

Data Sharing Statement

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| Article Info | https://dx.doi.org/10.21037/tcr-22-2608 | |
| Item | Question | Authors' Response (place "-" if not applicable) |
| 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? | The information of the eight genes should be specifically disclosed. This is because it is of great value in predicting patient prognosis. (Microarray data and sequence data from the FUSCC-TNBC cohort have been deposited in the Sequence Read Archive (WES and RNA-seq; SRA: SRP157974)) |
| 4 | Any other documents will be shared? Such as study protocol, statistical analysis plan, informed consent form, clinical study report, analytic code. | Statistical analysis plan, informed consent form, and clinical study report will also be shared if requested. |
| 5 | When will data availability begin? | From the publication date. |
| 6 | When will data availability end? | Two years within the publication date, since the technique or survival data may be updated over time. |
| 7 | To whom will you share the data? | Medical oncologists and breast surgeons who are interested in our study. |
| 8 | For what type of analysis or purpose? | For analysis to assess the accuracy of prognosis prediction using an eight-gene model in breast cancer patients. |
| 9 | How or where can the data/documents be obtained? | Emails could be sent to the address below to obtain the shared data: Suixinyi2021@163.com |
| 10 | Any other restrictions? | We may balance the potential benefits and risks for each request |

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| | | and then provide the data that could be shared. |
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