Date: _____Jul. 11th, 2022____

Your Name: _____Yu Pan_____

Manuscript Title: <u>Whole exome sequencing identifies novel inherited genetic variants in tetralogy of Fallot</u> 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.

The following questions apply to the author's relationships/activities/interests as they relate to the <u>current</u> <u>manuscript only</u>.

The author's relationships/activities/interests should be <u>defined broadly</u>. For example, if your manuscript pertains to the epidemiology of hypertension, you should declare all relationships with manufacturers of antihypertensive medication, even if that medication is not mentioned in the manuscript.

		Name all entities with whom you have this relationship or indicate none (add rows as needed)	Specifications/Comments (e.g., if payments were made to you or to your institution)
		Time frame: Since the initial	planning of the work
1	All support for the present manuscript (e.g., funding, provision of study materials, medical writing, article processing charges, etc.) No time limit for this item.	Guizhou Science and Technology Support Program (Guizhou Science and Technology Cooperation Support 2018-2783). National Natural Science Foundation of China (No.81860273)	
		Time frame: past	36 months
2	Grants or contracts from any entity (if not indicated in item #1 above).	None	

3	Royalties or licenses	None
3	Royalties of licenses	
4	Consulting fees	None
5	Payment or honoraria for	None
	lectures, presentations,	
	speakers bureaus,	
	manuscript writing or	
	educational events	
6	Payment for expert	None
	testimony	
7	Support for attending	None
	meetings and/or travel	
8	Patents planned, issued or	None
0	pending	None
	pending	
9	Participation on a Data	None
	Safety Monitoring Board or	
	Advisory Board	
10	Leadership or fiduciary role	None
	in other board, society, committee or advocacy	
	group, paid or unpaid	
11	Stock or stock options	None
12	Possint of aquinment	None
12	Receipt of equipment,	None
	materials, drugs, medical	
	writing, gifts or other	
	services	
13	Other financial or non- financial interests	None
		Letter and the second sec

This study was supported by the Guizhou Science and Technology Support Program (Guizhou Science and Technology Cooperation Support 2018-2783) and the National Natural Science Foundation of China (No.81860273).

Please place an "X" next to the following statement to indicate your agreement:

Date: _____Jul. 11th, 2022____

Your Name: <u>Manli Liu</u>

Manuscript Title: <u>Whole exome sequencing identifies novel inherited genetic variants in tetralogy of Fallot</u> Manuscript number (if known):

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		T :	
		Time frame: past	36 months
2	Grants or contracts from any entity (if not indicated in item #1 above).	None	
3	Royalties or licenses	None	
4	Consulting fees	None	
5	Payment or honoraria for	None	

6	lectures, presentations, speakers bureaus, manuscript writing or educational events Payment for expert testimony Support for attending meetings and/or travel	None None None None
8	Patents planned, issued or pending	None
9	Participation on a Data Safety Monitoring Board or Advisory Board	None
10	Leadership or fiduciary role in other board, society, committee or advocacy group, paid or unpaid	None
11	Stock or stock options	None
12	Receipt of equipment, materials, drugs, medical writing, gifts or other services	None
13	Other financial or non- financial interests	None

None.

Please place an "X" next to the following statement to indicate your agreement:

Date: _____Jul. 11th, 2022____

Your Name: <u>Songsong Zhang</u>

Manuscript Title: <u>Whole exome sequencing identifies novel inherited genetic variants in tetralogy of Fallot___</u> Manuscript number (if known):__

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3	Royalties or licenses	None	
4	Consulting fees	None	

5	Payment or honoraria for	None	
	lectures, presentations,		
	speakers bureaus,		
	manuscript writing or		
6	educational events	News	
6	Payment for expert	None	
	testimony		
7	Support for attending	None	
/	meetings and/or travel	None	
8	Patents planned, issued or	None	
-	pending		
	F		
9	Participation on a Data	None	
	Safety Monitoring Board or Advisory Board		
10	in other board, society, committee or advocacy	None	
	group, paid or unpaid		
11	Stock or stock options	None	
12	Receipt of equipment,	None	
	materials, drugs, medical		
	writing, gifts or other services		
42			
13	13 Other financial or non- financial interests	None	

None.

Please place an "X" next to the following statement to indicate your agreement:

Date: _____Jul. 11th, 2022____

Your Name:_____ Huaxian Mei _____

Manuscript Title: <u>Whole exome sequencing identifies novel inherited genetic variants in tetralogy of Fallot___</u> Manuscript number (if known): __

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7	Support for attending	None	
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	F		
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11	Stock or stock options	None	
12	Receipt of equipment,	None	
	materials, drugs, medical		
	writing, gifts or other services		
42			
13	13 Other financial or non- financial interests	None	

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Date: _____Jul. 11th, 2022____

Your Name:_____ Jing Wu _____

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	materials, drugs, medical		
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42			
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