

Modelling cancer risk – uneven outcomes

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Breast cancer risk modelling



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- Alex Jose (HWU)

Outline

- 1 Cancer rates trends over time
 - mainly all-cancer, lung, breast cancer
- 2 Stochastic modelling for incidence (& mortality) rates
- 3 Variation by region and deprivation
- 4 Projection into the future
- 5 Impact of diagnosis delays on mortality
 - also linked to delays relating to Covid-19
- 6 Deep learning methods for cancer rates

Cancer data

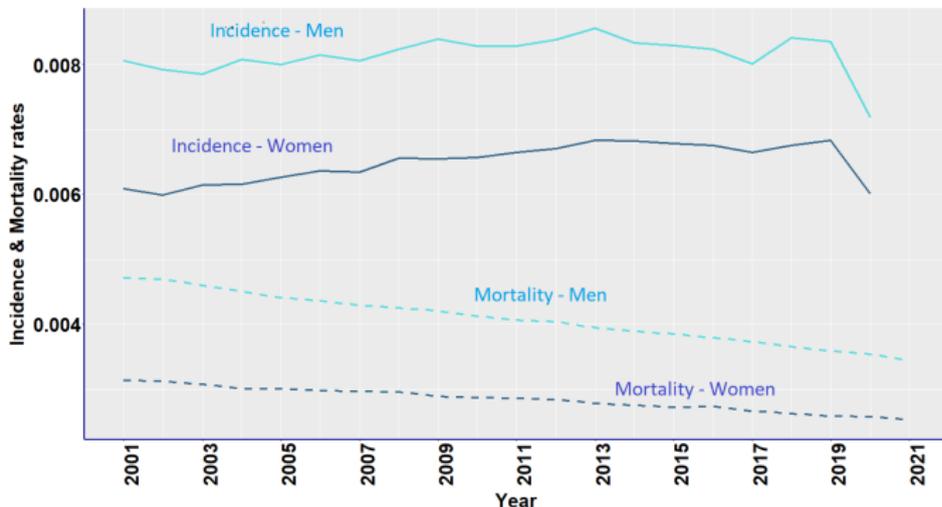
Cancer incidence and deaths data

England: Office for National Statistics (ONS)

- Age groups: 0, 1-4, 5-9, ..., 95+
Age-standardised results, based on the European Standard Population (ESP) 2013
- Gender
- Years: 2001 - 2017 (*some up to 2021*)
- Income Deprivation (ID) decile
1: most deprived; 10: least deprived
- Regions of England: North East, North West, Yorkshire and the Humber, East Midlands, West Midlands, East, London, South East and South West

Trend over time: 2001-2021

All-cancer incidence, mortality
Age standardised rates (no modelling)



Increasing trends for incidence

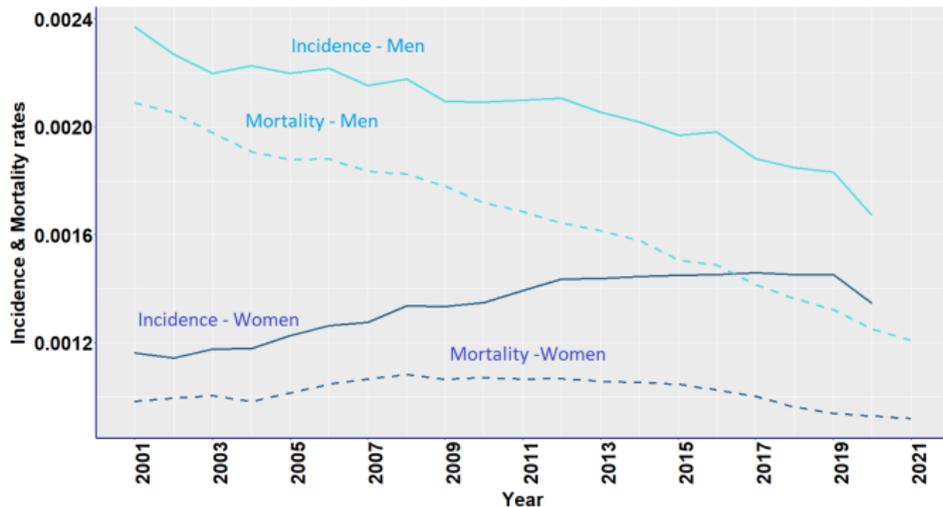
Decreasing mortality trends

Higher rates for men

Notable exception in trend:

Lung cancer, 2001-2021

Age standardised rates (no modelling)



Decreasing
incidence
for men

Increasing
for women

Mortality
relatively
close to
morbidity

Regional and/or socioeconomic differences in cancer rates?

- How big is the gap?
- Is it getting better? Worse?

We need modelling - to account for uncertainty and noise.

Stochastic modelling

- Stochastic modelling for cancer rates



- Transition characterised by underlying rate $\theta_{g,r,a,d,t}$
- $\theta_{g,r,a,d,t}$ depending on **gender, region, age, deprivation, time**
- Quantify uncertainty (probability intervals)

Bayesian models for incidence and mortality rates

$$C_{a,t,d,g,r} \sim \text{Poisson}(\theta_{a,t,d,g,r} E_{a,t,d,g,r})$$

$$\theta_{a,t,d,g,r} \sim \text{Lognormal}(\mu_{a,t,d,g,r}, \sigma^2)$$

$$\mu_{a,t,d,g,r} = \beta' \mathbf{X}$$

$$\beta\text{'s} \sim \text{Normal}(0, 10^4) \quad [\text{vague priors for risk factor effects}]$$

$$\sigma^2 \sim \text{Inv.Gamma}(1, 0.001)$$

- $C_{a,t,d,g,r}$: number of cancer registrations/deaths at **age** a , in **year** t , for **gender** g , **deprivation** level d and **region** r
- $E_{a,t,d,g,r}$: mid-year population estimates
- $\theta_{a,t,d,g,r}$: incidence/mortality rates
- \mathbf{X} : vector of covariates: **age**, **year**, **deprivation**, **gender**, **region**, average age-at-diagnosis + appropriate interaction(s)
- β : vector of coefficients

Also: change-point analysis, variable selection



Change points

- Allow change point(s) in time trends (and age)
 - E.g. different trend after new health/screening policy introduced
 - or after a certain age
- Change point analysis, based on BIC, is considered for detection of changes

$$\mu_{a,t,d,g,r} = \beta_0 + \beta_1 t + \beta_2 (t - \epsilon) \mathbf{I}(t \geq \epsilon)$$

with β_2 : change in trend after time point ϵ .

E.g.
$$\mu_{a,t,d,g,r} = \beta_0 + \beta_1 \text{ year} + \dots$$

may become

$$\mu_{a,t,d,g,r} = \beta_0 + \beta_1 \text{ year}_{<2006} + \beta_2 \text{ year}_{\geq 2007} + \dots$$

Model selection

- Bayesian variable selection methodology
- Chooses the **best** model for

$$\mu_{a,t,d,g,r} = \beta' \mathbf{X}$$

according to *marginal likelihood & Bayes factors*:

$$B_{jk} = \frac{\Pr(D|M_j)}{\Pr(D|M_k)}; j \neq k$$

or *deviance information criterion*:

$$\text{DIC} = -2E_{\beta|D}(\log f(D|\beta)) + 2 \log f(D|\hat{\beta}),$$

Initial findings and main trends (Arik et al, 2020)

Variable selection:

- All-cancer and *life-style* cancers, i.e. lung and bowel cancer: all main variables (age, time, deprivation, gender, region) are important
- Breast and prostate cancer mortality: deprivation is **not** important

Initial findings and main trends (cont.)

How do various factors affect rates? (in general ...)

- Age: higher rates at older ages
- Time:
 - higher incidence in more recent years
 - lower mortality
- Gender: higher rates for men
- Region?

Deprivation?

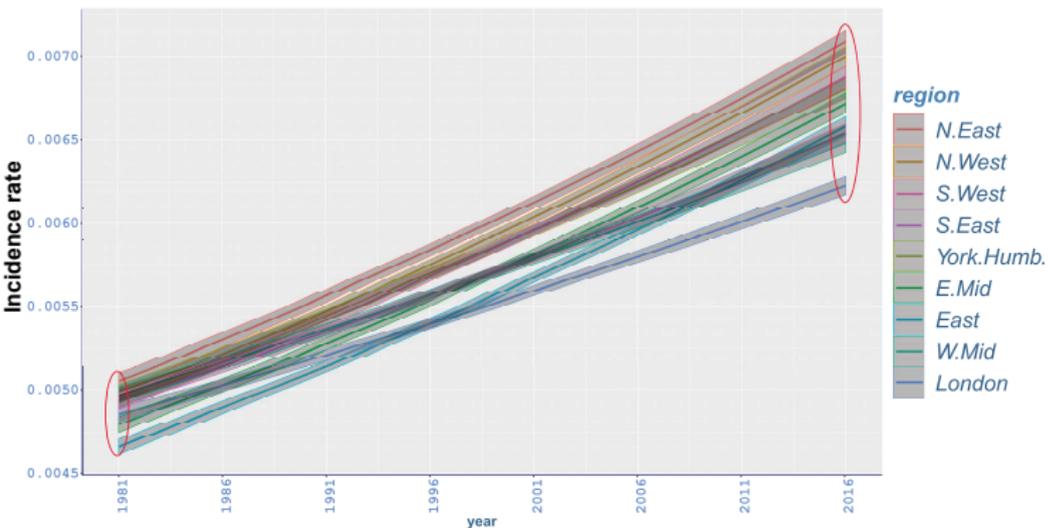
Regional variation in cancer rates?



- Is there a geographical pattern?
- Does variation change over time?
- Is variation the same for different types of cancer?

Regional variation

All cancer incidence – Females, 1981-2016



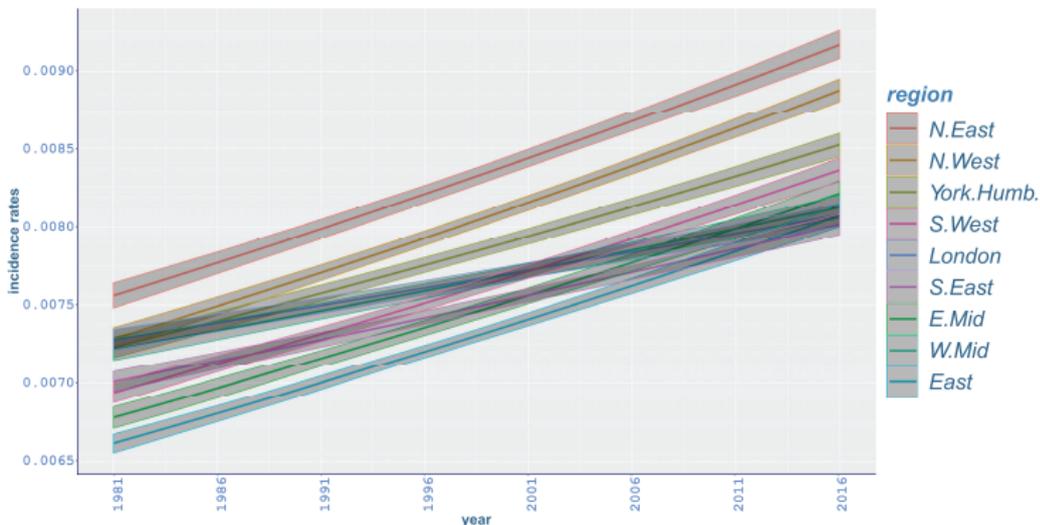
Increasing trend in all regions

Higher incidence in north

Gap widening with time

Regional variation

All cancer incidence – Males, 1981-2016



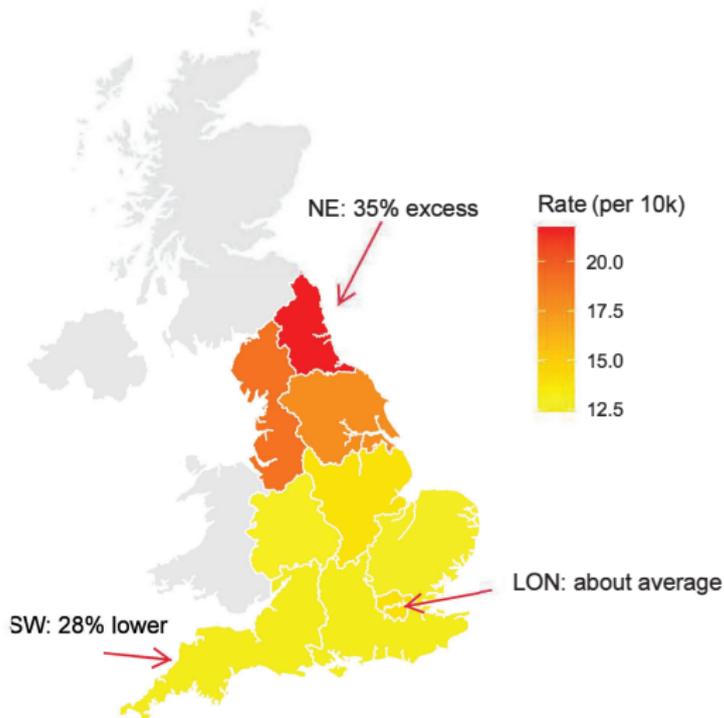
Higher incidence in north

Rates are higher than for women

Gap **not** widening for men

Regional effect

Lung cancer incidence – Females, 2017

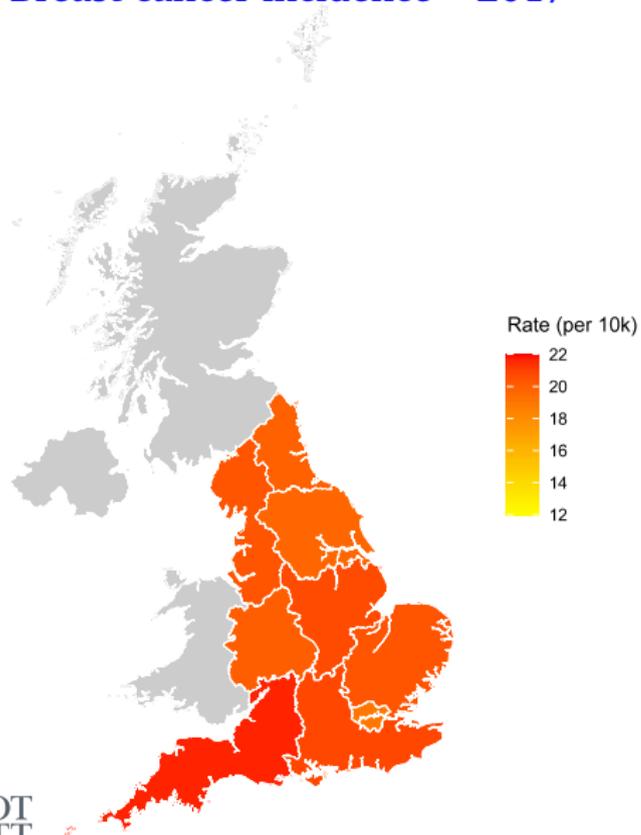


Regional
effect com-
pared to
average

North v.
south?

Regional effect

Breast cancer incidence – 2017



Not a 'life-style' cancer

Regional variation much lower

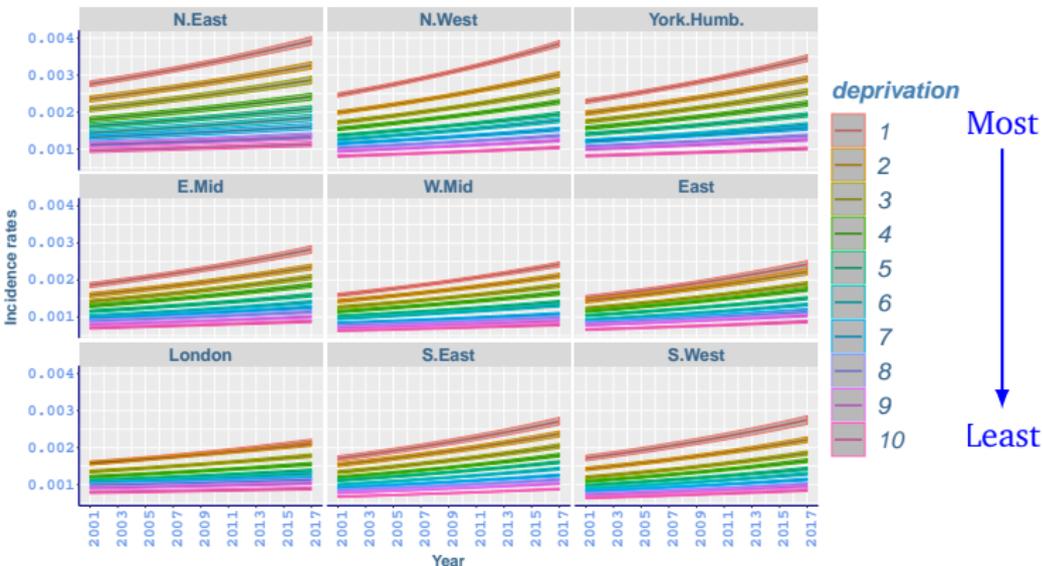
Socioeconomic inequality in cancer rates?



- Use Index of Income Deprivation (ID)
- Deciles: 1 (most deprived), 10 (least deprived)
- For projection (later): quintiles 1 – 5

Deprivation inequality in cancer rates

Lung cancer incidence – Females, 2001-2017



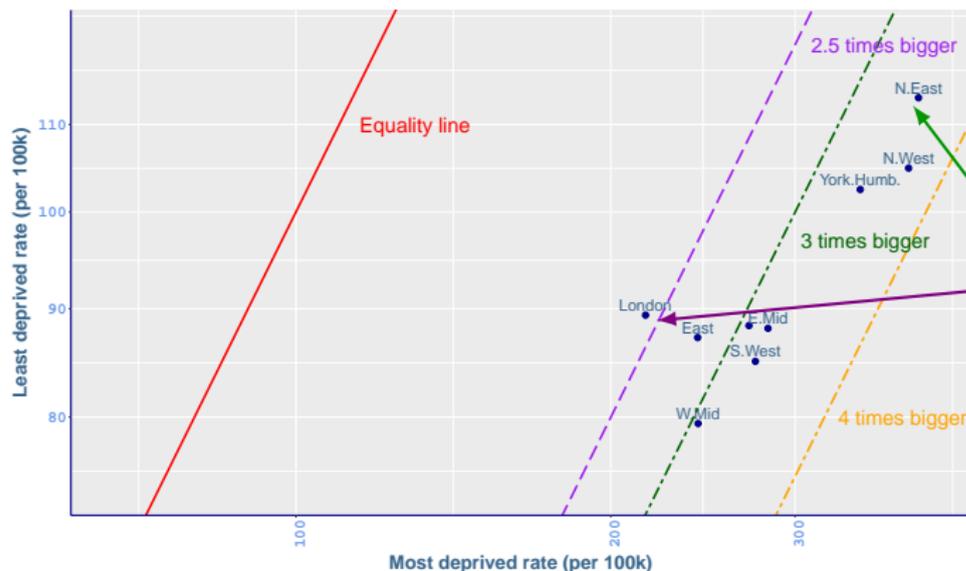
Higher rates for most deprived (1)

Variation by ID getting wider through time

Inequalities more evident in northern regions

Most v. least deprived by region

Lung cancer incidence – Females, 2017



Rates for **most deprived much higher:**

×3.5 N East
×2.5 London

Regional variation

Bayesian forecasting for mortality

$$C_{a,t,d,r} \sim \text{Poisson}(\theta_{a,t,d,r} E_{a,t,d,r})$$

$$\theta_{a,t,d,r} \sim \text{Lognormal}(\mu_{a,t,d,r}, \sigma^2)$$

$$\mu_{a,t,d,r} = \beta_0 + \beta_{1,a} + \beta_{2,t} + \beta_{3,r} + \beta_{4,d} + \beta_5 \text{AAD}_{r,d}$$

$$\sigma^2 \sim \text{Inv.Gamma}(1, 0.1)$$

$$\beta_0, \beta_1, \beta_3, \beta_4 \text{ and } \beta_5 \sim \text{Normal}(0, 10^4),$$

Add random walk with drift for ‘period’ effect:

$$\beta_{2,t} = \text{drift} + \beta_{2,t-1} + \epsilon_t$$

$$\text{drift} \sim \text{Normal}(0, \sigma_{\text{drift}}^2)$$

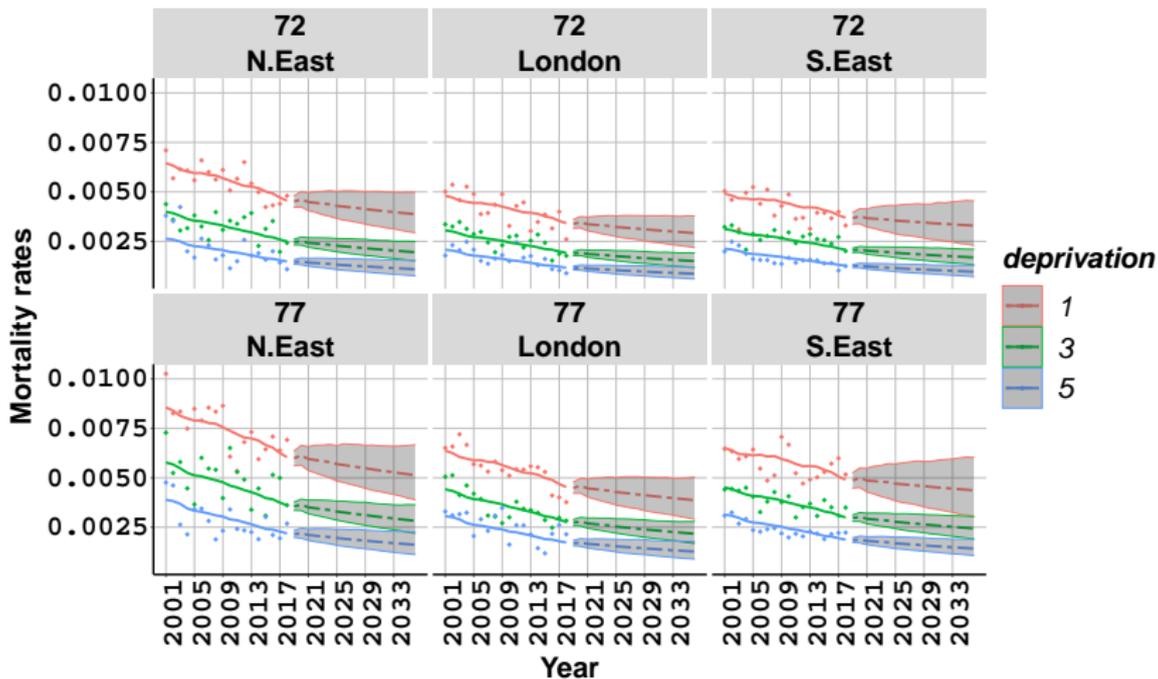
$$\epsilon_t \sim \text{Normal}(0, \sigma_{\beta_2}^2)$$

$$\sigma_{\beta_2}^2 \sim \text{Inv.Gamma}(1, 0.001),$$

for $t = 2001, 2002, \dots, 2018$, where $\hat{\sigma}_{\text{drift}}^2 = \frac{\hat{\sigma}_{\beta_2}^2}{2018-2001} \cdot$

Projected mortality – Lung cancer, 2001 - 2035

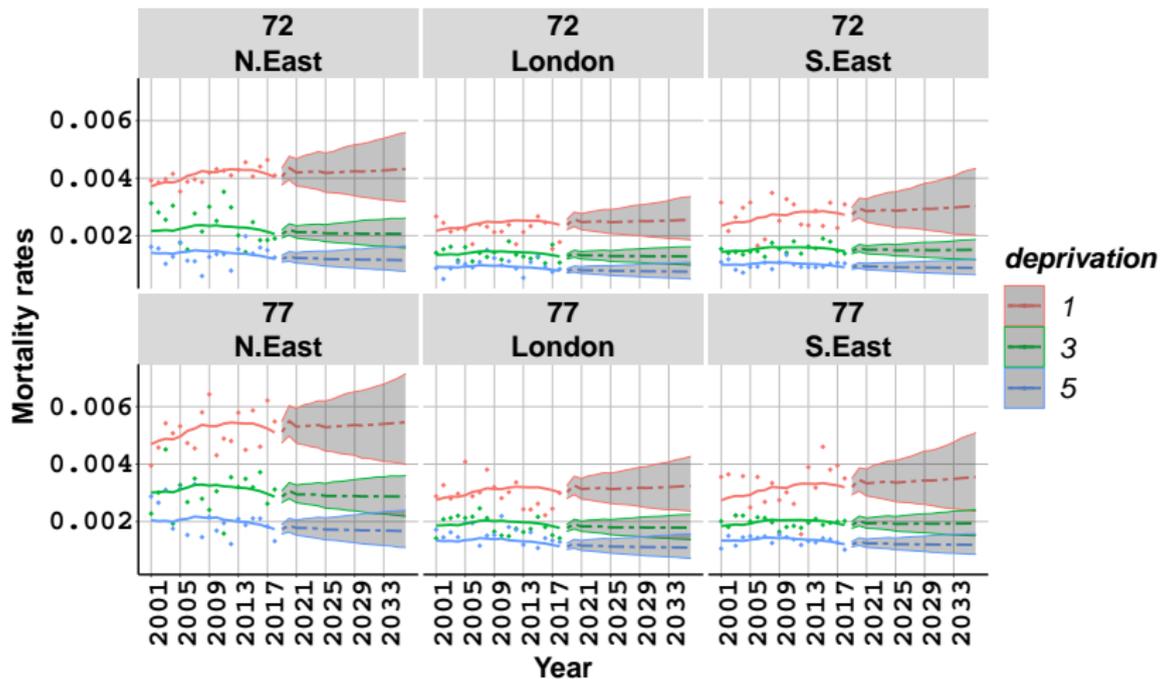
Men 72, 77 yo, deprivation quintiles



- Projected rates for most & least deprived NOT overlapping

Projected mortality – Lung cancer, 2001 - 2035

Women 72, 77 yo, deprivation quintiles



- Mortality for women *NOT* decreasing
- Still rates for most deprived not catching up

Impact of diagnosis delays on mortality



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Covid in Scotland: Cancer diagnoses fell 40% at start of pandemic

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Coronavirus pandemic



The number of people diagnosed with cancer fell by 40% at the start of the Covid pandemic, according to public health statistics.

Public Health Scotland (PHS) figures indicate cancer diagnoses fell by about

- Estimate average age-at-diagnosis (AAD) with incidence data

- Include AAD as risk factor in mortality model

e.g.

$$\mu_{a,t,d,r} = \beta_0 + \beta_{1,a} + \beta_{2,t} + \beta_{3,r} + \beta_{4,d} + \beta_5 \text{AAD}_{r,d}$$

- Estimate impact on mortality

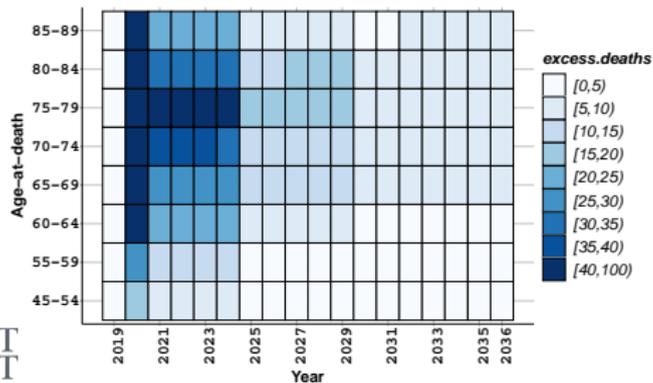
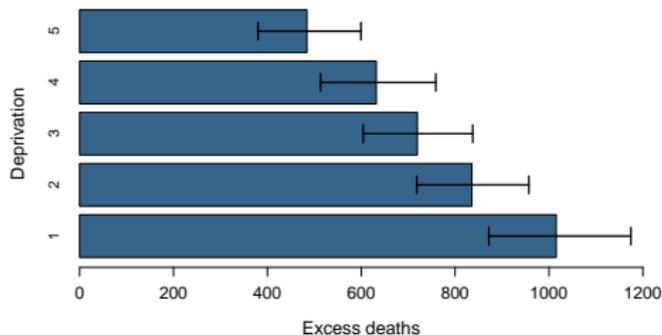
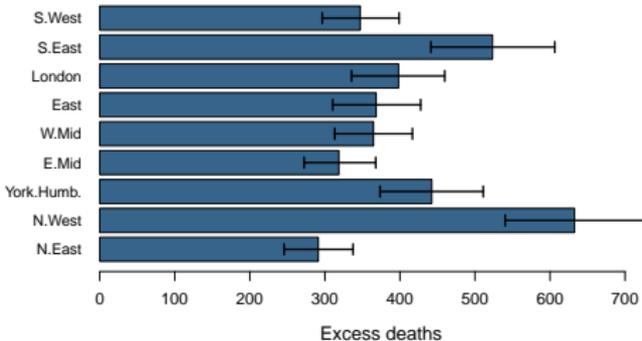
Projected mortality – Lung cancer, 2001 - 2035

Quantify Covid-19 impact on future mortality

- Assume increase in AAD: e.g. 1 month, 3 months etc.
 - Use ONS region future population estimates
 - Assume future deprivation structure unchanged
- Fit Bayesian forecasting model:
 - under no change in AAD (baseline scenario)
 - under 1-month AAD increase (Covid scenario)
 - estimate **excess deaths**

Projected mortality – Lung cancer, women, 2001 - 2035

Excess mortality due to 1-month increase in AAD



Total excess deaths: 3,687

Cancer admissions data (US, 2016-2019)

- Source: Merative (formerly IBM Watson Health)
- **Response:** number of hospital (or similar) admissions
- **Explanatory** variables:

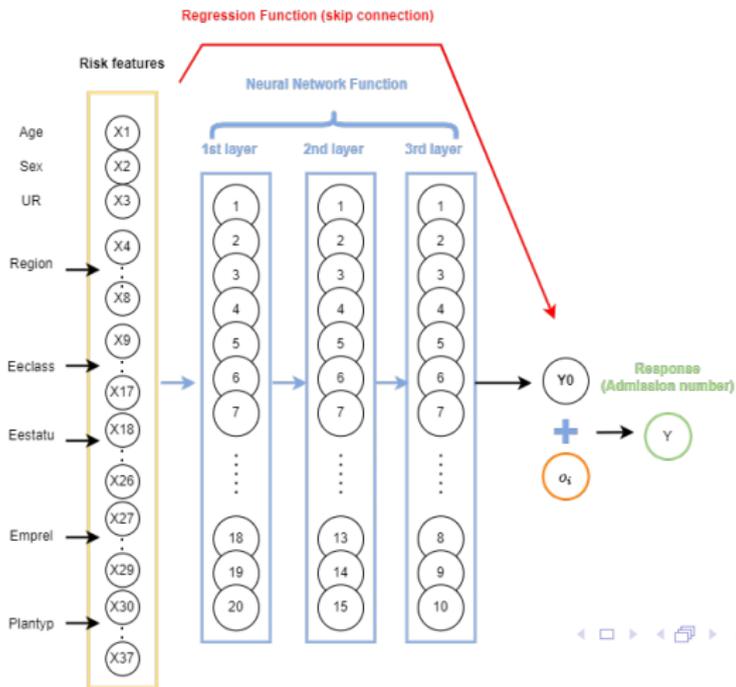
Variable	Description	Type
PLANTYP	Type of plan individual is part of	Factor w/8 levels
AGE	Age of the individual	num 30-65
REGION	Geographical region of residence	Factor w/5 levels
EGEoloc	Geographic location based on postal code	Factor w/53 levels
UR	Urban/rural indicator	Factor w/2 levels
EECLASS	Employee classification	Factor w/9 levels
EESTATUS	Status of employment	Factor w/9 levels
EMPREL	Relation to the primary beneficiary	Factor w/3 levels
SEX	Gender of patient	Factor w/2 levels

- 425,202 records

Artificial Neural Network (ANN) models

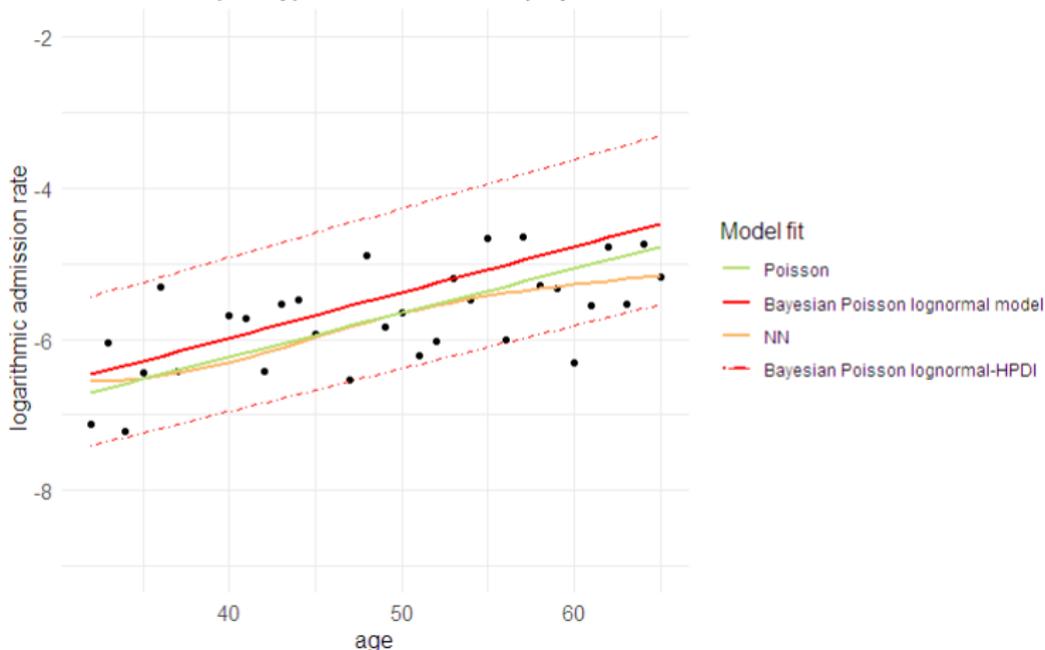
Replace predictor of GLM with **ANN predictor** – Poisson likelihood:

$$\mu^{CNNPoisR}(\mathbf{x}_i) = E_i \exp \left(\underbrace{\langle \boldsymbol{\beta}, \mathbf{x}_i \rangle}_{\text{Regression Function (skip connection)}} + \underbrace{\langle \mathbf{w}^{(d+1)}, (\mathbf{z}^{(d)} \circ \dots \circ \mathbf{z}^{(1)}) \rangle}_{\text{Neural Network Function}}(\mathbf{x}_i) \right)$$



Modelling results – learning data

Logarithmic admission rate for urban- female employee in south
under a PPO plan type-with unknown employee classification & status



NN fit more flexible

Predictive performance: GLM v Bayes v ANN

Table: Average loss over 10-fold validation

Model	Learning loss	Testing loss	Portfolio average
Observed			0.0027
GLM	16.747	16.849	0.0030
Bayes	16.771	16.785	0.0030
NN _{Pois} (20,15,10)	16.378	16.652	0.0027
CANN _{Pois} (20,15,10)	16.475	16.830	0.0027

- 90-10 training-testing split
- NN approach: better predictive performance over testing data
- Followed by Bayesian model

- 1 Regional and socioeconomic gap for cancer rates is widening in the UK
... but not for all cancer types
- 2 Covid-related delays in diagnoses can lead to significant increase in cancer deaths
– also region dependent
- 3 Projection for lung cancer mortality shows persistent deprivation gap
– and significant excess deaths due to covid-like disruptions
- 4 ANNs can provide enhanced rate predictions
– but we need to address interpretability
- 5 Can public health interventions at regional and deprivation level contribute to lower cancer incidence and deaths?

