

## ORIGINAL ARTICLE



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# Assessing heterogeneity of electronic health-care databases: A case study of background incidence rates of venous thromboembolism

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## Abstract

**Purpose:** Heterogeneous results from multi-database studies have been observed, for example, in the context of generating background incidence rates (IRs) for adverse events of special interest for SARS-CoV-2 vaccines. In this study, we aimed to explore different between-database sources of heterogeneity influencing the estimated background IR of venous thromboembolism (VTE).

**Methods:** Through forest plots and random-effects models, we performed a qualitative and quantitative assessment of heterogeneity of VTE background IR derived from 11 databases from 6 European countries, using age and gender stratified background IR for the years 2017–2019 estimated in two studies. Sensitivity analyses were performed to assess the impact of selection criteria on the variability of the reported IR.

**Results:** A total of 54 257 284 subjects were included in this study. Age-gender pooled VTE IR varied from 5 to 421/100 000 person-years and IR increased with increasing age for both genders. Wide confidence intervals (CIs) demonstrated considerable within-data-source heterogeneity. Selecting databases with similar characteristics had only a minor impact on the variability as shown in forest plots and the magnitude of the  $l^2$  statistic, which remained large. Solely including databases with primary care and hospital data resulted in a noticeable decrease in heterogeneity.

**Conclusions:** Large variability in IR between data sources and within age group and gender strata warrants the need for stratification and limits the feasibility of a meaningful pooled estimate. A more detailed knowledge of the data characteristics, operationalisation of case definitions and cohort population might support an informed choice of the adequate databases to calculate reliable estimates.

## **Key Points**

1. Using a multi-database approach provides a more accurate picture of true IRs, as there may be large clinical differences underpinning the variability in the estimates across different databases.

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- After mitigating unwanted heterogeneity through harmonization of database characteristics, there might still be some heterogeneity present, but this should be considered as a source of knowledge; our study confirmed prior knowledge that VTE backgrounds IRs were different dependents on the age and gender of the individual.
- 3. The level of the heterogeneity in estimates depends on differences in database characteristics. In our study, databases collecting data from different parts of the health-care systems were the largest contributors to heterogeneity in estimates.
- 4. When heterogeneity is present, a careful trade-off has to be made for the choice of IR, between stratified estimates or a pooled estimate, to support use in pharmacoepidemiological and regulatory evaluation.
- 5. To attenuate heterogeneity, a pre-screening of database characteristics through a metadataset and adequate analytical tools at study design stage might be considered.

#### **Plain Language Summary**

Real-world data collected in everyday clinical practice can complement information used in regulatory decision-making and provide evidence to support the benefit-risk assessment of medicines. To improve the added value of real-world data for regulatory decision-making, regulators pool information from multiple databases to provide a more accurate picture of the outcome of interest. However, there is regularly variability, also called heterogeneity, in study outcomes when using data from different databases and this poses challenges for interpretation and communication. In this study, we examined incidence rates of venous thromboembolism, identified as a potential side effect for some COVID-19 vaccines, derived from multiple databases. We investigated how differences in database characteristics might cause variation between rate estimates and concluded that the largest contributor to heterogeneity was the use of data from different health-care settings. Understanding which database characteristics contribute to variability can allow to mitigate variation. This can be done by selecting databases with similar data characteristics, such as harmonised codes to refer to clinical outcomes and comparable selection criteria for participants and by using appropriate statistical methods to analyse the variability. Overall, our study provides an overview of the complexity of real-world evidence and can be used to better understand and analyse sources of variability.

## 1 | INTRODUCTION

In the past two decades, the usage of large health-care databases has increased greatly.<sup>1</sup> Regulatory agencies such as the European Medicines Agency (EMA) and the US Food and Drug Administration (FDA) have highlighted the value of real-world data (RWD) in medicines regulation.<sup>2,3</sup> In Europe, the initiation of the Data Analysis and Real World Interrogation Network (DARWIN EU),<sup>4</sup> as well as the European Health Data Space are changing the landscape of real-world evidence (RWE) generation towards multi-database studies.

While there are already a number of advantages in using RWD in regulatory decision-making, those benefits can be improved by using more than one data source.<sup>5</sup> Trivially, incorporating data from multiple data sources in an analysis will increase sample size. This can be crucial in situations with low event counts, such as for estimating the incidence rates (IRs) of a rare disease. While observational data are more generalizable to the real world than randomised controlled trials, the level of generalizability can be even further increased, by covering a broader and more representative population, thus possibly mitigating selection biases that are specific to single databases and by allowing for the quantification of true differences between populations.

Even though the benefits of using multiple data sources cannot be denied, data from those sources should not be pooled without a preliminary assessment of the suitability of pooling data, due to inherent differences in their characteristics. Simulations have shown increased risks of false-positive and false-negative safety signals when pooling data from multiple databases.<sup>6</sup> This heterogeneity can have multiple forms, some of which are desirable for understanding true differences in outcomes event rates, while others make interpretation of results and decision-making regarding selection of suitable background rates for specific purposes such as observed-to-expected analyses for vaccines highly challenging.<sup>7</sup>

Sources of heterogeneity can be categorized into three types: measurement heterogeneity, information heterogeneity (both may be considered methodological heterogeneity), and true heterogeneity (also called clinical heterogeneity).<sup>8</sup> While measurement (e.g., clinical classification systems) and information heterogeneity (e.g., granularity of clinical codes) can generally be considered undesirable, clinical heterogeneity has its value, for example, by improving external validity of results or understanding differences in prescription patterns or impact of risk minimisation measures (RMMs) between different geographical regions, health-care systems or behaviours. However, to understand

heterogeneity, it is important to use appropriate tools to detect, report and account for it.

During the COVID-19 pandemic, RWD rapidly provided impactful evidence on safety and effectiveness of therapeutics and vaccines.<sup>9</sup> This included the generation of background IR for adverse events of special interest (AESIs) for COVID-19 vaccines.<sup>10</sup> Those background rates continue to be used in observed-to-expected analyses to estimate the expected number of cases in the general population prior introduction of COVID-19 vaccination, or during SARS-CoV2 circulation in non-vaccinated populations.

The list of AESIs included the concepts of deep vein thrombosis (DVT) and pulmonary embolism (PE). These two concepts make up the term venous thromboembolism (VTE).<sup>11</sup> EMA pharmacovigilance activities identified VTE as a possible adverse event of Jcovden (former COVID-19 Vaccine Janssen)<sup>12</sup> and listed VTE as an adverse event of Vaxzevria.<sup>13</sup> Background rates were used to calculate the excess number of cases potentially linked with these vaccines. However, the reported background rates showed large differences between EU countries as reflected through national health-care records.<sup>14,15</sup>

The objective of this study was to explore data characteristics that trigger heterogeneity in IR through both descriptive and statistical measures, using VTE as a case study. This investigation will further provide support in selecting adequate statistical methods for handling heterogeneity when pooling observations to derive meaningful pooled estimates to support regulatory decision-making.

## 2 | METHODS

## 2.1 | Data

To demonstrate an analytical workflow for handling heterogeneity between databases, we selected VTE, a safety concern listed for a class of COVID-19 vaccines, as a case study.

In order to assess potential adverse reactions related to approve COVID-19 vaccines in the EU, EMA-funded two studies through large research consortia: the ACCESS project with University Medical Center Utrecht and the EU PE&PV Research Network<sup>15,16</sup> and a study by ERAS-MUS University Medical Center<sup>17</sup> to generate aggregated background IRs of AESIs, including VTE. Both consortia reported background IRs from multiple databases stratified by age group and gender, using the same eight age categories. IRs were estimated by dividing the number of incident cases by the total person-time at risk, with individuals entering the study cohort on their first visit after January 1, 2017, and being followed until the outcome, exit from the database or end of the study period. The study period covered 2017-2020. In the ACCESS protocol, the study population included all individuals who were observed in the databases for at least 1 day during the study period (January 1, 2017 to last date available) and who had at least 1 year of data availability before cohort entry, except for individuals <1 year of age with data available since birth. In the ERASMUS protocol, the study population was defined slightly different people observed on January 1 2017, January 1, 2018, or January 1, 2019 had to be observed continuously for at least 365 days with no event before this observation date. Ninety-five percent of CIs were calculated using an exact method described by Ulm.  $^{18}\,$ 

Case definitions for VTE were developed by the researchers independently. ACCESS utilized the CodeMapper tool<sup>19</sup> to find harmonized definitions across coding systems. The full list of included concepts and details on its generation process is publicly available.<sup>20</sup> Through using the OMOP common data model for its analyses, clinical codes in databases to which ERASMUS has access to were mapped to the SNOMED system, ensuring harmonized case definitions. The list of clinical codes included by ERASMUS can be accessed in the ATLAS application.<sup>21</sup> Table A1 in the appendix shows included ICD10 codes for both ACCESS and ERASMUS. Only ICD10 codes are shown since both research organisations harmonized their definitions across coding systems. For both PE and DVT, the definition by ACCESS includes a broader range of concepts. For PE, the additional concepts are related to septic PE. The additional concepts for DVT mostly correspond to phlebitis, thrombophlebitis and DVT related to pregnancy. As described in a report by the FDA,<sup>22</sup> there are no clear guidance on whether these concepts should be included or not.

A short overview of the databases is provided in Table 1. Further details, including demographic characteristics and total population, are provided in the corresponding published reports.<sup>15,16</sup> All databases are listed in the ENCePP research database, which also shows a list of relevant research publications they have been used in. For three databases (PHARMO, BIFAP, and SIDIAP), the IR had been estimated both on the total population and on the subset of subjects with linked primary care and hospital records (PC-H linkage). For the primary analysis, the total population estimates were used.

The data included the years 2017–2019. ACCESS reported IRs by year; hence, rates were pooled based on counts to match the data structure of ERASMUS, who reported only combined estimates for all years. Data from the Danish registries (DCE-AU) were reported only for the years 2010–2013 and thus were not included in the analysis. For the PHARMO database, only data for 2017 and 2019 was reported, due to an error in the imputation of a subset of data for 2018; BIFAP data with hospital linkage were only reported for 2017–2018.

## 2.2 | Analysis

We used forest plots to visualize heterogeneity, displaying estimated IRs as squares with CIs for each database. The size of the squares is proportional to the precision of the estimate.

A random-effects meta-analysis, using the restricted maximum likelihood (REML) estimation method, was performed on the log scale of the IRs to calculate a summary estimate and to quantify the level of heterogeneity, thereby allowing for heterogeneity between databases, which is more realistic than assuming that the true value of the estimand is exactly the same for each database.

To quantify the absolute value of this heterogeneity, we reported estimates of  $\tau^2$  measuring the 'dispersion of true effect sizes between studies in terms of the scale of the effect size'<sup>23</sup> and  $l^2$  measuring

## TABLE 1 Overview of main characteristics by data source.

	Consortium	Region	Type of data source/health care system: Primary care (PC) or Hospital (H) data	Study population	Clinical Classification Coding system
PHARMO	ACCESS	Netherlands	Н	9 184 832	ICD10
PHARMO	ACCESS	Netherlands	PC and H	496 197	ICPC (PC), ICD10 (H)
BIFAP	ACCESS	Spain	PC	10 266 468	SNOMED, ICD9
BIFAP	ACCESS	Spain	PC and H	4 423 843	SNOMED, ICD9 (PC), ICD10 (H)
SIDIAP	ACCESS	Catalonia (Spain)	PC	6 205 573	ICD10
SIDIAP	ACCESS	Catalonia (Spain)	PC and H	1 758 239	ICD10
FISABIO	ACCESS	Valencia region (Spain)	PC and H	5 886 560	ICD9, ICD10
PEDIANET	ACCESS	Italy	PC	181 290	ICD9
ARS	ACCESS	Tuscany (Italy)	PC and H	3 067 602	ICD9
CPRD GOLD	ACCESS	United Kingdom	PC	4 688 710	READ, SNOMED
CPRD GOLD	ERASMUS	United Kingdom	PC	3 913 071	READ
IQVIA DA Germany	ERASMUS	Germany	PC	8 459 098	ICD10
IQVIA LPD France	ERASMUS	France	PC	3 951 633	ICD10
IPCI	ERASMUS	Netherlands	PC	1 299 288	ICPC
IQVIA LPD Italy	ERASMUS	Italy	PC	1 066 230	ICD9
SIDIAP	ERASMUS	Catalonia (Spain)	PC and H	1 909 814	ICD10

Abbreviations: ARS, Agenzia Regionale di Sanita Toscana; BIFAP, Base de Datos para la Investigacion Farmacoepidemiologica en Atencian Primaria; CPRD, clinical practice research datalink; DA, disease analyzer; FISABIO, Fundación para el Fomento de la Investigación Sanitaria y Biomédica de la Comunitat Valenciana; IPCI, integrated primary care function; LPD, longitudinal patient data; SIDIAP, Sistema d'Informació per al Desenvolupament de la Investigació en Atenció Primària.

what proportion of variation in the observed effects is due to variation in true effects, that is, due to inherent differences between the investigated data sources rather than sampling error.<sup>24</sup> Borenstein et al.<sup>18</sup> stress that l<sup>2</sup> represents a proportion rather than an absolute value. Therefore, we estimated the level of heterogeneity in comparison with statistical variability rather than heterogeneity itself. In addition, a prediction interval was calculated.<sup>25</sup> Such an interval combines uncertainty due to sampling variation and due to heterogeneity to provide an approximate range of true values.

Finally, based on the available metadata characteristics of each of the included databases (see Table 1), several supplementary analyses were performed omitting or selecting a set of databases meeting selected criteria, aiming to reduce potential unwanted measurement and information heterogeneity, in order to assess more accurately true differences in subject-level data (i.e., true heterogeneity). These exploratory analyses allowed determining the contribution of each database characteristic to the heterogeneity in IR across databases.

The following supplementary analyses were performed:

- a. Restricting the analysis to a subpopulation; only those databases with linkage between primary care and hospital data can provide information on the influence of the health-care setting (i.e., type of data source). Including data sources with only primary care data might lead to underestimation in case of in-patient diagnosis. Data sources with only hospital data may underestimate the events in case of out-patient diagnosis.
- b. Restricting the analysis to only those databases using the same clinical classification system for diseases. In this study we included

only databases that used the International Clinical Classification of Disease (ICD10)<sup>26</sup> to diagnose VTE as it is the most widely used vocabulary among the available data sources.

- c. Restricting the analysis to databases with homogeneous case definition. Table A1 in the appendix specifies the ICD10 codes used to diagnose VTE in the two studies. We performed separate analyses by study, that is, ACCESS and ERASMUS, to explore differences in case definitions and population selection criteria.
- d. The analyses were performed using the R software  $^{\rm 27}$  package meta.  $^{\rm 28}$

## 3 | RESULTS

A total of 60 080 169 subjects contributed to the 13 databases. See Tables A2 and A3 for an overview of the reported IRs stratified by age category and gender, by database and by study.

Two databases (CPRD GOLD and SIDIAP) were used in both studies by both consortia. Differences in defining the study cohort resulted in the cohort entry criteria not being identical. After removing the duplicated databases, a total of 54 257 284 subjects derived from 11 databases were included in the main analysis, representing collectively all age and gender subgroups from six countries.

As the first step, we display the age-gender-database-specific IR estimates in a forest plot (Figure 1) using total population estimates. The forest plot showed a relatively large amount of heterogeneity between databases and within strata of age groups and gender. While for the 0–19 age group the IRs appeared to be in the same order of



magnitude, with increasing age and increasing IRs the heterogeneity increased. In the 80+ age group, estimates ranged from <100 to >1 000 per 100 000 person-years. There did not seem to be a large difference in heterogeneity between gender categories. Two databases, LPD France and PHARMO, showed considerably lower estimates than the other databases in most age groups. The Dutch IPCI database, on the other hand, showed the highest estimates, especially for younger age groups and women. We generally observed consistent ranking of IR estimates across age groups, that is, across strata estimates are consistently high or low relative to the other databases.

Next to the age-gender-database-specific IRs, age-gender IRs from meta-analyses were calculated. In our study, the meta-analysis estimated IR of VTE from 5 to 421 per 100 000 person-years depending on age-gender strata. The wide confidence interval of the summary (i.e., pooled) measure identified even within each stratum large patient-level differences. Table A4 in the appendix displays the age-gender IR estimates and CIs of the pooled measure for VTE from meta-analyses.

Figure A1 in the appendix shows the calculated prediction interval for the primary analysis. The prediction intervals for each agegender group were notably high confirming the substantial population-level heterogeneity observed across data sources.

Table 2 shows the estimated  $l^2$  and  $\tau^2$  values by age-gender stratum. The values for  $l^2$  indicated that a majority of the observed variability is due to differences between databases rather than random sampling error. Supporting the impression from the forest plot, we observed an increasing estimate of  $l^2$  with increasing age in both sexes. There did not seem to be an age-related trend in the estimates of  $\tau^2$ , but estimates for  $\tau^2$  appear to be lower for males than for females.

#### 3.1 | Sensitivity analyses

In the first sensitivity analysis, we restricted the databases to those with PC-H linkage (Figure 2). The forest plot demonstrates a relatively large decrease in heterogeneity when restricting the analysis to databases with PC-H linkage, across all age groups. It became apparent that this restriction of databases primarily leads to low estimates being excluded from the analysis. In Table A5 in the appendix, which lists  $l^2$  and  $\tau^2$  estimates for all sensitivity analyses, we noticed lowered  $l^2$  estimates specially for younger age groups and considerably lowered  $\tau^2$  values for all age groups. Figure 3 did not imply any reduction in heterogeneity when restricting the analysis to databases using ICD 10 codes to diagnose VTE. Both range and distribution of estimates were similar to the primary analysis. The same was true for estimates of  $l^2$  and  $\tau^2$ . Figure 4a, b showed forest plots of the analysis considering data from ACCESS and ERASMUS separately. Note that due to

**TABLE 2** Age-gender-stratified  $I^2$  and  $\tau^2$  estimates from metaanalyses.

Age	Gender	l <sup>2</sup>	$\tau^2$
0-19	Female	0.872	0.253
0-19	Male	0.921	0.376
20-29	Female	0.983	0.531
20-29	Male	0.954	0.381
30-39	Female	0.992	0.626
30-39	Male	0.981	0.423
40-49	Female	0.996	0.600
40-49	Male	0.991	0.367
50-59	Female	0.997	0.517
50-59	Male	0.995	0.299
60-69	Female	0.998	0.473
60-69	Male	0.996	0.309
70-79	Female	0.998	0.527
70-79	Male	0.997	0.358
80+	Female	0.998	0.735
80+	Male	0.997	0.625

**FIGURE 2** Age-gender-databasespecific IR estimates and 95% CIs for VTE in databases with PC-H linkage. \*Due to the CIs being too small compared with the size of the square, some of the CIs are not noticeable in the figure.

including the estimates from both the total population and the subpopulation with PC-H linkage, there was some dependence between the estimates of BIFAP, SIDIAP, and PHARMO. For ACCESS, the hospital database PHARMO showed far lower estimates than all other databases included. This could be linked to an oversampling of the denominator. Apart from this, visually there seemed to be some reduction in heterogeneity. Data from ERASMUS and ACCESS showed a similar amount of heterogeneity; the forest plots indicate that estimates from ERASMUS are spread more equally, while the PHARMO hospital data differ strongly from the other estimates within ACCESS.

## 4 | DISCUSSION

This study explored heterogeneity in background IRs of VTE reported from 11 data sources spanning six EU countries and derived from two observational studies, by focusing on the database as a source of heterogeneity. Through investigating data source characteristics potentially introducing differences in estimated IRs, our aim was to investigate the amount of unwanted (i.e., methodological)



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heterogeneity or uncertainty between data sources to provide more valid conclusions for safety surveillance activities. Data sources used in this study were mostly from primary care settings, partly with linkage to hospital data. The study used aggregated background IRs of VTE, considered a relevant AESI for a class of EU-approved COVID-19 vaccines.

Substantial heterogeneity in the background IRs was observed between all included data sources, in addition to observed within-datasource differences across age groups and genders. Age was the main contributor to the heterogeneity as shown in our study. Overall, it was observed that background rates increased with increasing age with no clear pattern in IR between males and females. The observation of increased IRs with increasing age is in line with another study on VTE,<sup>29</sup> with the same study also suggesting a difference in IR between genders: IRs increase markedly with age for men and women; the overall ageadjusted IR is higher for men (130 per 100 000) than women (110 per 100 000). The observed heterogeneity in the different age-gender strata is a source of information that leads to a better understanding of the burden of VTE in the general population. However, as demonstrated through the summary estimate and CIs, we still found substantial heterogeneity between data sources within each stratum, suggesting that there still might be unobserved patient-level heterogeneity and therefore a single estimate for each stratum might be inaccurate.

In an attempt to understand the contribution of database characteristics to the reported heterogeneity, we performed several exploratory analyses. Our databases included data derived from both hospital and primary care settings. In all data sources, when estimating background rates, it is important to consider how the population denominator was derived. When linking data between the two settings, depending on the mechanism of linkage, there is a risk of only capturing those subjects that had a hospital visit recorded, which could lead to biased estimates. Restricting the databases that included a link with hospital (PC-H linkage) resulted in a moderate decrease in the reported variability. Alongside, we did not see a decrease in heterogeneity using only databases that used the ICD-10 vocabulary demonstrating that the type of vocabulary used for clinical classification of VTE could not be identified as a major source of heterogeneity. Differences in background rates remained between the two studies even if the time at risk in which the rates were collected and agegender subgroup definitions and analytical methods were similar. Comparing more closely the methodology applied in the two studies, differences in case definitions were noted, with some clinical codes only included in one of the studies (Table A1 in the appendix). The inclusion and exclusion criteria for individuals also differed, leading to non-identical study populations even when within the same data source. Since we did not find any systematic differences in estimates

**FIGURE 4** a. Age-gender-database specific IR estimates and 95% Cls for VTE in databases from ACCESS. \*Due to the Cls being too small compared with the size of the square, some of the Cls are not noticeable in the figure. b. Agegender-database-specific IR estimates and 95% Cls for VTE in databases from ERASMUS. \*Due to the Cls being too small compared with the size of the square, some of the Cls are not noticeable in the figure.



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between the two consortia, it is unlikely that differences in case definition or inclusion criteria had a large influence on observed heterogeneity.

When quantifying heterogeneity using the statistical measure  $l^2$ , a considerable amount of heterogeneity (close to 100%) is reported. The large values of  $l^2$  are not surprising, as the large sample sizes in every database imply small variance estimates. In particular the  $l^2$  estimates in the 0–19 age group seem to be influenced by this fact: due to a larger sample size, the variance is lower than for the other age groups, leading to  $l^2$  estimates that appear too high in comparison with the other age groups when looking at the forest plot.

In this study, we have calculated pooled estimates for the primary analyses. However, when large heterogeneity is present, focusing on a pooled estimate is not advisable, given that the pooled estimate will derive largely from the particular choice of databases and the relative weights associated with each database.<sup>30</sup> Following the classification in Deeks et al.<sup>8</sup> it is not advised to combine the estimates if the value of *I*<sup>2</sup> was estimated to be larger than 90%. In addition, when reporting the results after combining estimates, attention needs to be given to uncertainty quantification. In addition to the common risk of misinterpretation for Cls,<sup>31</sup> Cls for random-effects meta-analyses are easily misinterpreted to quantify dispersion of study effects. However, Cls only represent the uncertainty in estimating the mean effect size, not taking into account variability due to different database characteristics. This means, that Cls are always smaller than the range of observed estimates.

A major strength of this study is the use of data aggregated from a large number of data sources independently provided by two research consortia using the same calculation method to estimate the IRs. This enabled the exploration of database-specific aspects related to heterogeneity. From 11 databases, 54 257 284 subjects contributed to the main analysis; with the databases spanning a large part of Europe, this can be considered a representative sample of the total population.

The above exploratory analyses show that even when certain database characteristics are harmonized, significant heterogeneity is still present. A limitation of our study is that only aggregated data was available which prevented us from investigating potential sources of heterogeneity attributable to patient characteristics. For instance, comorbidities may have an influence on IRs.<sup>32</sup> A final limitation is that clinical validation of the diagnosis codes was not performed in these studies which might have led to different frequencies of misclassification or underreporting of VTE cases, which may affect estimates of heterogeneity.<sup>33</sup>

This study highlights the challenges regarding the varying levels of available information about database characteristics and the difficulty to identify sufficiently detailed information about the data sources. For example, some differences can only be explored through subject matter expertise about the corresponding health-care systems. Health-care systems might differ between regions, implying possible differences in the probability of recording certain events even in the same health-care setting. The process of clinical coding could also influence the quality of recording, with different levels of quality control or incentives for correct coding. With the large level of observed heterogeneity, an important recommendation is to use the same databases when comparing estimates at different time-points, for example, for pre- or post-exposure IRs of AESI.

The above considerations highlight the necessity for careful assessment of the suitability of databases to include in multi-database studies. In the two studies, a variety of databases was included because many different AESIs were considered simultaneously. For a study on a specific outcome, more specific restrictions on the databases should be placed a priori. In our study we have observed that the type of data source is one of the most important considerations. Based on subject matter knowledge or available validation studies, it should be evaluated in which type of setting the most accurate estimation is possible.

Our analysis also highlights the importance of unified case definitions and inclusion and exclusion cohort criteria to select adequate data sources in multi-database studies. This is evidenced in this study by the CPRD data source used by both study groups to address the same objective, but with a difference of 17% in the total number of individuals included in the study cohort, most likely related to differences in the operationalization of case definitions.

It is important to note that the methods used for detecting and addressing heterogeneity should be specified before starting any meta-analysis. When the forest plot show outliers among observed rates, it can be tempting to exclude the corresponding databases from the analysis without further investigating causes for outliers. This practice is, however, likely to introduce bias and should be avoided in most situations. Criteria for excluding certain databases should be specified prior to performing the analysis, but even then, it is advisable to also present results with the excluded databases, as a sensitivity analysis. In parallel, the choice of method for handling heterogeneity should also be prespecified, conditional of the outcome of the method for detecting heterogeneity. Also, it is preferable that the method for estimating the meta-analysis model and its statistical heterogeneity (e.g., REML), the methods for quantifying CIs (e.g., the Hartung-Knapp and Sidik-Jonkman modifications to the Wald method) and prediction intervals are prespecified.<sup>34</sup>

More specifically, exploring the level of heterogeneity using multiple databases must be considered if these rates are intended to be used to support safety signal detection activities and to avoid misleading recommendations. One of the current initiatives is the DIVERSE project with the aim to develop guidelines for the identification, collection and reporting of heterogeneity in multi-database studies.<sup>35</sup> In addition, EMA's list of metadata for Real World Data catalogues,36 which will be the basis of a catalogue of RWD sources, will provide researchers with standardized, relevant information about databases to use for RWE studies. Another approach would be to develop a set of metrics to measure database heterogeneity or to develop phenotype libraries to identify important variables in different databases. For instance, Ostropolets et al.<sup>22</sup> quantify factors influencing IRs through a set of sensitivity analysis using patient level data. Finally, the use of SNOMED CT (systematized nomenclature of medicine - clinical terms),<sup>37</sup> a terminology that can cross-map to other

classifications and code systems, may reduce the variability among estimates derived from different data sources. Nonetheless, the mapping of original coding systems to SNOMED may not reduce the heterogeneity as such, but may merely conceal possible heterogeneity introduced by different classification systems to operationalize the case definition of VTE. This is evidenced in this study by the ERAS-MUS data sources, where still large variability in the estimates is seen even when converted to SNOMED.

## 5 | CONCLUSION

Our study revealed large variability in estimated age-gender-stratified background IRs of VTE between different databases, demonstrating the presence of one or several sources of heterogeneity. Restricting the databases to similar health-care settings contributed to less variability in the reported rates. Still, variability was present, triggered most likely by presence of analytical heterogeneity through differences in the case definitions and population cohorts as defined in the protocols used by the two study groups. The use of HARPER (harmonized protocol template to enhance reproducibility)<sup>38</sup> to operationalize code definitions will improve the creation of unambiguous clinical codes in studies integrating data from multiple data sources.

Our study can be utilized to better understand the complexity of RWE and to illustrate the importance of a cautious selection of databases, based on their characteristics, so that the observed heterogeneity represents true differences, to ultimately improve the reliability of RWE. Our findings should be considered in context of similar analyses with other databases and in other settings.

## AUTHOR CONTRIBUTIONS

All authors meet ICMJE criteria, they have all contributed to the development of the study, as well as writing and approval of the manuscript.

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## APPENDIX A

**FIGURE A1** Age-gender-stratified prediction interval\* and 95% CIs for VTE by database and pooled. \* Due to the prediction interval being too small compared with the size of the square, some of the prediction intervals are not noticeable in the figure.



**TABLE A1**ICD-10 codes used by ACCESS and ERAMUS todiagnose VTE.

Coding system ACCESS <sup>a</sup>	Code	Code name
ICD10CM	126	pulmonary (acute) (artery)(vein) thromboembolism
ICD10CM	126	Pulmonary embolism
ICD10CM	180	Phlebitis and thrombophlebitis
ICD10CM	181	Portal vein thrombosis
ICD10CM	182	Other venous embolism and thrombosis
ICD10CM	O08.2	Embolism following ectopic and molar pregnancy
ICD10CM	O22.3	Deep phlebothrombosis in pregnancy
ICD10CM	O87.1	Deep phlebothrombosis in the puerperium
Coding system ERASMUS <sup>b</sup>	Code	Code name
ICD10	126	Pulmonary embolism
ICD10CM	126	Pulmonary embolism
ICD10	126.0	Pulmonary embolism with mention of acute cor pulmonale
ICD10CM	126.0	Pulmonary embolism with acute cor pulmonale

TABLE A1	(Continu	ed)
Coding system ERASMUS <sup>b</sup>	Code	Code name
ICD10CM	126.02	Saddle embolus of pulmonary artery with acute cor pulmonale
ICD10CM	126.09	Other pulmonary embolism with acute cor pulmonale
ICD10	126.9	Pulmonary embolism without mention of acute cor pulmonale
ICD10CM	126.9	Pulmonary embolism without acute cor pulmonale
ICD10CM	126.92	Saddle embolus of pulmonary artery without acute cor pulmonale
ICD10CM	126.93	Single subsegmental pulmonary embolism without acute cor pulmonale
ICD10CM	126.94	Multiple subsegmental pulmonary emboli without acute cor pulmonale
ICD10CM	126.99	Other pulmonary embolism without acute cor pulmonale
ICD10CM	180.21	Phlebitis and thrombophlebitis of iliac vein
ICD10CM	180.219	Phlebitis and thrombophlebitis of unspecified iliac vein
ICD10	182.2	Embolism and thrombosis of vena cava
		(Continues)

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#### TABLE A1 (Continued)

Coding system ERASMUS <sup>b</sup>	Code	Code name
ICD10CM	182.2	Embolism and thrombosis of vena cava and other thoracic veins
ICD10CM	182.21	Embolism and thrombosis of superior vena cava
ICD10CM	182.210	Acute embolism and thrombosis of superior vena cava
ICD10CM	182.211	Chronic embolism and thrombosis of superior vena cava
ICD10CM	182.22	Embolism and thrombosis of inferior vena cava
ICD10CM	182.220	Acute embolism and thrombosis of inferior vena cava
ICD10CM	182.221	Chronic embolism and thrombosis of inferior vena cava
ICD10	182.3	Embolism and thrombosis of renal vein
ICD10CM	182.3	Embolism and thrombosis of renal vein
ICD10CM	182.4	Acute embolism and thrombosis of deep veins of lower extremity
ICD10CM	182.40	Acute embolism and thrombosis of unspecified deep veins of lower extremity
ICD10CM	182.401	Acute embolism and thrombosis of unspecified deep veins of right lower extremity
ICD10CM	182.402	Acute embolism and thrombosis of unspecified deep veins of left lower extremity
ICD10CM	182.403	Acute embolism and thrombosis of unspecified deep veins of lower extremity, bilateral
ICD10CM	182.409	Acute embolism and thrombosis of unspecified deep veins of unspecified lower extremity
ICD10CM	182.41	Acute embolism and thrombosis of femoral vein
ICD10CM	182.411	Acute embolism and thrombosis of right femoral vein
ICD10CM	182.412	Acute embolism and thrombosis of left femoral vein
ICD10CM	182.413	Acute embolism and thrombosis of femoral vein, bilateral
ICD10CM	182.419	Acute embolism and thrombosis of unspecified femoral vein
ICD10CM	182.42	Acute embolism and thrombosis of iliac vein
ICD10CM	182.421	Acute embolism and thrombosis of right iliac vein
ICD10CM	182.422	Acute embolism and thrombosis of left iliac vein
ICD10CM	182.423	Acute embolism and thrombosis of iliac vein, bilateral
ICD10CM	182.429	Acute embolism and thrombosis of unspecified iliac vein
ICD10CM	182.43	Acute embolism and thrombosis of popliteal vein
ICD10CM	182.431	Acute embolism and thrombosis of right popliteal vein

TABLE A1

ERASMUS<sup>b</sup>

ICD10CM

Coding system

(Continued)

Code name

popliteal vein

vein, bilateral

vein

vein

tibial vein

vein. bilateral

extremity

bilateral

extremity

lower extremity

lower extremity

extremity, bilateral

lower extremity

extremity, bilateral

extremity

Acute embolism and thrombosis of left

Acute embolism and thrombosis of unspecified popliteal vein

Acute embolism and thrombosis of tibial

Acute embolism and thrombosis of right

Acute embolism and thrombosis of tibial

Acute embolism and thrombosis of unspecified tibial vein

Acute embolism and thrombosis of

Acute embolism and thrombosis of left tibial

other specified deep vein of lower extremity

Acute embolism and thrombosis of other

Acute embolism and thrombosis of other specified deep vein of left lower extremity

Acute embolism and thrombosis of other specified deep vein of lower extremity,

Acute embolism and thrombosis of other specified deep vein of unspecified lower

Acute embolism and thrombosis of unspecified deep veins of proximal lower extremity

Acute embolism and thrombosis of unspecified deep veins of right proximal

Acute embolism and thrombosis of unspecified deep veins of left proximal

Acute embolism and thrombosis of

Acute embolism and thrombosis of unspecified deep veins of unspecified

proximal lower extremity

unspecified deep veins of proximal lower

Acute embolism and thrombosis of unspecified

unspecified deep veins of left distal lower

deep veins of distal lower extremity

Acute embolism and thrombosis of unspecified deep veins of right distal

Acute embolism and thrombosis of

Acute embolism and thrombosis of unspecified deep veins of distal lower

specified deep vein of right lower

Acute embolism and thrombosis of popliteal

Code

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182.439

182.44

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182.49

182.491

182.492

182.493

182.499

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182.4Y2

182.4Y3

182.4Y9

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182.4Z2

182.4Z3

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#### TABLE A1 (Continued)

Coding system ERASMUS <sup>b</sup>	Code	Code name
ICD10CM	182.4Z9	Acute embolism and thrombosis of unspecified deep veins of unspecified distal lower extremity
ICD10CM	182.5	Chronic embolism and thrombosis of deep veins of lower extremity
ICD10CM	182.50	Chronic embolism and thrombosis of unspecified deep veins of lower extremity
ICD10CM	182.501	Chronic embolism and thrombosis of unspecified deep veins of right lower extremity
ICD10CM	182.502	Chronic embolism and thrombosis of unspecified deep veins of left lower extremity
ICD10CM	182.503	Chronic embolism and thrombosis of unspecified deep veins of lower extremity, bilateral
ICD10CM	182.509	Chronic embolism and thrombosis of unspecified deep veins of unspecified lower extremity
ICD10CM	182.59	Chronic embolism and thrombosis of other specified deep vein of lower extremity
ICD10CM	182.591	Chronic embolism and thrombosis of other specified deep vein of right lower extremity
ICD10CM	182.592	Chronic embolism and thrombosis of other specified deep vein of left lower extremity
ICD10CM	182.593	Chronic embolism and thrombosis of other specified deep vein of lower extremity, bilateral
ICD10CM	182.599	Chronic embolism and thrombosis of other specified deep vein of unspecified lower extremity
ICD10CM	182.5Y	Chronic embolism and thrombosis of unspecified deep veins of proximal lower extremity
ICD10CM	I82.5Y1	Chronic embolism and thrombosis of unspecified deep veins of right proximal lower extremity
ICD10CM	182.5Y2	Chronic embolism and thrombosis of unspecified deep veins of left proximal lower extremity
ICD10CM	I82.5Y3	Chronic embolism and thrombosis of unspecified deep veins of proximal lower extremity, bilateral
ICD10CM	I82.5Y9	Chronic embolism and thrombosis of unspecified deep veins of unspecified proximal lower extremity
ICD10CM	182.62	Acute embolism and thrombosis of deep veins of upper extremity
ICD10CM	182.621	Acute embolism and thrombosis of deep veins of right upper extremity
ICD10CM	182.622	Acute embolism and thrombosis of deep veins of left upper extremity
ICD10CM	182.623	Acute embolism and thrombosis of deep veins of upper extremity, bilateral

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Coding system ERASMUS <sup>b</sup>	Code	Code name
ICD10CM	182.629	Acute embolism and thrombosis of deep veins of unspecified upper extremity
ICD10CM	182.81	Embolism and thrombosis of superficial veins of lower extremities
ICD10CM	182.811	Embolism and thrombosis of superficial veins of right lower extremity
ICD10CM	182.812	Embolism and thrombosis of superficial veins of left lower extremity
ICD10CM	182.813	Embolism and thrombosis of superficial veins of lower extremities, bilateral
ICD10CM	182.819	Embolism and thrombosis of superficial veins of unspecified lower extremity
ICD10CM	182.A	Embolism and thrombosis of axillary vein
ICD10CM	182.A1	Acute embolism and thrombosis of axillary vein
ICD10CM	l82. A11	Acute embolism and thrombosis of right axillary vein
ICD10CM	l82. A12	Acute embolism and thrombosis of left axillary vein
ICD10CM	l82. A13	Acute embolism and thrombosis of axillary vein, bilateral
ICD10CM	l82. A19	Acute embolism and thrombosis of unspecified axillary vein
ICD10CM	182.B	Embolism and thrombosis of subclavian vein
ICD10CM	l82.B1	Acute embolism and thrombosis of subclavian vein
ICD10CM	l82. B11	Acute embolism and thrombosis of right subclavian vein
ICD10CM	l82. B12	Acute embolism and thrombosis of left subclavian vein
ICD10CM	l82. B13	Acute embolism and thrombosis of subclavian vein, bilateral
ICD10CM	l82. B19	Acute embolism and thrombosis of unspecified subclavian vein
ICD10CM	182.C	Embolism and thrombosis of internal jugular vein
ICD10CM	l82.C1	Acute embolism and thrombosis of internal jugular vein
ICD10CM	l82. C11	Acute embolism and thrombosis of right internal jugular vein
ICD10CM	l82. C12	Acute embolism and thrombosis of left internal jugular vein
ICD10CM	l82. C13	Acute embolism and thrombosis of internal jugular vein, bilateral
ICD10CM	182.	Acute embolism and thrombosis of

TABLE A1

(Continued)

<sup>a</sup>ACCESS included all subcodes, whereas ERASMUS only used the listed codes and not any unlisted codes.

unspecified internal jugular vein

C19

<sup>b</sup>ICD-10 codes not included by ERASMUS as compared to ACCESS are: 126.01, 126.90, 180 and subcodes except 180.21 and 180.219, 181 and subcodes, 182, 182.0, 182.1, 182.29, 182.290, 182.291, 182.51-182.56 and 182.5Z and subcodes, 182.6-182.7 and 182.9 and subcodes except 182.62 and subcodes, 182.B2 and 182.C2 and subcodes O08.02, O22.3, O87.1.

H_ADIAP_H	IR_upr IR_lwr IR	8.401 3.683 5.691	8.558 3.935 5.921	42.831 27.519 34.551	34.043 20.090 26.391	65.147 49.371 56.852	50.089 35.527 42.344	83.982 66.827 75.039	114.790 94.446 104.247	142.046 117.867 129.535	214.288 184.456 198.953	257.495 222.594 239.569	351.061 309.225 329.646	479.374 425.596 451.886	495.524 437.671 465.925	708.428 6 637.169 5 672.090 5	11.455 28.101 68.635
SIDIAP	IR_upr	3.965	2.158	26.031	16.999	35.844	33.165	52.101	71.404	80.333	129.702	156.578	209.455	325.792	319.963	506.255 4	132.463
	IR_lwr	2.114	0.874	19.069	11.348	28.958	26.360	44.267	62.204	69.831	115.971	140.480	189.719	298.792	290.318	469.686	87.497
	≌	2.933	1.412	22.349	13.962	32.264	29.616	48.065	66.685	74.945	122.692	148.365	199.404	312.073	304.870	487.713 4	409.517
ΓΡΟ ΙΤΑΓΥ	IR_upr	18.533	20.361	29.521	35.718	56.030	60.506	108.185	102.275	129.371	173.693	222.919	254.114	482.714	409.460	728.965	333.780
	IR_lwr	3.125	3.973	13.952	15.924	35.379	34.264	83.512	72.635	103.875	136.809	186.667	206.169	424.357	341.139	648.693	522.077
	≌	8.515	9.882	20.679	24.377	44.830	46.019	95.254	86.515	116.101	154.431	204.192	229.207	452.832	374.134	687.952	575.907
LPD	IR_upr	3.078	2.431	14.368	12.986	20.481	22.188	28.467	43.741	36.220	65.696	61.570	90.441	100.744	114.658	153.878	50.461
FRANCE	IR_lwr	0.803	0.474	7.661	5.259	13.299	12.565	20.533	31.514	27.612	51.922	49.968	74.395	83.309	93.896	127.632	18.459
	≌	1.674	1.180	10.629	8.496	16.604	16.875	24.259	37.256	31.699	58.507	55.543	82.125	91.717	103.890	140.297	33.747
IPCI	IR_upr	18.781	9.74	124.661	53.590	213.464	95.729	326.663	207.130	310.978	333.740	424.226	481.229	740.676	603.106	886.734	337.853
	IR_lwr	10.165	3.853	92.163	33.065	171.910	66.879	280.467	168.438	268.142	288.068	369.025	421.215	655.350	522.530	761.507	89.366
	≌	13.990	6.308	107.500	42.416	191.848	80.346	302.907	187.038	288.966	310.275	395.905	450.474	697.035	561.737	822.337 7	60.906
DA German)	/ IR_upr	8.974	4.870	50.190	41.029	78.340	71.979	102.490	121.669	118.153	166.851	175.389	226.173	278.467	278.472	394.823	319.480
	IR_lwr	5.575	2.439	40.803	30.197	67.861	58.776	91.478	106.356	108.009	152.551	161.889	208.809	260.228	258.221	367.633	90.473
	Щ	7.126	3.502	45.315	35.305	72.960	65.128	96.867	113.819	112.995	159.581	168.538	217.361	269.231	268.203	381.046	304.717
CPRD	IR_upr	13.048	4.148	94.268	32.978	118.780	90.853	133.897	145.978	156.620	197.799	268.255	324.970	477.566	488.495	717.938	522.249
	IR_lwr	8.636	1.849	77.761	23.299	101.457	74.995	116.096	126.803	138.019	176.771	240.892	294.459	435.655	443.471	652.590	547.731
	≌	10.674	2.831	85.718	27.827	109.863	82.640	124.759	136.138	147.099	187.064	254.297	309.433	456.250	465.575	684.679	84.100
	gender	. Female	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female N	1ale
	age_gr	0-19	0-19	20-29	20-29	30-39	30-39	40-49	40-49	50-59	50-59	60-69	69-69	62-02	70-79	80+ 8	+0

TABLE A2 Age-gender-stratified IRs per 100 000 person-years (with 95% CIs) for VTE for the databases provided by ERASMUS.

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3 Age
<b>TABLE A</b>

IT_PEDIANET	IR_upr 2. IR_lwr 0 r IR 0	781 3. 0.	872 018 695														
UK_CPRD	IR_upr	4.880	3.650	64.120	30.740	128.380	80.630	168.040	140.790	200.070	196.850	325.070	305.510	541.160	416.750	639.560 4	157.340
	IR_lwr	2.580	1.740	51.520	22.360	111.950	67.700	149.330	123.930	180.140	177.260	296.480	277.500	499.000	377.750	583.400	399.390
	Я	3.600	2.570	57.570	26.300	119.960	73.950	158.480	132.160	189.910	186.860	310.530	291.250	519.760	396.890	610.990 4	ł27.630
SIDIAP	R	2.930	1.410	22.350	13.960	32.260	29.620	48.060	66.690	74.940	122.690	148.370	199.400	312.070	304.870	487.710 4	t09.520
NL_PHARMO	IR_upr	33.180	26.760	117.900	79.800	216.990	127.630	300.050	224.150	321.150	325.390	436.910	549.590	768.650	613.540	850.010	332.610
PCHOSP	IR_lwr	10.810	7.470	60.230	33.710	136.590	66.960	211.410	148.350	234.340	238.820	327.790	426.980	601.030	459.250	625.020	555.830
	R	19.780	14.960	85.540	53.180	173.360	93.730	252.860	183.360	275.210	279.620	379.440	485.410	681.000	532.240	731.090	683.890
NL_PHARMO	IR_upr	5.490	3.320	14.800	7.100	19.500	13.270	29.410	28.410	40.710	65.520	73.750	109.220	124.250	146.060	87.750	94.100
HOSP	IR_lwr	3.240	1.660	10.540	4.110	14.850	9.190	23.840	22.650	34.100	56.820	63.780	96.600	109.440	128.960	75.850	78.940
	Щ	4.260	2.390	12.530	5.460	17.050	11.090	26.520	25.410	37.300	61.050	68.630	102.760	116.670	137.310	81.640	86.280
IT_ARS	IR_upr	7.960	9.680	24.190	31.090	63.250	46.430	88.570	96.200	113.730	179.620	221.040	308.280	483.490	521.680	884.070	328.570
	IR_lwr	3.600	4.950	13.700	19.510	47.120	32.580	73.010	79.640	96.410	157.230	194.180	274.950	442.240	475.000	824.970 7	755.530
	R	5.470	7.030	18.400	24.800	54.740	39.050	80.510	87.630	104.800	168.140	207.290	291.260	462.520	497.930	854.140 7	791.420
ES_SIDIAP_	IR_upr	13.200	12.910	66.990	42.210	131.210	83.290	172.120	163.870	253.610	301.970	448.040	467.950	850.560	727.520 1	1154.280	24.290
PCHOSP	IR_lwr	6.920	6.880	46.800	26.820	106.880	64.330	146.590	139.830	219.720	265.000	397.970	413.990	771.920	648.100 1	1055.760	311.170
	щ	9.690	9.550	56.220	33.870	118.580	73.350	158.970	151.490	236.210	283.030	422.450	440.350	810.520	686.950 1	1104.190	366.350
ES_SIDIAP_P(	C IR_upr	8.370	7.380	55.390	33.200	102.730	65.050	146.040	131.130	203.940	232.820	370.020	360.970	677.690	515.690	924.290	688.350
	IR_lwr	5.730	4.970	45.710	25.880	91.470	56.240	133.670	119.820	187.910	215.680	345.810	335.930	640.150	479.820	878.070	536.240
	R	6.960	6.090	50.380	29.370	96.980	60.520	139.750	125.380	195.800	224.130	357.760	348.280	658.720	497.510	900.960	61.910
ES_FISABIO	IR_upr	8.930	11.360	44.490	30.200	84.110	66.510	111.720	122.540	157.240	246.670	303.970	405.680	591.720	614.000	995.190	35.500
	IR_lwr	6.110	8.240	35.450	22.890	73.280	56.930	100.300	110.840	142.870	228.450	282.010	378.810	557.590	576.050	943.420	371.940
	Щ	7.420	9.700	39.780	26.350	78.560	61.580	105.900	116.580	149.930	237.430	292.840	392.070	574.470	594.800	969.050	903.300
ES_BIFAP_	IR_upr	10.140	9.450	58.450	37.620	118.620	72.350	167.140	134.890	228.930	246.740	420.460	392.630	722.730	578.150	891.520 6	684.130
PCHOSP	IR_lwr	5.900	5.460	44.020	26.210	100.760	58.420	147.770	117.620	205.620	222.610	384.980	358.170	670.840	528.900	839.780	\$27.510
	Я	7.810	7.250	50.850	31.530	109.420	65.110	157.230	126.030	217.040	234.440	402.430	375.100	696.430	553.110	865.360	55.360
ES_BIFAP_PC	: IR_upr	7.980	5.980	49.790	28.510	100.760	63.040	147.680	117.380	204.810	206.660	362.580	320.880	625.820	456.320	798.300	583.870
	IR_lwr	5.890	4.230	42.430	23.010	91.690	55.840	137.640	108.470	192.540	194.280	344.240	303.160	598.780	431.280	768.780	551.730
	씸	6.880	5.050	46.000	25.650	96.150	59.360	142.590	112.860	198.600	200.400	353.320	311.920	612.190	443.670	783.430	567.630
	gender Fe	imale M	lale F	emale	Male	Female N	1ale F	<sup>-</sup> emale N	Male F	emale	Male F	emale N	1ale F	<sup>-</sup> emale N	Vale F	<sup>-</sup> emale N	۱ale
	age_gr 0-	-19 0-	-19 2	0-29	20-29	30-39 3	0-39 4	10-49 2	t0-49 5	0-59	50-59 6	69-69	0-69 7	70-79 7	70-79	80+ 8	+0

**TABLE A4** Age-gender IR estimates and CIs for VTE from metaanalyses.

Age	Gender	Estimate	Lower	Upper
0-19	Female	5.697	4.095	7.924
0-19	Male	4.322	2.911	6.418
20-29	Female	32.773	20.772	51.707
20-29	Male	21.825	14.778	32.231
30-39	Female	62.316	38.087	101.957
30-39	Male	44.485	29.621	66.807
40-49	Female	93.155	57.581	150.707
40-49	Male	89.761	61.579	130.841
50-59	Female	118.074	75.574	184.476
50-59	Male	157.258	111.909	220.984
60-69	Female	203.979	133.110	312.582
60-69	Male	242.021	171.372	341.798
70-79	Female	372.142	237.184	583.891
70-79	Male	340.595	234.922	493.800
80+	Female	495.147	290.933	842.703
80+	Male	421.791	258.186	689.069

**TABLE A5** Age-gender-stratified  $l^2$  and  $\tau^2$  estimates from meta-analyses for the different sensitivity analyses.

		PC-H linka	PC-H linkage		ICD-10 code		ACCESS		ERASMUS	
Age	Gender	l <sup>2</sup>	$\tau^2$	l <sup>2</sup>	$\tau^2$	l <sup>2</sup>	$\tau^2$	l <sup>2</sup>	τ <sup>2</sup>	
0-19	Female	0.771	0.145	0.894	0.506	0.851	0.177	0.944	0.521	
0-19	Male	0.539	0.019	0.928	0.588	0.927	0.309	0.896	0.479	
20-29	Female	0.942	0.293	0.961	0.328	0.975	0.345	0.988	0.573	
20-29	Male	0.731	0.042	0.936	0.245	0.950	0.351	0.966	0.290	
30-39	Female	0.971	0.178	0.985	0.481	0.990	0.443	0.994	0.570	
30-39	Male	0.904	0.084	0.954	0.208	0.980	0.383	0.983	0.250	
40-49	Female	0.985	0.184	0.991	0.485	0.993	0.406	0.996	0.554	
40-49	Male	0.958	0.070	0.976	0.204	0.990	0.322	0.990	0.235	
50-59	Female	0.988	0.149	0.994	0.463	0.995	0.366	0.995	0.392	
50-59	Male	0.970	0.042	0.991	0.218	0.994	0.213	0.992	0.199	
60-69	Female	0.989	0.086	0.997	0.445	0.996	0.321	0.995	0.305	
60-69	Male	0.968	0.035	0.995	0.286	0.995	0.205	0.994	0.215	
70-79	Female	0.988	0.045	0.998	0.468	0.997	0.335	0.996	0.304	
70-79	Male	0.951	0.014	0.997	0.320	0.999	0.226	0.994	0.232	
80+	Female	0.967	0.019	0.998	0.368	0.998	0.631	0.994	0.228	
80+	Male	0.970	0.018	0.997	0.343	0.997	0.521	0.991	0.222	