

SUPPLEMENTARY INFORMATION

A register-based cohort study on the effectiveness and Safety of anti-PCSK9 treatment in persons with hyperlipidemia

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Supplementary Methods

WebI

SAP Web Intelligence (WebI) is a tool that is integrated into the EPIC electronic medical record system. It allows users to create custom reports and analyses of data stored in the EPIC Clarity database. With WebI, users can build queries using a wide range of predefined variables and filters. WebI can generate reports in a variety of formats including tables, graphs, and maps.

Mgcv

In the present study, we used generalized additive models (GAMs) as implemented in the *mgcv* package in R to analyze the data. GAMs are a flexible class of regression models that allow for the incorporation of non-linear relationships between the response variable and predictor variables. They are particularly useful for modeling complex, non-linear trends in data and can be seen as an extension of linear regression models. The *mgcv* package is a popular and widely used tool for fitting GAMs, and it provides a range of options for specifying and estimating model parameters. Overall, the use of GAMs and the *mgcv* package allowed us to accurately model and analyze the data, leading to insights and conclusions that would not have been possible with traditional linear regression models. To check the model fit of a GAM fitted with the *mgcv* package, the `gam.check()` function can be used. This function produces a plot that displays the residuals of the model, as well as a number of diagnostic plots that can be used to assess the model fit. The `gam.check()` function produces a plot that displays the residuals of the model on the left hand side, and a number of diagnostic plots on the right hand side. The residuals plot shows the residuals of the model as a function of the fitted values. Ideally, the residuals should be randomly distributed around zero, with no discernible pattern. If there is a pattern present in the residuals plot, it may indicate that the model is not adequately fitting the data. In *mgcv*, most smooths are required to satisfy a sum-to-zero identifiability constraint, which allows for the inclusion of an intercept in the model, especially when the model includes factor parametric terms. This constraint centers the smooth around 0 on the axis. The 0 line represents the average value of the response (or the reference levels for factor parametric terms) on the link scale.

Definitions

Atherosclerotic cardiovascular disease (ASCVD):

Hypercholesterolemia criteria:

Familial hypercholesterolemia (FH):

Procedures prior to inclusion:

Procedure:	Procedure code:	Operation code:
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Percutaneous coronary intervention (PCI)	None	KFNG and KZFFX01
Coronary artery bypass grafting (CABG)	None	KFNA, KFNB, KFNC, KFND, KFNE, KFNH20 and KFNF
Angioplasty and stent placement	None	KPAN, KPAP, KPBN, KPBP, KPCN, KPCP, KPDN, KPDP, KPEN, KPEP, KPFN and KPFP
Endarterectomy	None	KPEF, KPAF, KPBF, KPCF and KPDF
Thrombectomy	None	KAAL11, KFAB10, KPAE10, KPAE25, KPAE30, KPAE99, KPAU74, KPBE10, KPBE20, KPBE30, KPBE99, KPCE30, KPCE40, KPCE99, KPDE10, KPDE30, KPDU74, KPDE10, KPDE11, KPDE12, KPFE10, KPFE30, KPHE22, KPHE23, KPHE25, KPHE30, KPHE31 and KPHE99

Comorbidities prior to inclusion:

Cardiovascular comorbidities:

Disease:	Diagnostic code:	Operation code:	ATC-code:
Diabetes (DM)	E10, E11 and N083	None	A10
Diabetes target organ damage	H360, E103, E113, G590, E104, E114, N083, E102 and E112	None	None
Chronic kidney disease (CKD)	E102, E112, I120, N03, N04, N05, N06, N07, N08, N11, N12, N162, N163, N164, N182, N183, N184, N185, N26 and Q61	None	None
Chronic obstructive pulmonary disease	J43 and J44	None	None
Hypertension	I10, I11, I12, I13 and I15	None	C07, C08, C09, C03A, C03B, C03C, C03EA, C03EB, C02L, C10BX, C03DA, C03DB, C03E, C02DB, C02DC, C02DD, C02DG, C02A, C02B and

			C02C (excluding C02AC02)
Heart failure	I099A, I110, I130, I132, I420 and I50	None	None
Cancer, ex. non-melanoma skin cancer	C00, C01, C02, C03, C04, C05, C06, C07, C08, C09, C10, C11, C12, C13, C14, C15, C16, C17, C18, C20, C21, C22, C23, C24, C25, C26, C30, C31, C32, C33, C34, C37, C38, C39, C40, C41, C43, C45, C46, C47, C48, C49, C50, C51, C52, C53, C54, C55, C56, C57, C58, C60, C61, C62, C63, C64, C65, C66, C67, C68, C69, C70, C71, C72, C73, C74, C75, C76, C77, C78, C79, C80, C81, C82, C83, C84, C85, C86, C88, C90, C91, C92, C93, C94, C95 and C96	None	None
Liver disease	K70, K71, K72, K73, K74, K75, K76, K77, B15, B16, B17, B18, B19 and B581	None	None
Rheumatoid arthritis	M05, M060, M069 and M080	None	None

Laboratory values:

Biomarker:	Laboratory code:
Low-density lipoprotein cholesterol (LDL-C)	NPU01568, NPU10171, AAB00101, AAB00102 and DNK35308
High-density lipoprotein cholesterol (HDL-C)	NPU01567 and NPU10157
Total cholesterol (TC)	NPU18412 and NPU01566
Triglycerides (TGs)	NPU04094 and NPU03620
Lipoprotein a (Lp[a])	NPU21687, NPU19840 and NPU58475
Apolipoprotein B (Apo-B)	NPU22299 and NPU19697

Cholesterol-lowering drug group classification:

Contains the ATC-codes used to define the cholesterol-lowering drug groups used in multiple endpoints.

Drug group:	ATC-code:
Statin	C10AA, C10BA01, C10BA02, C10BA03, C10BA04, C10BA05, C10BA06, C10BA07, C10BA08, C10BA09 and C10BX
PCSK9i	C10AX13 and C10AX14
Ezetimibe	C10AX09, C10BA02, C10BA05, C10BA06 and C10BA10
Bile acid sequestrants	C10AC01, C10AC02, C10AC03 and C10AC04
Fibrates	C10AB, C10BA03, C10BA04 and C10BA09