = \sum_{k=1}^{n_0} R_k \sim Poisson(n_0\lambda_0)$$

Non-identifiable model \Rightarrow External data required to estimate $\theta = (A, \alpha, \beta)$



Conclusion & Perspectives

Dose-response model $\mathcal{M}_{\textit{C}}$ for calibration data

Standard approaches

Let's consider a given experimental (in-vitro) irradiation i $\in \{1, \ldots, I\}$

- D_i: Fixed absorbed dose (in Gray) received by each cell
- $Z_{i,l}$: Number of dicentrics observed in each cell $I \in \{1, ..., n_i\}$ at dose D_i

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In case of LOW-LET radiation and homogeneous irradiation

At a given dose D_i :

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$$(\mathcal{M}_{C}) \qquad Z_{i,l} \sim^{i.i.d} \mathsf{Poisson}(\lambda_{i}) \\ \lambda_{i} = \mathsf{A} + \alpha \mathsf{D}_{i} + \beta \mathsf{D}_{i}^{2}$$

 $\Rightarrow Y_i = \sum_{l=1}^{n_i} Z_{i,l} \sim Poisson(n_i \lambda_i)$

where Y_i is the total number of dicentrics observed at dose D_i and n_i the total number of analyzed cells

 \bullet Fit $\mathcal{M}_{\textit{C}}$ to calibration data using maximum likelihood estimation

• Plug
$$\hat{\theta} = (\hat{A}, \hat{\alpha}, \hat{\beta})$$
 into $\mathcal{M}_{\mathcal{A}}$

• Derive point estimate $\hat{D_0}$ of the absorbed dose D_0 (inverse regression)

$$\hat{D_0}=g(\hat{A},\hat{lpha},\hat{eta})=rac{-\hat{lpha}+\sqrt{\hat{lpha}^2+4\hat{eta}(\hat{\lambda_0}-\hat{A})}}{2\hat{eta}}$$

where $\hat{\lambda_0} = \frac{Y_0}{n_0}$



Standard approaches

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Answering Q_2 - Estimation of the dose

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Id	Circumstances of accident	MLE for the dose D ₀	Id	Circumstances of accident	MLE for the dose D ₀
06-11	Exposure to γ -rays	4.40	06-13	Colleague of 06-11	0.02
11-08	Medical context; 10 minutes located next to a γ-source (Co 60)	1.88	06-14	Colleague of 06-11	0.02
08-03	Put the γ-source (lr) in his hand then in his pocket (10 minutes to 1 hour) -> Hand burn	0.23	06-15	Colleague of 06-11	-0.03
05-03	Exposure head and chest : 15-30 seconds Shoulders 5cms away from the X source Neck 20cms away from the X source	0.11	06-16	Colleague of 06-11	0.02
06-63	Exposure to γ -rays (10-15 minutes)	0.15	04-14	Positive dosimeter	-0.03
06-70	Spent the night 25 centimeters away from a γ-source	0.25	13-09	Positive dosimeter	-0.03

Potential drawbacks:

- If $\hat{\lambda_0} = \frac{Y_0}{n_0} = 0$ then $\hat{D}_0 < 0$ (Context: Small signal in the data)
- Prior information on the dose not accounted for
- Modular approach : Disjoint estimation of θ and D_0



• Approach 1: Multivariate delta-method

$$\begin{split} \sigma_{\hat{D}_{0}}^{2} &= \sigma_{\hat{A}}^{2} \left(\frac{\partial g}{\partial A}\right)_{A=\hat{A}}^{2} + \sigma_{\hat{\alpha}}^{2} \left(\frac{\partial g}{\partial \alpha}\right)_{\alpha=\hat{\alpha}}^{2} + \sigma_{\hat{\beta}}^{2} \left(\frac{\partial g}{\partial \beta}\right)_{\beta=\hat{\beta}}^{2} + \sigma_{\hat{\lambda}_{0}}^{2} \left(\frac{\partial g}{\partial \lambda_{0}}\right)_{\lambda_{0}=\frac{Y_{0}}{r_{0}}}^{2} \\ &+ 2 \left(\frac{\partial g}{\partial A}\right)_{A=\hat{A}} \left(\frac{\partial g}{\partial \alpha}\right)_{\alpha=\hat{\alpha}} \operatorname{cov}(\hat{A}, \hat{\alpha}) + 2 \left(\frac{\partial g}{\partial \alpha}\right)_{\alpha=\hat{\alpha}} \left(\frac{\partial g}{\partial \beta}\right)_{\beta=\hat{\beta}} \operatorname{cov}(\hat{\alpha}, \hat{\beta}) \\ &+ 2 \left(\frac{\partial g}{\partial A}\right)_{A=\hat{A}} \left(\frac{\partial g}{\partial \beta}\right)_{\beta=\hat{\beta}} \operatorname{cov}(\hat{A}, \hat{\beta}) \end{split}$$

 \Rightarrow Asymptotical 95% confidence interval on dose estimate: $\hat{D}_0 \pm 1.96\hat{\sigma}_{D_0}$

Approach 2: Bootstrap

Potential drawbacks:

- Is the asymptotic assumption correct?
- Bootstrap \Rightarrow Strong data redundancy if small signal in data
- Uncertainty on the dose estimation may depend on the statistical method used to compute the confidence interval

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Hypothesis testing: $H_0: D_0 = 0$ vs $H_1: D_0 = d_1$ (with $d_1 > 0$)

- Test statistic: $Y_0 = \sum_{k=1}^{n_0} R_k$
- Under H_0 , $Y_0 \sim Poisson(n_0A)$
- Critical region: [y₀^{*}, +∞] with y₀^{*} = 0.95 quantile of Poisson(n₀Â)
 y₀^{*} is called "Decision threshold"
- If $y_0^{obs} > y_0^*$, H_0 is rejected with error (of the first kind) = 0.05
- Statistical power: 1 Frd_{H1}(y₀^{*}) where Frd_{H1} cumulative distribution function of a Poisson distribution with intensity = n₀(Â + âd₁ + βd₁²)
 - Detection Limit: The smallest value of dose *d*₁ from which the statistical power of the test is greater or equal to 0.95

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Answering Q1 - Strictly positive absorbed dose received?



$\mathsf{DL}=\mathsf{Detection}\ \mathsf{Limit}$

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Potential drawbacks:

- Binary answer to Q_1 : Rejection of H_0 or not
- D₀ is unknown ! : Statistical power?
- The statistical power may be very small for small doses $D_0...$
- Uncertainty on the estimation of the background expected number of dicentrics per cell A not accounted for
- Does not allow to test if **only a fraction** of the analyzed cells have received a strictly positive radiation dose



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Aim of the	work			

• Can Bayesian statistical methods offer relevant alternative answers to questions Q_1 and Q_2 in biological retrospective dosimetry ?

- To account for expert knowledge when assigning a prior distribution on the unknown absorbed dose *D*₀
- To propose a **unique**, **flexible and coherent framework** allowing to simultaneously answer to questions Q_1 and Q_2



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Approach 1: the one previously described.... Directed Acyclic Graph of the full model $(M_A + M_C)$



- $\theta = (A, \alpha, \beta)$: shared parameters
- \bullet Possibility for the in-vivo data to be accounted for when fitting A, $\alpha,\,\beta$

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• The Bayesian framework allows fitting this model in one step

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The prior distributions

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- $A \sim Unif[0, +\infty[$
- $\alpha \sim Unif[-2\sqrt{A\beta}, +\infty[$
- $\beta \sim \textit{Unif}[0, +\infty[$
- Prior probability distribution on D_0
 - $D_0 \sim Unif(0, 10) \Rightarrow$ Vague prior
 - $D_0 \sim Gamma(a, b) \Rightarrow$ Informative prior

• Hyperparameters a and b of the Gamma prior may be fixed by expert knowledge given the accident scenario

Id	Circumstances of accident	Clinical signs	Physical dosimetry	Prior distribution on D_0
06-11	Exposure to γ-rays	Vomiting (4h30), nausea, hair loss, Lymphocytes: 0.8 × 10 ^{.3}	No	D_0 .median=2.5 D_0 max = 10 (q99-10) D_0 -Gamma(a=1.98 , b=0.66)
11-08	Medical context; 10 minutes located next to a γ- source (Co 60)	Hematopoetic syndrom 7 days after exposure	No	D_0 .median=2.5 D_0 max = 10 (q99~10) D_0 -Gamma(a=1.98 , b=0.66)
08-03	Put the γ-source (lr) in his hand then in his pocket (10 minutes to 1 hour) -> Hand burn	lymphocytes: 1.05 × 10 ⁻³	0.25 Sv	D ₀ .median=0.25 D ₀ max = 5 (q99-5) D ₀ -Gamma(a=0.4, b=0.6)
05-03	Exposure head and chest : 15- 30 seconds Shoulders 5cms away from the X source Neck 20cms away from the X source	Erythema (collarbone) Lymphocytes: 2.39×10 ⁻³	0.045 Sv	D ₀ .median=0.045 D ₀ max = 5 (q99-5) D ₀ -Gamma(a=0.2, b=0.44)

For individuals for which no clinical sign was observed: $D_0 \sim Unif(0,2)$ \Rightarrow Not enough informative ! To improve!



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 Answering
 Q2
 - Bayesian estimation of the dose
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MCMC algorithm - Package R "rjags"



GUM= Multivariate Delta-Method

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BUT				

Given the prior distribution assigned to D_0 , we are assuming that $D_0 > 0$

\Rightarrow Is this assumption relevant for all the considered individuals?



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Answering Q_1 and Q_2 under the Bayesian framework

Question Q₁

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Can it be stated that a strictly positive radiation dose has been received by :

- **1** all of the analyzed cells (whole-body irradiation)?
- **3** only a fraction of the analyzed cells (partial irradiation)?
- **one** of the analyzed cells ? (Relevant for unclear exposure scenarios)

The above sub-questions 1 and 3 can be formalized as :

A Bayesian model selection problem				
$\mathcal{M}_0: R_k \sim^{i.i.d} \textit{Poisson}(A)$	vs	$\mathcal{M}_{A}: \textit{R}_{k} \sim^{i.i.d} \textit{Poisson}(\textit{A} + \alpha\textit{D}_{0} + \beta\textit{D}_{0}^{2})$		
given in-vivo data and calibr	ation da	ata following model $\mathcal{M}_{\mathcal{C}}$ $(\mathcal{D}_0>0)$		

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Answering Q_1 and Q_2 under the Bayesian framework





Credible interval at 95%

[0,0006; 0.0015]

[0.032; 0.048]

[0.044: 0.052]

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 \Rightarrow A Bayes factor (Jeffreys, 1939) can be efficiently approximated (e.g., Monte-Carlo estimate)

But what about sub-question 2 about partial irradiation?



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Idea: using a mixture model (Kamary et al. (2014) - arXiv)

Let's consider a given individual - potentially exposed - with n_0 analyzed cells:

- p_0 : unknown probability for each cell to have received a dose > 0
- D₀ : unknown absorbed dose (in Gray) received by each irradiated cell

A mixture model for in-vivo data (LOW LET + homogeneous irradiation)

 $\mathcal{M}_{\textit{mix}}$: $R_k \sim^{i.i.d} (1 - p_0) \textit{Poisson}(A) + p_0 \textit{Poisson}(A + \alpha D_0 + \beta D_0^2)$

- $D_0 > 0$ and $p_0 \in [0, 1]$
- $\theta = (A, \alpha, \beta)$: unknown parameters with A > 0, $\beta > 0$, $\alpha > -2\sqrt{A\beta}$
- A: common parameter shared by both mixture components
- p0 can also be interpreted as the proportion of irradiated cells
- D_0 and p_0 assumed to be identical for each irradiated cell
- $\bullet~\mathcal{M}_0$ and \mathcal{M}_A are very special cases of the mixture model





- $\theta = (A, \alpha, \beta)$: shared parameters
- The Bayesian framework allows fitting this model in one step



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• If $p_0 = 0$, model \mathcal{M}_0 is selected given the available count data

- \Rightarrow Response to Q_1 is NO= "There is no evidence that a strictly positive radiation dose has been received".
- If $p_0 = 1$, model M_A is selected given the available count data
 - \Rightarrow Response to Q_1 is YES= "A strictly positive radiation dose has been received by all the analyzed cells".
- If $p_0 \in]0, 1[$, neither model \mathcal{M}_0 nor model \mathcal{M}_A is selected given the available count data
 - \Rightarrow Response to Q_1 is YES= "A strictly positive radiation dose has been received BUT only by a fraction of the analyzed cells" (partial body exposure).
 - The fraction of the body irradiated is defined as (IAEA report 2001):

$$F_0 = \frac{p_0 \times exp(D_0/\tilde{D})}{(1-p_0) + p_0 \times exp(D_0/\tilde{D})} \qquad \tilde{D} \sim \textit{Unif}(2.7, 3.5)$$

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- Posterior distribution on $p_0 \Rightarrow$ Probabilistic answer to Q_1
- $\bullet\,\,\Rightarrow\,$ Decision criterion to define the range of acceptance, rejection and indecision conclusions



- Let's c_1 , c_2 , U be fixed decision thresholds (to calibrate by simulation)
- Compute $\pi_1 = P(p_0 > c_1 | Y_i, R_k)$ and $\pi_2 = P(p_0 < c_2 | Y_i, R_k)$
 - If $\pi_1 > U \Rightarrow YES=$ "There is strong evidence that a strictly positive radiation dose has been received by all of the analyzed cells".
 - If $\pi_2 > U \Rightarrow NO=$ "There is no evidence that a strictly positive radiation dose has been received".
 - Else YES= "A strictly positive radiation dose has been received **BUT** only by a fraction of the analyzed cells" (partial body exposure).

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The prior distributions

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- $A \sim Unif[0, +\infty[$
- $\alpha \sim Unif[-2\sqrt{A\beta}, +\infty[$
- $\beta \sim \textit{Unif}[0, +\infty[$
- $D_0 \sim \textit{Gamma}(a, b)$ or $D_0 \sim \textit{Unif}(0, 10)$
- p₀ ∼ Beta(c, d)
- Hyperparameters a,b,c,d may be fixed by expert knowledge given the accident scenario
- Default choice (Rousseau and Mengersen (2011)): c=0.5,d=0.5

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Bayesian inference

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Adaptive Metropolis-Hastings algorithm

- Block updating for (A, α, β) using a Gaussian random walk (20% acceptation rate)
- Gaussian random walk for D_0 (40% acceptation rate)
- For the mixture weight p₀:
 - Iteration t: Independent proposal $\Rightarrow p_0^{cand} \sim Beta(0.5, 0.5)$
 - Iteration t+1: Random walk $\Rightarrow p_0^{cand} \sim Beta(1 + p_0^t, 2 p_0^t)$
 - 40% acceptation rate
- Implemented in Python (2.7.10) (100000 iterations = 30 seconds)

Asymptotic consistency of the proposed mixture testing procedure

- Proved by Kamary et al. (2014) in the specific case of embedded mixture components
 - "If one model is indeed correct, the posterior medians of the corresponding weight in the mixture settles very quickly near the boundary values of 1 as the sample size increases"

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Remark				

 Equivalent formulation of M_{mix} pointing out the latent allocation variables
 M_{mix}: R_k ∼ⁱ Poisson(λ_k) with λ_k = A + αD_{0k} + βD_{0k}²
 D_{0k} = γ_k × D₀ with γ_k ∼ Bern(p₀)

 Easy implementation in WinBUGS or JAGS but inefficient Gibbs sampler!!!



Convergence diagnostics on the weight p_0

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Gibbs sampler (Left) vs Adaptive Metropolis-Hastings (Right)



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Posterior statistics, Bayes factor and posterior probability of \mathcal{M}_1

		Bayes Factor M ₁ vs M ₀ (Kass & Raftery (1995)	P(M ₁ y)				
	D₀ posterior median 95%Cl	p₀ posterior median 95%Cl	F ₀ posterior median 95%Cl	P(p ₀ >0.8)	P(p ₀ <0.2)		
06-11	4.61 [4.14; 5.19]	0.91 [0.76,1.00]	0.97 [0.90; 1.00]	0.93	0.0	+∞ (very strong)	1 [1.0; 1.0]
11-08	2.09 [1.76; 2.69]	0.84 [0.56; 1.00]	0.90 [0.69; 1.00]	0.60	0.0	1.75 ^e +185 (very strong)	1 [1.0; 1.0]
08-03	0.32 [0.15; 1.25]	0.67 [0.10; 1.00]	0.69 [0.11; 1.00]	0.39	0.11	>10 [*] 7 (very strong)	1 [1.0; 1.0]
05-03	0.13 [0.0002; 1.29]	0.54 [0.011; 1.0]	0;55 [0.01; 1.0]	0.31	0.25	4 (Positive)	0.67 [0.63; 0.70]
06-63	0.47 [0.08; 1.84]	0.23 [0.02; 0.99]	0.26 [0.02; 0.99]	0.16	0.46	8.3 (Positive)	0.86 [0.83; 0.88]
06-70	0.55 [0.16; 1.84]	0.36 [0.04; 1.00]	0.40 [0.06; 1.00]	0.21	0.33	303.03 (Very Strong)	1.00 [1.0; 1.0]

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Comparison of dose estimations

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 $\begin{array}{l} \mbox{Posterior medians} + 95\% \mbox{ credible intervals} \\ \mbox{ISO2014} = \mbox{Multivariate Delta Method} \end{array}$

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Prior probability distribution on p_0 : Beta(0.5,0.5)



From left to right : Victims 06-11 (Estimated dose: 4.61 Gy), 08-03 (Estimated dose: 0.32Gy), 05-03 (Estimated dose: 0.13Gy)

- Weak influence of the prior choice on D_0 (results not shown)
- Lack of information in the data to infer p_0 especially when dose is small \Rightarrow More data needed to infer p_0 (and then answer Q_1)?

Sensitivity to the prior choice on p_0

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• Informative Beta priors defined from expert knowledge

Standard approaches

• Jeffrey's prior Beta(0.5,0.5)



Bayesian contributions

Posterior distribution on the dose D_0

Posterior distribution on the weight p_0

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 \Rightarrow Sensitivity is clearly present but should naturally vanish as the number \cite{IRSN} of analyzed blood lymphocytes increases

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Conclusion & Perspectives

2 Data

3 Standard approaches

Bayesian contributions

5 Conclusion & Perspectives

Introduction	Data 000	Standard approaches	Bayesian contributions	Conclusion & Perspectives ●○○
Conclusion	s			

- First fully Bayesian approach proposed to simultaneously answer to two main questions of interest in biological retrospective dosimetry
 - \Rightarrow New insights to the European Radiation Dosimetry (EURADOS) Working Group 10, task 10.6
- Using the proposed mixture model \mathcal{M}_{mix} allows to get rich probabilistic answers to questions Q_1 and Q_2
 - \Rightarrow Relevant input data for decision-making in the contexts of clinical management of patients, rapid triage after large-scale radiation incident, reassuring the 'worried-well'...
- In case of low suspected dose, the number of analyzed blood lymphocytes should be higher to obtain more precise answers to question Q₁

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Introduction	Data 000	Standard approaches	Bayesian contributions	Conclusion & Perspectives ○●○
Perspective	es			

- Simulation studies to validate the whole methodology and calibrate the decision thresholds (c₁,c₂,U)
- Validate the whole methodology from new experimental data for which D₀ and p₀ are known
- Bayesian optimal design to define the number of analyzed cells n_0 required to optimally answer to question Q_1 and Q_2 under budget constraint
- Extend the proposed approach to other chromosome aberrations
- Provide operational tools to dosimetrists

IRSN and and a construction for # Monoral Construction [1] IAEA 2011 report (2011) Cytogenetic dosimetry: applications in preparedness anfor and response to radiation emergencies. International Atomic Energy Agency: Vienna.

[2] Merkle W. (1983) Statistical Methods in Regression and Calibration Analysis of Chromosome Aberration Data. Radiat Environ Biophys. 1:217-233
[3] Ainsbury et al. (2014) Review of Bayesian statistical analysis methods for cytogenetic radiation biodosimetry with a practical example. Radiation Protection Dosimetry. 162(3):185-96

[4] Higueras et al. (2016) A new Bayesian model applied to cytogenetic partial body irradiation estimation. Radiation Protection Dosimetry. 168(3):330-6
[5] Kamary K., Mengersen K., Robert CP., Rousseau J. (2014) Testing hypotheses via a mixture estimation model. ArXiv:1412.2044v2

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