

Additional file 4: Mitochondrial genetic distances (*D_{xy}*) between and within Eastern tree frog taxa

Between taxa

	<i>H. immaculata</i> / <i>H. suweonensis</i>	<i>H. japonica</i> A	<i>H. japonica</i> B
<i>H. immaculata</i> / <i>H. suweonensis</i>	-		
<i>H. japonica</i> A	0.135	-	
<i>H. japonica</i> B	0.128	0.104	-

Within *H. japonica* taxon A

	A1	A2	A3	A4
A1	-			
A2	0.036	-		
A3	0.066	0.055	-	
A4	0.070	0.064	0.036	-

Within *H. japonica* taxon B

	B1	B2	B3	B4	B5
B1	-				
B2	0.036	-			
B3	0.039	0.021	-		
B4	0.041	0.035	0.034	-	
B5	0.042	0.036	0.036	0.017	-
