

ICMJJE DISCLOSURE FORM

Date: **Apr. 24th, 2022**

Your Name: **Jiang-Tao Mou**

Manuscript Title: **Identification of genetic polymorphisms in unexplained recurrent spontaneous abortion based on whole exome sequencing**

Manuscript number (if known): _____

In the interest of transparency, we ask you to disclose all relationships/activities/interests listed below that are related to the content of your manuscript. "Related" means any relation with for-profit or not-for-profit third parties whose interests may be affected by the content of the manuscript. Disclosure represents a commitment to transparency and does not necessarily indicate a bias. If you are in doubt about whether to list a relationship/activity/interest, it is preferable that you do so.

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9	Participation on a Data Safety Monitoring Board or Advisory Board	<input type="checkbox"/> <u>√</u> <input type="checkbox"/> None	
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11	Stock or stock options	<input type="checkbox"/> <u>√</u> <input type="checkbox"/> None	
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ICMJE DISCLOSURE FORM

Date: Apr. 24th, 2022

Your Name: Shi-Xing Huang

Manuscript Title: Identification of genetic polymorphisms in unexplained recurrent spontaneous abortion based on whole exome sequencing

Manuscript number (if known): _____

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ICMJJE DISCLOSURE FORM

Date: Apr. 24th, 2022

Your Name: Li-Li Yu

Manuscript Title: Identification of genetic polymorphisms in unexplained recurrent spontaneous abortion based on whole exome sequencing

Manuscript number (if known): _____

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ICMJE DISCLOSURE FORM

Date: Apr. 24th, 2022

Your Name: Jing Xu

Manuscript Title: Identification of genetic polymorphisms in unexplained recurrent spontaneous abortion based on whole exome sequencing

Manuscript number (if known): _____

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ICMJE DISCLOSURE FORM

Date: **Apr. 24th, 2022**

Your Name: **Qiao-Ling Deng**

Manuscript Title: **Identification of genetic polymorphisms in unexplained recurrent spontaneous abortion based on whole exome sequencing**

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Date: **Apr. 24th, 2022**

Your Name: Yi-Shan Xie

Manuscript Title: Identification of genetic polymorphisms in unexplained recurrent spontaneous abortion based on whole exome sequencing

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Date: Apr. 24th, 2022

Your Name: Kun Deng

Manuscript Title: Identification of genetic polymorphisms in unexplained recurrent spontaneous abortion based on whole exome sequencing

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