Table S4 Gene flow among all populations of *P.serrulata* and *P.xueluoensis*

	QLFxD	QLFxU	LSxD	LSxU	MYSxD	MYSxU	DWSx	Esx	QLFs	LuSs	MYSs	LaoSs
QLFxU	3.647											
LSxD	2.560	2.804										
LSxU	2.359	2.403	4.599									
MYSxD	1.659	1.429	1.864	1.290								
MYSxU	1.273	1.232	1.202	1.095	2.928							
DWSx	1.874	1.536	1.846	1.310	1.668	1.931						
Esx	1.951	1.693	1.850	1.338	1.078	1.108	2.941					
QLFs	1.472	1.283	1.648	1.227	1.571	0.814	1.879	1.129				
LuSs	1.233	1.111	1.163	0.905	1.461	0.677	1.447	1.166	4.133			
MYSs	1.070	0.903	1.132	0.826	1.149	0.603	1.455	1.285	2.123	1.932		
LaoSs	0.784	0.539	0.433	0.570	0.404	0.871	0.495	0.834	1.671	1.677	1.081	
FHSs	0.590	0.921	0.980	0.755	0.909	0.614	0.698	0.689	1.261	0.980	0.789	1.440