

**Table S8.** Genome-wide pairwise sequence divergence, estimated using equation E8.2 (see Text S8) using all the genomic sites that passed the genomic quality filters outlined in Text S8.

	<b>Boxer</b>	<b>Basenji</b>	<b>Dingo</b>	<b>Israeli wolf</b>	<b>Croatian wolf</b>	<b>Chinese wolf</b>
<b>Boxer</b>						
<b>Basenji</b>	0.00087					
<b>Dingo</b>	0.00094	0.00100				
<b>Israeli wolf</b>	0.00111	0.00112	0.00116			
<b>Croatian wolf</b>	0.00113	0.00117	0.00116	0.00115		
<b>Chinese wolf</b>	0.00114	0.00117	0.00115	0.00118	0.00115	
<b>Golden jackal</b>	0.00211	0.00214	0.00214	0.00214	0.00214	0.00214