S1 Checklist

STROBE Statement—Checklist of items that should be included in reports of *cohort studies*

	Item No	Recommendation	Section
Title and abstract	1	(a) Indicate the study's design with a	YES
		commonly used term in the title or the	Title
		abstract	
		(b) Provide in the abstract an informative	YES
		and balanced summary of what was done	Abstract
		and what was found	
Introduction			
Background/rationale	2	Explain the scientific background and	YES
		rationale for the investigation being reported	Intro,
			paragraphs 1 & 2
Objectives	3	State specific objectives, including any	YES
J		prespecified hypotheses	Intro,
			paragraphs 3 & 4
Methods			
Study design	4	Present key elements of study design early	YES
		in the paper	M&M, Study design and patient
			characteristics, paragraph 1
Setting	5	Describe the setting, locations, and relevant	YES
		dates, including periods of recruitment,	M&M, Study design and patient
		exposure, follow-up, and data collection	characteristics
Participants	6	(a) Give the eligibility criteria, and the	YES
		sources and methods of selection of	M&M, Study design and patient
		participants. Describe methods of follow-up	characteristics, paragraphs 2-5
		(b) For matched studies, give matching	Not applicable
		criteria and number of exposed and	
		unexposed	
Variables	7	Clearly define all outcomes, exposures,	YES
		predictors, potential confounders, and effect	M&M, Study design and patient
		modifiers. Give diagnostic criteria, if	characteristics, paragraphs 3-5
		applicable	Time-to-event outcomes,
			Survival analysis
Data sources/	8*	For each variable of interest, give sources	YES
measurement		of data and details of methods of assessment	M&M, Study design and patient
		(measurement). Describe comparability of	characteristics, paragraphs 2 & 6,
		assessment methods if there is more than	Anti-drug antibody assays,
		one group	CXCL12 laboratory test, DNA
			extraction and genotyping
Bias	9	Describe any efforts to address potential	YES
		sources of bias	M&M, Survival analysis,
			paragraph 3
Study size	10	Explain how the study size was arrived at	Not applicable (real-life
		•	prospective study)
Quantitative variables	11	Explain how quantitative variables were	YES
		handled in the analyses. If applicable,	M&M, Survival analysis,
		describe which groupings were chosen and	paragraph 1
		3 1 0	

	why	
12	(a) Describe all statistical methods,	YES
	including those used to control for	M&M, Survival analysis,
	confounding	Mediation analysis
	(b) Describe any methods used to examine	Not applicable
	subgroups and interactions	
	(c) Explain how missing data were	YES
	addressed	M&M, Survival analysis,
		paragraph 4
	(d) If applicable, explain how loss to follow-	YES
	up was addressed	M&M, Time-to-event outcomes
	(e) Describe any sensitivity analyses	Not applicable
13*	(a) Report numbers of individuals at each	YES
	- · · ·	M&M
		Flowchart Fig 1
		6
		YES
		M&M
		Flowchart Fig 1
	(c) Consider use of a flow diagram	YES
	(e) consider use of a now diagram	M&M
		Flowchart Fig 1
14*	(a) Give characteristics of study participants	YES
	• • •	Results, Demographic and
		clinical characteristics,
	• •	paragraphs 1-3
	Comoditation of the company of the c	Table1
		S1 Table
	(b) Indicate number of participants with	YES
	- · ·	Flowchart Fig 1
	missing data for each variable of interest	S2 Table
	(c) Summarise follow-up time (ag average	YES
		Table 1
15*	·	YES
15.	_	Results, Demographic
	summary measures over time	and clinical characteristics,
		paragraph 4 Table 2
	(a) Give unadjusted estimates and, if	YES
16	(a) Orve unaujusteu estimates and, n	I L'O
16	applicable confounder adjusted astimates	Paculte Association between
16	applicable, confounder-adjusted estimates	Results, Association between
16	and their precision (eg, 95% confidence	bio-clinical variables and ADA
16	and their precision (eg, 95% confidence interval). Make clear which confounders	bio-clinical variables and ADA occurrence, Genetic variants
16	and their precision (eg, 95% confidence interval). Make clear which confounders were adjusted for and why they were	bio-clinical variables and ADA occurrence, Genetic variants associated with ADAs, CXCL12
16	and their precision (eg, 95% confidence interval). Make clear which confounders	bio-clinical variables and ADA occurrence, Genetic variants associated with ADAs, CXCL12 serum level analysis
16	and their precision (eg, 95% confidence interval). Make clear which confounders were adjusted for and why they were	bio-clinical variables and ADA occurrence, Genetic variants associated with ADAs, CXCL12
	13* 14*	subgroups and interactions (c) Explain how missing data were addressed (d) If applicable, explain how loss to follow-up was addressed (e) Describe any sensitivity analyses 13* (a) Report numbers of individuals at each stage of study—eg numbers potentially eligible, examined for eligibility, confirmed eligible, included in the study, completing follow-up, and analysed (b) Give reasons for non-participation at each stage (c) Consider use of a flow diagram 14* (a) Give characteristics of study participants (eg demographic, clinical, social) and information on exposures and potential confounders (b) Indicate number of participants with missing data for each variable of interest (c) Summarise follow-up time (eg, average and total amount)

		(b) Report category boundaries when	YES
		continuous variables were categorized	M&M, Survival analysis,
			paragraph 1
		(c) If relevant, consider translating estimates	YES
		of relative risk into absolute risk for a	Results, Association between
		meaningful time period	bio-clinical variables and ADA
			occurrence, paragraph 4
			(mediation analysis)
Other analyses	17	Report other analyses done—eg analyses of	YES
		subgroups and interactions, and sensitivity	Results, CXCL12 serum level
		analyses	analysis
Discussion			
Key results	18	Summarise key results with reference to	YES
		study objectives	Disc., paragraph 1
Limitations	19	Discuss limitations of the study, taking into	YES
		account sources of potential bias or	Disc., paragraphs 12-13
		imprecision. Discuss both direction and	
		magnitude of any potential bias	
Interpretation	20	Give a cautious overall interpretation of	YES
		results considering objectives, limitations,	Disc., paragraphs 5-11
		multiplicity of analyses, results from similar	
		studies, and other relevant evidence	
Generalisability	21	Discuss the generalisability (external	YES
		validity) of the study results	Disc., paragraph 12
Other information			
Funding	22	Give the source of funding and the role of	YES
		the funders for the present study and, if	Funding statement
		applicable, for the original study on which	
		the present article is based	

^{*}Give information separately for exposed and unexposed groups.

Note: An Explanation and Elaboration article discusses each checklist item and gives methodological background and published examples of transparent reporting. The STROBE checklist is best used in conjunction with this article (freely available on the Web sites of PLoS Medicine at http://www.plosmedicine.org/, Annals of Internal Medicine at http://www.annals.org/, and Epidemiology at http://www.epidem.com/). Information on the STROBE Initiative is available at http://www.strobe-statement.org.