# nature portfolio

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## **Reporting Summary**

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our Editorial Policies and the Editorial Policy Checklist.

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101	an statistical analyses, commit that the following items are present in the figure regend, table regend, main text, or interious section.
n/a	Confirmed
	$\square$ The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	🔀 A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	The statistical test(s) used AND whether they are one- or two-sided  Only common tests should be described solely by name; describe more complex techniques in the Methods section.
	A description of all covariates tested
	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i> ) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable.</i>
$\boxtimes$	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
$\boxtimes$	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
$\boxtimes$	Estimates of effect sizes (e.g. Cohen's <i>d</i> , Pearson's <i>r</i> ), indicating how they were calculated
	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.
So	ftware and code

Policy information about availability of computer code

Data collection Microsoft Excel version 16.56

Data analysis

R version 3.5.1 statistical software (R Foundation for Statistical Computing, Vienna, Austria), GraphPad Prism 9 software (GraphPad Software

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

#### Data

Policy information about availability of data

All manuscripts must include a <u>data availability statement</u>. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

Provide your data availability statement here.

Human rese	arch part	icipants			
Policy information	about <u>studies</u>	involving human research participants and Sex and Gender in Research.			
Reporting on sex	and gender	sex			
Population chara	acteristics	Patients with systemic autoimmune rheumatic disease, 18-60 year old, who had received $\geq$ 4 weeks of $\geq$ 1 immunosuppressive drug at a stable dose [prednisolone <20 mg per day, MTX >10 mg per week, leflunomide (LEF) 20 mg per day, azathioprine (AZA) >50 mg per day, and MMF >1,000 mg per day]			
		Consecutive patients with SARDs who regular following up at rheumatology unit, of Songklanagarind hospital healthy groups were recruited by poster annoucement at vaccine center.			
Ethics oversight		Human Research Ethics Committee of the Faculty of Medicine, Prince of Songkla University (REC.64–421–14–1)			
Note that full informa	ation on the app	roval of the study protocol must also be provided in the manuscript.			
Field-spe					
	ne below that	is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.			
Life sciences		Behavioural & social sciences			
For a reference copy of	the document with	all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>			
Life scier	nces st	udy design			
All studies must dis	sclose on these	points even when the disclosure is negative.			
Sample size	mRNA vaccine titers were sign calculated this group with allo	aple size was determined by the data of a previous study by Geisen, U. M. et al. that evaluated the immunogenicity of anti-SARS-CoV-2 vaccines in 42 healthy controls and 26 patients with chronic inflammatory conditions undergoing immunosuppressive therapy. The IgG ere significantly lower in patients, compared with controls (2053 BAU/mL ±1218 vs. 2685±1102 BAU/mL; 24% reduction). We ed this using n4Studies program for testing finite population proportions. The calculated sample size (n) was 27 participants in each vith allow for a 10% dropout. The ratio between case and control is 1:1, and the final calculated sample size is 60 participants, 1:1 ratio, all participants are 60.			
Data exclusions	no				
Replication	no				
Randomization		his study was not a randomized control trial study. Age- and sex-matched healthy individuals with no medical history, nor any current nedication, and vaccinated under this regimen were recruited as controled groups.			
Blinding	this is a prospe	prospective case-control study. the blinding was not relevant to our study.			
We require informati	ion from authors	pecific materials, systems and methods about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material,			
system or method lis	ted is relevant to	your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.			
Materials & experimental systems		systems Methods			
n/a Involved in the study		n/a Involved in the study			
☐ ☐ Antihodies		MID Chip-sea			

Flow cytometry

MRI-based neuroimaging

Eukaryotic cell lines

Clinical data

Palaeontology and archaeology
Animals and other organisms

Dual use research of concern

#### **Antibodies**

Antibodies used We have provided details of all the antibodies used in this study in the online methods

Validation We have done a flow panel optimisation before carrying out the study

#### Clinical data

Policy information about clinical studies

All manuscripts should comply with the ICMJE guidelines for publication of clinical research and a completed CONSORT checklist must be included with all submissions.

Clinical trial registration | this study was not registered to clinical trial.gov

Study protocol The protocols were report in online methods

Data collection The trial was conducted according to the principles of Good Clinical Practice by Clinical Research Center of Faculty of Medicine,

Prince of Songkhla University

Estimated Study Start Date: October, 2021

Estimated Primary Completion Date: March31, 2022 Estimated Study Completion Date: April 30, 2022

Outcomes Primary outcome

- 1. Compared the seropositivity rate between SARD and healthy group
- 2 .Comapared the Anti-RBD Ab levels between SARD and healthy group
- 3. Evauated the neutralization of the emerging Omicron BA.2 VOC

Secondary outcome

1. Comapare the cellular immunogenicity of heterologous vaccination with CoronaVac (Sinovac Life Sciences, Beijing, China) followed

by ChAdOx1-nCoV-19 (Oxford-AstraZeneca) between SARD and healthy group

#### Flow Cytometry

### Plots

Confirm that:

 $\nearrow$  The axis labels state the marker and fluorochrome used (e.g. CD4-FITC).

The axis scales are clearly visible. Include numbers along axes only for bottom left plot of group (a 'group' is an analysis of identical markers).

All plots are contour plots with outliers or pseudocolor plots.

A numerical value for number of cells or percentage (with statistics) is provided.

#### Methodology

Sample preparation Flow cytometry analysis were carried out on cryopreserved PBMCs. Cells were thawed in media containing 5 U/mL of

benzonase and resuspended in complete RPMI media supplemented with 10% FCS, L-glutamine and penicillin-streptomycin

(R10).

Instrument The stained cell were analysed using CytoflexS Beckman

Software The acquired data was analysed using FlowJo Software (Version 10)

Cell population abundance 200,000 events were recorded. Data was exported and analysed using FlowJo Software

Gating strategy Gating strategy is provided in The supplementary information

X Tick this box to confirm that a figure exemplifying the gating strategy is provided in the Supplementary Information.