

Supplemental Table 1. Matrix of genetic similarity among 32 *Stephania rotunda* Lour. accessions using ISSR markers calculated by Jaccard's similarity coefficient

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
1	1.000															
2	0.833	1.000														
3	0.822	0.922	1.000													
4	0.811	0.822	0.789	1.000												
5	0.856	0.844	0.856	0.822	1.000											
6	0.733	0.700	0.667	0.722	0.744	1.000										
7	0.589	0.667	0.656	0.644	0.644	0.811	1.000									
8	0.589	0.600	0.544	0.556	0.556	0.744	0.756	1.000								
9	0.611	0.711	0.656	0.689	0.667	0.833	0.867	0.822	1.000							
10	0.589	0.667	0.633	0.644	0.622	0.744	0.800	0.733	0.867	1.000						
11	0.822	0.856	0.800	0.833	0.789	0.711	0.656	0.611	0.678	0.611	1.000					
12	0.789	0.800	0.767	0.800	0.778	0.678	0.622	0.600	0.644	0.600	0.856	1.000				
13	0.778	0.722	0.711	0.811	0.789	0.711	0.589	0.478	0.589	0.567	0.756	0.744	1.000			
14	0.678	0.733	0.722	0.756	0.756	0.611	0.622	0.556	0.644	0.667	0.789	0.733	0.700	1.000		
15	0.700	0.733	0.789	0.733	0.778	0.700	0.644	0.556	0.667	0.622	0.767	0.733	0.789	0.733	1.000	
16	0.700	0.756	0.722	0.756	0.756	0.700	0.689	0.667	0.733	0.667	0.767	0.733	0.767	0.778	0.822	1.000
17	0.789	0.800	0.833	0.822	0.800	0.722	0.689	0.622	0.711	0.689	0.811	0.756	0.744	0.778	0.822	0.844
18	0.733	0.833	0.756	0.833	0.789	0.711	0.611	0.589	0.722	0.656	0.822	0.744	0.711	0.767	0.744	0.811
19	0.789	0.822	0.767	0.800	0.800	0.678	0.644	0.511	0.622	0.622	0.811	0.800	0.789	0.778	0.733	0.800
20	0.744	0.844	0.767	0.800	0.778	0.744	0.667	0.644	0.778	0.689	0.811	0.756	0.722	0.756	0.778	0.844
21	0.722	0.778	0.767	0.756	0.733	0.700	0.644	0.578	0.667	0.622	0.767	0.733	0.700	0.733	0.733	0.800
22	0.700	0.778	0.722	0.822	0.756	0.678	0.644	0.622	0.733	0.689	0.767	0.733	0.700	0.733	0.733	0.844
23	0.700	0.778	0.767	0.778	0.756	0.700	0.711	0.578	0.711	0.667	0.789	0.756	0.744	0.733	0.822	0.867
24	0.767	0.778	0.744	0.756	0.733	0.744	0.667	0.600	0.711	0.667	0.722	0.689	0.678	0.622	0.689	0.711
25	0.622	0.722	0.667	0.722	0.656	0.644	0.633	0.589	0.700	0.700	0.667	0.722	0.689	0.656	0.700	0.767
26	0.611	0.711	0.656	0.711	0.667	0.633	0.622	0.600	0.689	0.667	0.656	0.644	0.656	0.644	0.689	0.778
27	0.811	0.756	0.722	0.733	0.733	0.722	0.622	0.600	0.689	0.689	0.744	0.711	0.700	0.644	0.689	0.689
28	0.722	0.778	0.744	0.800	0.756	0.656	0.644	0.578	0.667	0.667	0.767	0.733	0.722	0.756	0.778	0.822
29	0.700	0.778	0.722	0.756	0.733	0.656	0.644	0.533	0.667	0.644	0.767	0.733	0.700	0.756	0.733	0.822
30	0.744	0.822	0.767	0.800	0.778	0.700	0.689	0.578	0.711	0.667	0.811	0.756	0.744	0.800	0.778	0.867
31	0.733	0.744	0.756	0.789	0.789	0.667	0.633	0.544	0.656	0.611	0.756	0.744	0.733	0.767	0.789	0.833
32	0.656	0.667	0.656	0.667	0.600	0.567	0.511	0.533	0.622	0.578	0.656	0.600	0.567	0.578	0.644	0.667

Supplemental Table 2. Matrix of genetic similarity among 32 *Stephania rotunda* Lour. accessions using ISSR markers calculated by Jaccard's similarity coefficient

8																
9																
10																
11																
12																
13																
14																
15																
16																
17	1.000															
18	0.903	1.000														
19	0.847	0.889	1.000													
20	0.861	0.931	0.875	1.000												
21	0.847	0.861	0.806	0.847	1.000											
22	0.903	0.861	0.917	0.847	0.861	1.000										
23	0.819	0.917	0.806	0.903	0.833	0.778	1.000									
24	0.778	0.819	0.819	0.861	0.819	0.819	0.792	1.000								
25	0.681	0.750	0.722	0.764	0.750	0.722	0.722	0.792	1.000							
26	0.722	0.764	0.764	0.778	0.792	0.764	0.708	0.806	0.819	1.000						
27	0.764	0.806	0.750	0.792	0.861	0.778	0.750	0.819	0.750	0.819	1.000					
28	0.875	0.917	0.833	0.903	0.889	0.833	0.889	0.847	0.806	0.792	0.833	1.000				
29	0.847	0.889	0.833	0.875	0.861	0.806	0.861	0.847	0.778	0.792	0.806	0.972	1.000			
30	0.875	0.889	0.833	0.875	0.861	0.833	0.833	0.819	0.778	0.819	0.806	0.944	0.944	1.000		
31	0.875	0.861	0.833	0.903	0.833	0.861	0.806	0.875	0.750	0.819	0.833	0.917	0.889	0.889	1.000	
32	0.722	0.708	0.764	0.778	0.736	0.792	0.708	0.778	0.736	0.722	0.764	0.764	0.736	0.708	0.819	1.000

Supplemental Table 3. Matrix of genetic similarity among 32 *Stephania rotunda* Lour. accessions using combined RAPD and ISSR markers calculated by Jaccard's similarity coefficient

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
1	1.000															
2	0.815	1.000														
3	0.821	0.920	1.000													
4	0.802	0.852	0.833	1.000												
5	0.858	0.858	0.852	0.833	1.000											
6	0.691	0.716	0.698	0.716	0.747	1.000										
7	0.636	0.735	0.716	0.722	0.704	0.821	1.000									
8	0.617	0.679	0.636	0.642	0.636	0.802	0.809	1.000								
9	0.630	0.741	0.710	0.716	0.698	0.852	0.883	0.864	1.000							
10	0.623	0.710	0.679	0.698	0.679	0.784	0.852	0.809	0.883	1.000						

11	0.802	0.827	0.784	0.815	0.809	0.679	0.673	0.630	0.667	0.648	1.000			
12	0.778	0.778	0.759	0.790	0.772	0.642	0.623	0.605	0.617	0.623	0.877	1.000		
13	0.772	0.784	0.753	0.846	0.815	0.722	0.691	0.586	0.660	0.667	0.772	0.747	1.000	
14	0.716	0.778	0.772	0.790	0.809	0.667	0.698	0.654	0.691	0.722	0.790	0.753	0.772	1.000
15	0.735	0.784	0.827	0.772	0.802	0.698	0.716	0.636	0.698	0.679	0.772	0.735	0.790	0.784
16	0.747	0.809	0.802	0.821	0.827	0.722	0.728	0.698	0.747	0.704	0.796	0.747	0.802	0.833
17	0.790	0.827	0.870	0.852	0.821	0.741	0.735	0.691	0.753	0.722	0.802	0.741	0.772	0.790
18	0.741	0.840	0.809	0.864	0.784	0.728	0.698	0.679	0.753	0.698	0.778	0.728	0.772	0.802
19	0.784	0.833	0.802	0.809	0.802	0.722	0.716	0.623	0.698	0.679	0.759	0.747	0.815	0.784
20	0.765	0.877	0.833	0.827	0.796	0.741	0.735	0.704	0.778	0.722	0.790	0.741	0.772	0.802
21	0.759	0.796	0.790	0.784	0.790	0.710	0.691	0.673	0.710	0.679	0.759	0.722	0.741	0.796
22	0.735	0.821	0.790	0.833	0.790	0.722	0.716	0.710	0.784	0.741	0.747	0.710	0.741	0.759
23	0.722	0.796	0.790	0.809	0.790	0.710	0.753	0.660	0.722	0.704	0.784	0.747	0.778	0.809
24	0.728	0.802	0.772	0.765	0.735	0.765	0.722	0.691	0.741	0.722	0.679	0.654	0.710	0.691
25	0.617	0.741	0.685	0.716	0.660	0.679	0.698	0.679	0.716	0.722	0.654	0.679	0.698	0.679
26	0.654	0.728	0.710	0.716	0.673	0.667	0.673	0.679	0.704	0.698	0.654	0.654	0.673	0.704
27	0.809	0.784	0.741	0.759	0.753	0.722	0.667	0.685	0.710	0.716	0.722	0.710	0.741	0.710
28	0.735	0.833	0.802	0.846	0.802	0.710	0.741	0.698	0.735	0.728	0.759	0.722	0.790	0.821
29	0.710	0.821	0.778	0.809	0.778	0.710	0.728	0.660	0.722	0.704	0.747	0.722	0.765	0.809
30	0.747	0.846	0.815	0.833	0.802	0.722	0.753	0.685	0.747	0.716	0.772	0.735	0.778	0.833
31	0.741	0.815	0.821	0.815	0.796	0.716	0.722	0.667	0.728	0.698	0.741	0.716	0.772	0.802
32	0.679	0.728	0.698	0.679	0.673	0.630	0.611	0.630	0.679	0.648	0.679	0.617	0.648	0.642

