

## Data Sharing Statement

<b>Article Info</b>	https://dx.doi.org/10.21037/tlcr-21-729	
<b>Item</b>	<b>Question</b>	<b>Authors' Response (place "-" if not applicable)</b>
1	Would you like to share data collected for your study to others?	Yes
2	If not, would you like to share the reason for your decision?	-
3	What data in particular will be shared?	All raw RNA-sequencing data.
4	Any other documents will be shared? Such as study protocol, statistical analysis plan, informed consent form, clinical study report, analytic code.	-
5	When will data availability begin?	From the publication date.
6	When will data availability end?	-
7	To whom will you share the data?	Scholars who are interested in studies of extracellular vesicle and tumor biomarkers.
8	For what type of analysis or purpose?	For analysis to evaluate the performance of exLRs based diagnostic signature for early lung cancer.
9	How or where can the data/documents be obtained?	All raw RNA-sequencing data have been deposited in the National Genomics Data Center (NGDC) under the accession code HRA002156 ( <a href="https://bigd.big.ac.cn/gsa-human/browse/HRA002156">https://bigd.big.ac.cn/gsa-human/browse/HRA002156</a> ). The deposited and publicly available data are compliant with the regulations of the Ministry of Science and

		Technology of the People's Republic of China. The raw sequencing data contain information unique to individuals and are available under controlled access.
10	Any other restrictions?	We may balance the potential benefits and risks for each request and then provide the data that could be shared.