

Table S7. Uncorrected genetic distances (p-distances) within and between *Pareremias* and *E. nikolskii* based on 13 concatenated PCGs.

	1	2	3	4	5	6	7	8	9	10	11
1. <i>E. nikolskii</i>											
2. Viviparous group	0.108										
3. <i>E. argus</i> - <i>E. brenchleyi</i>	0.123	0.123									
4. <i>E. argus</i>	0.122	–	–	–							
5. <i>E. brenchleyi</i>	0.124	–	–	–							
6. <i>E. szczerbaki</i>	0.107	–	–	–	–						
7. <i>E. yarkamdensis</i>	0.105	–	–	–	–	0.087					
8. <i>E. dzungarica</i>	0.106	–	–	–	–	0.088	0.039				
9. <i>E. przewalskii</i>	0.109	–	–	–	–	0.091	0.043	0.051			
10. <i>E. stummeri</i>	0.109	–	–	–	–	0.067	0.084	0.087	0.089		
11. <i>E. multiocellata</i>	0.109	–	–	–	–	0.092	0.079	0.087	0.084	0.091	