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Supplement to:
Effectiveness of mammography screening on breast cancer mortality - a study protocol for emulation of target trials using German health claims data

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18 1 Details on the statistical analysis

19 Discrete-time cumulative incidence functions (CIFs) will be estimated using the following approach. Let
 20 A_{never} (reference), A_{once} , and $A_{regular}$ be indicator variables for the screening strategies “never
 21 screened”, “screened at least at baseline”, and “screened at baseline and every two years afterwards”.
 22 The discrete-time (cause specific) hazard is modelled using pooled logistic regression adjusted for
 23 baseline covariates:

$$\begin{aligned}
 24 \quad & \text{logit} \left(\mathbb{P}(Y_{t+1} | \bar{Y}_t = 0, \bar{C}_t = 0, \bar{D}_t = 0, A_{once}, A_{regular}, X) \right) \\
 25 \quad & = f_1(\theta'_1, t) + f_2(\theta'_2, t, A_{once}) + f_3(\theta'_3, t, A_{regular}) + \theta_4 A_{once} + \theta_5 A_{regular} + \theta'_6 X'.
 \end{aligned}$$

26 The above model includes flexible functions $f(\cdot)$ of time t , regression coefficients θ for (transformed)
 27 time and, possibly, interaction terms between time and screening strategy. The functions $f(\cdot)$ will be
 28 determined by visual inspection so that the unadjusted parametric CIF estimated via pooled logistic
 29 modelling approximates the non-parametric Aalen-Johansen curves reasonably well. The binary
 30 variable Y_t denotes the outcome event breast cancer death at time t . The binary variable C_t denotes
 31 censoring status at time t and the binary variable D_t contains the event status of the competing event
 32 (death by other causes) at time t . Baseline covariates and interactions between covariates are denoted
 33 by X . The prime notation $(\cdot)'$ denotes vectors. The history of a variable is denoted by overbars as $\bar{(\cdot)}$.
 34 The above model is a marginal structural model and contains baseline covariates, but no time-varying
 35 covariates. Adjustment for time-varying confounding by X_t is achieved by inverse probability
 36 weighting, where time-varying weights are calculated for each screening strategy
 37 $A \in \{A_{never}, A_{once}, A_{regular}\}$ separately as

$$38 \quad W_t^A = \prod_{k=1}^t \frac{1}{\hat{\mathbb{P}}(A_k | \bar{A}_{k-1}, \bar{X}_k, \bar{Y}_{k-1} = \bar{C}_{k-1} = 0)},$$

39 truncating weights at the 99th percentile. Here A_k is the actual screening status at time k and is, by
 40 definition, consistent with the strategy A as individuals will otherwise be censored. For efficiency the
 41 above weights can be replaced by stabilized weights (see Cain et al. (2010) for a description of
 42 stabilized weights). Analogous weights are used for censoring due to competing events when
 43 estimating the direct effect. Below, upper indices refer to counterfactuals, e.g. the probability of breast
 44 cancer death under screening even if a portion of the study subjects did not experience screening, i.e.
 45 exposure is set to a value possibly contrary to the observed exposure (Hernan & Robins, 2020). The
 46 cumulative incidence function $\widehat{CIF}_{i,t}^{A=a}$ for clone $i = 1, \dots, m$, at time point t under screening strategy
 47 $A = a$ will then be estimated using one of the approaches (i.e. based on modelling either
 48 subdistribution or cause-specific hazard) described in Young et al. (2020), depending on computational

49 cost. This cumulative incidence will be standardized to the empirical distribution of baseline
50 confounders as

51
$$\widehat{CIF}_t^{A=a} = \frac{1}{m} \sum_{i=1}^m \widehat{CIF}_{i,t}^{A=a}.$$

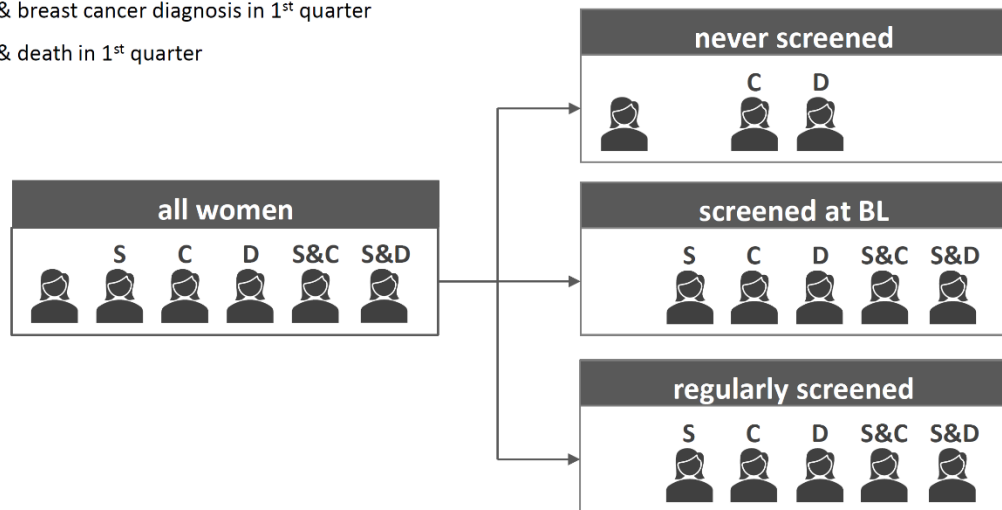
52 As a function of time t , the above cumulative incidence function allows an assessment of how the
53 effect of screening evolves over the whole of follow-up.

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55 2 Illustration of assignment to screening strategies

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- S: screening in 1st quarter
- C: breast cancer diagnosis in 1st quarter
- D: death in 1st quarter
- S&C: screening & breast cancer diagnosis in 1st quarter
- S&D: screening & death in 1st quarter



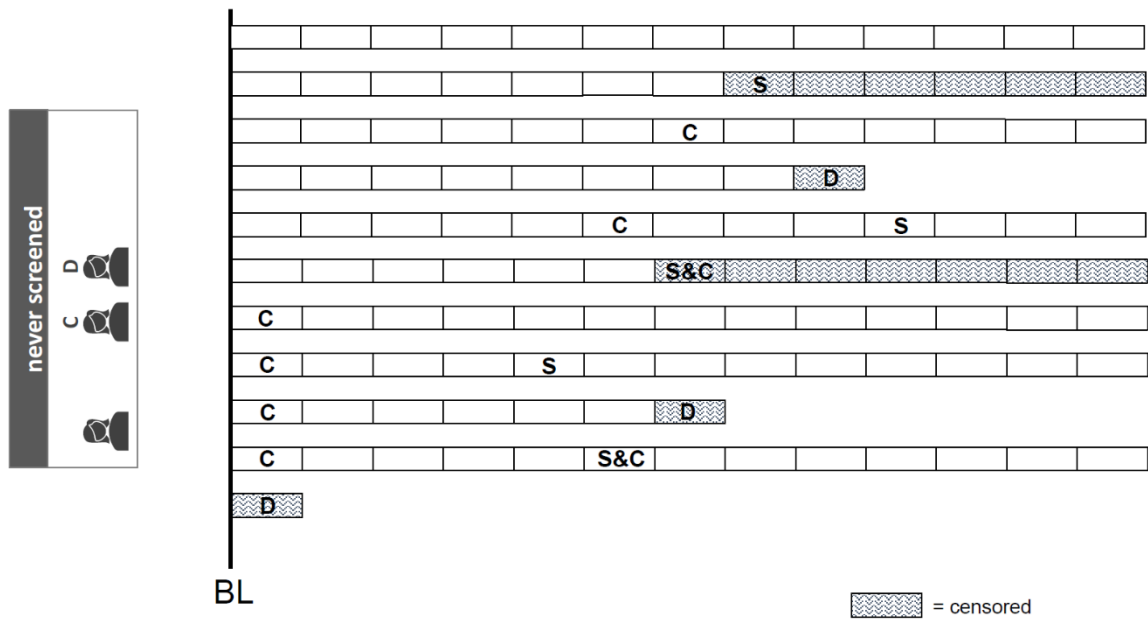
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Figure S1: Illustration of cloning of women into the screening strategies. Assignment of clones to screening strategies is based on screening behaviour from the calendar quarter of baseline. Women with a breast cancer diagnosis or recorded death in the first quarter are cloned into all screening strategies, since they were compliant with all screening strategies until the diagnosis/death occurred.

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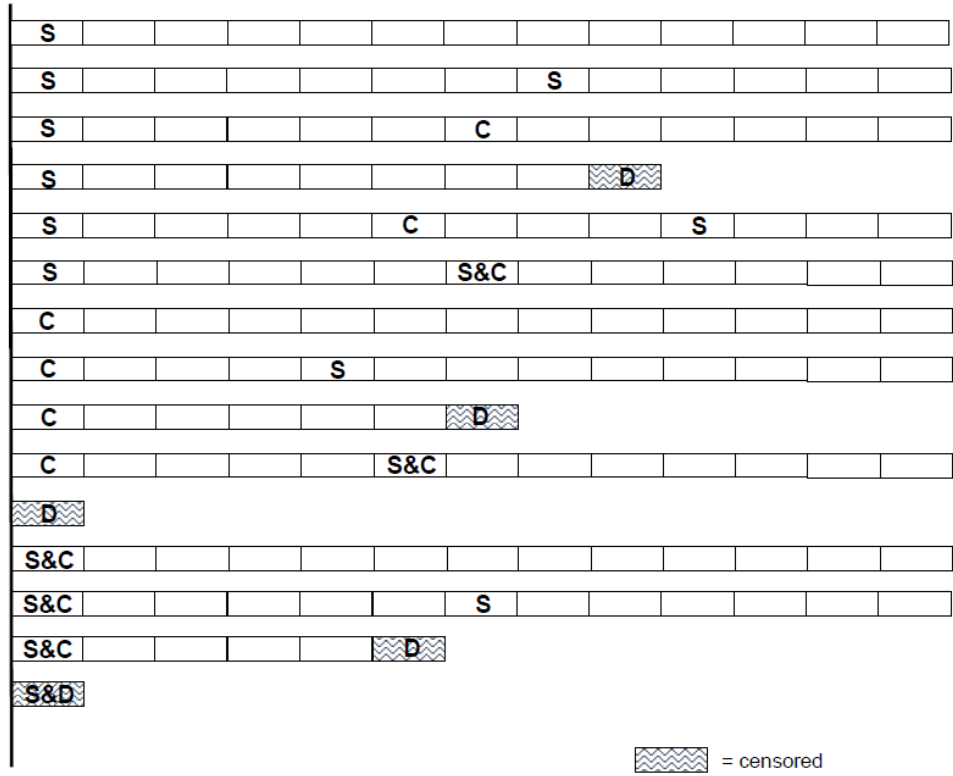
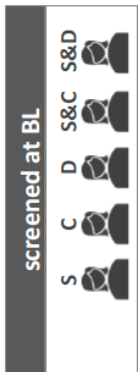
59 3 Illustrations of artificial censoring schemes per screening strategy

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Figure S2: Illustration of artificial censoring scheme under screening strategy “never screened”. Follow-up time is discretized into calendar quarters, with rectangles denoting individual quarters. The rationale for censoring is described in depth in the main body of the paper. Note that when a woman is censored, the time of censoring is set to the beginning of the calendar quarter that led to censoring. In the above illustration, the last woman is censored at baseline because she dies in the baseline quarter, i.e. she is censored at time point 0 with reason of censoring being death. C = breast cancer, D = death, S = screening, S&C = screening and cancer in the same quarter.



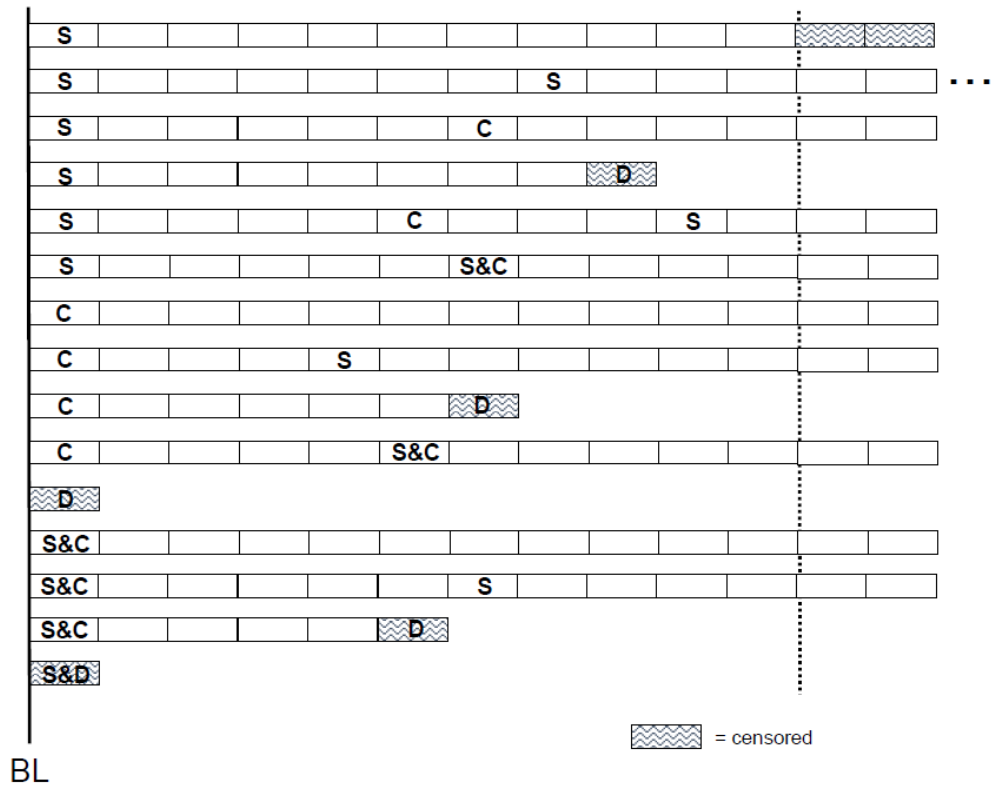
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Figure S3: Illustration of artificial censoring scheme under screening strategy “screened at baseline”. Follow-up time is discretized into calendar quarters, with rectangles denoting individual quarters. The rationale for censoring is described in depth in the main body of the paper. C = breast cancer, D = death, S = screening, S&C = screening and cancer in the same quarter, S&D = screening and death in the same quarter.

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Figure S4: Illustration of artificial censoring scheme under screening strategy “regularly screened every two years”. Follow-up time is discretized into calendar quarters, with rectangles denoting individual quarters. A regular screening is defined as having taken place between one year to ten quarters after the previous screening. The rationale for censoring is described in depth in the main body of the paper. C = breast cancer, D = death, S = screening, S&C = screening and cancer in the same quarter, S&D = screening and death in the same quarter. The dotted line indicates the end of the time period in which the second screening would need to take place.

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68 4 Addressing potential sources of bias

69 *Confounding:* Covariates used to adjust for confounding will be derived at baseline and during follow-
70 up. Their selection is based on subject matter knowledge and available literature. Risk factors for
71 breast cancer were considered relevant, even though the outcome variable is breast cancer mortality,
72 since developing breast cancer is a necessary antecedent for breast cancer death. Figure S5 illustrates
73 the causal considerations for covariate selection. Adjustment for confounding will be carried out via
74 standardization and inverse probability weighting.

75 Given that claims data are not collected for research purposes, direct information on relevant
76 confounders is not always available or only available for extreme cases (e.g. heavy smoking, alcohol
77 abuse). We aim to minimize this problem by using indirect information on these confounders (e.g.
78 diseases resulting from exposure to these risk factors such as smoking-related diseases, or diseases
79 resulting mainly from unhealthy behaviour such as obesity) as well as proxy variables for a health-
80 seeking behaviour (e.g. utilization of preventive services, educational attainment). With respect to
81 family history of breast cancer, the information is restricted to the ICD-10-GM code Z80.3 (“malignant
82 neoplasm of the breast in the family”). It is not clear whether it is primarily coded in patients with a
83 hereditary breast cancer syndrome rather than in those with a “simple” family history. The observed
84 low proportion of women with Z80 codes (Braitmaier et al. 2022) indicates that it might only be used
85 in high-risk subjects who would not be the target group of normal MSP screening. We therefore plan
86 to conduct sensitivity analyses excluding women with this code. In addition, we will conduct a
87 quantitative bias analysis to estimate the impact of unmeasured confounding regarding a “simple”
88 family history of breast cancer.

89 For some risk factors, no information will be available in our data, for example age at menarche, parity,
90 age at first full-term pregnancy, breastfeeding, age at menopause, height, breast density, exposure to
91 radiation (unrelated to mammography). However, we argue that these risk factors are relatively
92 unknown to the public and it is therefore reasonable to assume that they do not influence the decision
93 to undergo screening.

94 *Healthy screenee bias:* Individuals volunteering for screening are generally healthier than individuals
95 who choose not to undergo screening (Weiss & Rossing 1996). In addition to adjustment for
96 confounding, we will address this specific issue by carrying out a subgroup analysis within screening-
97 affine women, defined by their pre-baseline use of other preventive services (research question 2).
98 This subpopulation is more homogenous regarding health seeking behaviour, and we expect an
99 increased internal validity albeit at the cost of generalizability. Therefore, both effects, the one in the

100 full study sample and the one in the subgroup of screening-affine women, will be important for the
101 evaluation of the screening programme.

102 *Competing events:* Death due to causes other than breast cancer is a competing event for the outcome
103 of interest. We will compare the total effect (where death due to other causes is not treated as
104 eliminable) with the direct effect of screening (where the competing event is treated as eliminable and
105 thus censored with appropriate inverse probability of censoring weights, IPCW). Note that adjustment
106 for confounding of the direct effect must also include common causes of the competing event and the
107 study outcome, e.g. by including comorbidities (Young et al. 2020).

108 *Time-related biases:* Immortal time and other biases will be minimized by aligning eligibility checks and
109 treatment assignment at time zero, i.e. baseline (Dickerman et al. 2019). Furthermore, women whose
110 screening behaviour in the first quarter after trial start is consistent with more than one screening
111 strategy will be copied and one clone will be assigned to each eligible screening strategy, i.e. women
112 who undergo screening in the baseline quarter will be assigned to all active screening strategies. An
113 alternative, but less efficient approach would be to randomly assign each person to exactly one of the
114 eligible strategies (Garcia-Albeniz et al. 2020). Given that some information in the database used for
115 this study is only available on a quarterly basis (e.g. outpatient diagnosis codes), it is impossible to
116 break down the information into smaller time intervals than quarters. However, the length of follow-
117 up required to observe the effect of screening is large (approx. 7 - 10 years) (Jatoi & Miller 2003). We
118 therefore argue that the extent of bias due to the time units is negligible, as a delay of diagnosis of
119 three months is unlikely to influence the screening effect.

120 *Misclassification:* Health claims data is primarily generated for reimbursement purposes and,
121 therefore, some diagnosis codes might be used inappropriately for the underlying condition or over-
122 used (e.g. diagnosis codes in the outpatient setting). To minimize misclassification, we define most of
123 the diseases based on algorithms that, for example, combine different sources of information (e.g.
124 diagnosis codes in combination with therapy), only use codes with a high validity (such as inpatient
125 diagnosis codes) or only consider codes if recorded repeatedly. There may still be some
126 misclassification of morbidity, but we consider this type of misclassification unlikely to differ between
127 groups and negligible in our analysis. Risk factors that have a delayed impact on breast cancer may not
128 be measured adequately due to a limited length of the available look-back period. For instance, HRT
129 might influence breast cancer risk only after several years. Thus, a woman who stopped HRT treatment
130 five years before baseline would be misclassified as “no HRT use” if her look-back period in the data is
131 only three years. We will systematically describe the available look-back period (stratified by age at
132 baseline) to assess whether this could be a relevant misclassification. Finally, misclassification of the
133 outcome variable of breast cancer related deaths might occur since this variable is not directly

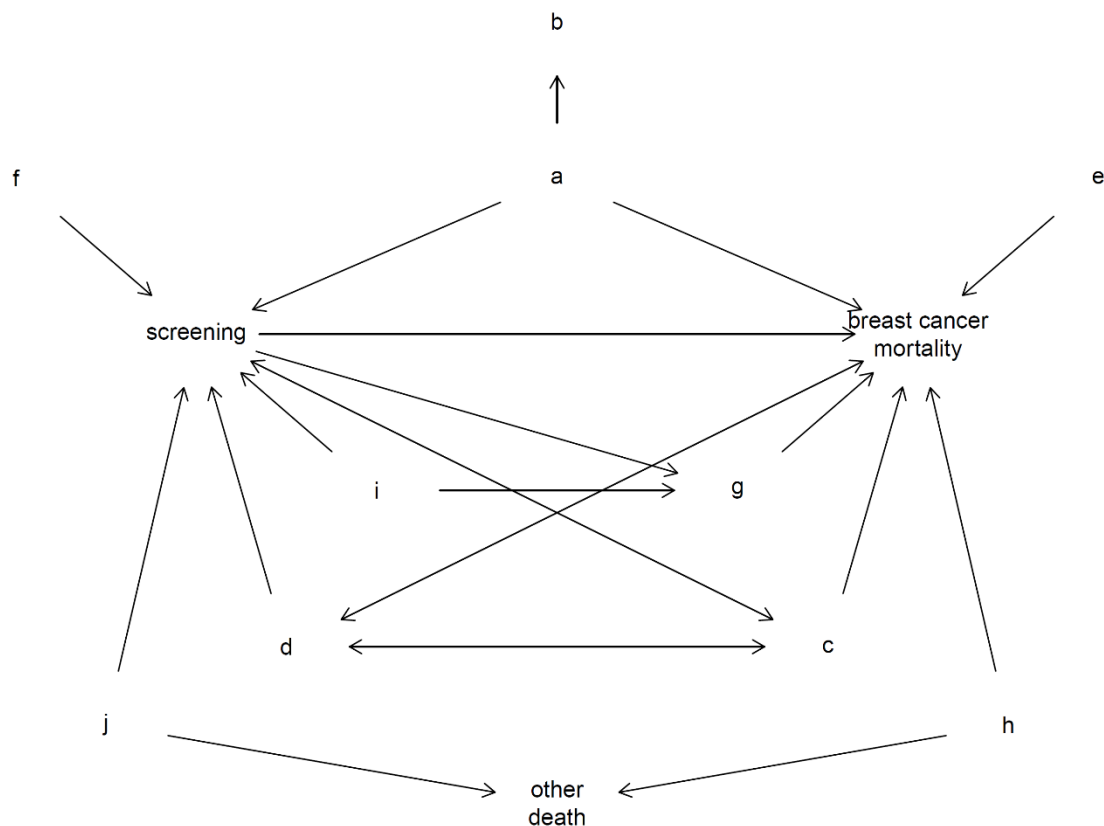
134 available for much of the data and must be derived based on an algorithm. Langner et al. (2019)
135 reported a sensitivity of 91.3 % and a specificity of 97.4 % for a former version of this algorithm, which
136 is currently being further optimized and will be validated again based on a sample for which the official
137 cause of death is available.

138 *Identifying assumptions:* We make the usual assumptions for causal inference from observational data,
139 namely consistency, sequential exchangeability given observed covariates, and positivity. Consistency
140 is fulfilled when the screening strategies being assessed are well-defined and correspond to the
141 screening behaviour observed in the data, e.g. the outcome for a woman who happens to never
142 undergo screening is the same as if she had been assigned to never undergo screening in the target
143 trial. Sequential exchangeability is fulfilled when the observed screening behaviour of a woman at time
144 t is independent of her potential outcomes under the strategies given the measured covariates prior
145 to t ; this can be thought of as no unmeasured baseline or time-varying confounding. Positivity is
146 fulfilled when the probability of observing a screening strategy is greater than zero for all strategies in
147 all covariate strata (Young et al. 2020, Hernan & Robins 2020). Furthermore, censoring competing
148 events to obtain the direct effect requires an assumption of no unmeasured common causes of the
149 different event types.

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151 5 Illustration of causal considerations for covariate selection

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Figure S5: Illustration of variable groups considered for covariate adjustment and their causal connections. Note that this is a simplified graph, ignoring the longitudinal aspect of the study. A directed edge from one variable to another means that the first variable is a direct cause of the second. Screening is the exposure, breast cancer death is the outcome, and other death is a competing event. A bi-directed edge can be interpreted as presence of latent variables between the two connected variables. Variables “a” are common causes of screening and outcome. Variables “b” are proxies for those of category “a”. Variables “c” are causes of the outcome that are associated with exposure. Variables “d” are causes of the exposure that are associated with the outcome. Variables “e” are causes of the outcome that are not associated with exposure. Variables “f” are causes of the exposure that are not associated with the outcome. Variables “g” are post-screening variables that are mediators between exposure and outcome. Variables “h” have a causative effect both on the competing event and the outcome. Variables “i” are causes of exposure and mediators. Variables “j” are confounders between exposure and the competing event. Variables “f” should not be included for adjustment, as this can lead to bias-amplification in case of residual unobserved confounding. Variables “g” (e.g. treatment after screening) should not be included for adjustment, as they are on the causal path from exposure to outcome. Variables “a”, “b” (if “a” is unmeasured), “c”, “d”, “h” (only for estimating the direct effect, not for the total effect), “i”, and “j” should be included for adjustment to mitigate confounding. Variables “e” are not needed for adjustment but can be included to increase precision of estimation. The variable groups (except “f”) are not mutually exclusive, and in fact many variables will fit into more than one of these groups. An example of a covariate of the category “a” would be previous use of menopausal hormone therapy, as this is a known risk factor for breast cancer and physicians might advise women with this risk factor to attend screening. An example of a covariate of the category “j” would be presence of palliative care. An example for “d” might be educational attainment as it may affect awareness of screening and is strongly associated with direct risk factors “c” of breast cancer mortality; educational attainment can also be seen as type “b” proxy for further unmeasured confounders.

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156 **6 List of covariates**

157 In Table S1 below, we give an overview of variables used to adjust for confounding. Time-varying
 158 covariates will be re-assessed on a quarterly basis. Variables might be added to this list of covariates,
 159 if indicated by subject matter knowledge. The list of covariates used in the final analysis will be finalized
 160 before data on the study outcome becomes available. Note that this is just an alphabetical list of
 161 covariates that will be defined based on the information in the database. Content-wise, a discussion
 162 on how confounding as a potential source of bias is considered and how relevant covariates are
 163 captured in the data is provided in Supplement 4. Furthermore, Figure S5 illustrates the causal
 164 considerations for covariate selection.

165 The covariates in Table S1 are mostly implemented as binary (time-dependent) variables. For most of
 166 the variables, algorithms considering different types of information (e.g. diagnosis codes in
 167 combination with therapy) will be developed or have been developed, with the aim of maximizing
 168 validity and thus minimizing misclassification (see also Supplement 4).

169 **Table S1:** Relevant covariates for confounder adjustment.

variable/variable group	time-varying
Acute hemorrhagic stroke	yes
Acute ischemic stroke	yes
Acute myocardial infarction	yes
Age at baseline	no
Alcohol abuse	yes
Anaemia	yes
Anticoagulant therapy	yes
Antihypertensive therapy	yes
Antiplatelet therapy	yes
Benign neoplasm of breast	yes
Breast disorders (benign mammary dysplasia, inflammatory disorders of breast, hypertrophy of breast, unspecified lump in breast, other disorders)	yes
Bronchial asthma	yes
Cachexia	yes
Chronic obstructive pulmonary disease (COPD)	yes
Coronary heart disease	yes
Dementia	yes
Diabetes with end organ damage	yes
Drug abuse	yes
Drug-treated (arterial) hypertension	yes
Educational attainment	no
Family history of breast cancer*	yes
Glaucoma	yes
Heart failure	yes
Hemiplegia	yes
Hepatitis B or C	yes
Hip fracture	yes
HIV therapy	yes

Hormone replacement therapy	yes
Lipid-lowering therapy	yes
Liver diseases including chronic viral hepatitis	yes
Mental diseases	yes
Number of hospitalizations	yes
Number of non-screening mammographies	yes
Number of outpatient physician contacts	yes
Number of prescriptions	yes
Number of screening mammographies	yes
Obesity/adiposity	yes
Other cancers	yes
Palliative care	yes
Severe liver disease	yes
Terminal renal disease	yes
Tobacco abuse	yes
Treated diabetes	yes
Treatment for hypothyroidism	yes
Treatment for osteoporosis	yes
Treatment with antidepressants	yes
Treatment with antipsychotics	yes
Treatment with immunosuppressive drugs	yes
Treatment with opioids	yes

170 * Given that information on family history is limited, additional methods will be taken to consider
171 this confounder (see manuscript and Supplement 4).

172

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