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Corresponding author(s): JUNG KYOON CHOI

Last updated by author(s): Nov 14, 2019

Reporting Summary

Nature Research 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 Research policies, see <u>Authors & Referees</u> and the <u>Editorial Policy Checklist</u>.

Statistics

For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.					
n/a	Cor	firmed			
		The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
	\square	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.			
	\square	A description of all covariates tested			
	\square	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)			
	\boxtimes	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 Give <i>P</i> values as exact values whenever suitable.			
\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			
	\square	Estimates of effect sizes (e.g. Cohen's d, Pearson's r), indicating how they were calculated			
	1	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.			

Software and code

Policy information about <u>availability of computer code</u>							
Data collection	All the softwares used in this study were described in Methods section.						
Data analysis	SMMPMBEC, ANN, NetMHCcon, NetMHCpan, BWA, GATK, MuTect, CHAT, CNVkit, Seq2HLA, SIFT, PROVEAN, randomForest (R package), Enrichr, Panther, glmnet (R package), ROCR (R package)						

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/reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.

Data

Policy information about availability of data

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

- Accession codes, unique identifiers, or web links for publicly available datasets
- A list of figures that have associated raw data
- A description of any restrictions on data availability

Results of neoantigen calling for our SMC cohort samples by the CNN and NetMHCpan methods were made available at the authors' webpage (http:// omics.kaist.ac.kr/resources).

Field-specific reporting

K Life sciences

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

Behavioural & social sciences Ecological, evolutionary & environmental sciences

For a reference copy of the document with all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>

Life sciences study design

 All studies must disclose on these points even when the disclosure is negative.

 Sample size
 Sample size was not calculated in this study.

 Data exclusions
 None.

 Replication
 Not relevant in this study.

 Randomization
 Not relevant in this study.

 Blinding
 Not relevant in this study.

Reporting for specific materials, systems and methods

Methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed 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

n/a	Involved in the study	n/a	Involved in the study
\boxtimes	Antibodies	\boxtimes	ChIP-seq
\boxtimes	Eukaryotic cell lines	\boxtimes	Flow cytometry
\boxtimes	Palaeontology	\boxtimes	MRI-based neuroimaging
\boxtimes	Animals and other organisms		
	Human research participants		
\boxtimes	Clinical data		
	•		

Human research participants

Policy information about studies involving human research participants

Population characteristics	A total of 122 advanced non-small cell lung carcinoma patients who were treated with anti-PD-1/PD-L1 from 2014 to 2017 at Samsung Medical Center were enrolled for this study
Recruitment	This study was done retrospectively and no patients were directly recruited.
Ethics oversight	This study was approved from the institutional review board at Samsung Medical Center (2018-03-130 and 2013-10-112).

Note that full information on the approval of the study protocol must also be provided in the manuscript.