

**Table S7.** Genome-wide pairwise sequence divergence, estimated using equation E8.1 (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	0.00097				
<b>Israeli wolf</b>	0.00111	0.00105	0.00111			
<b>Croatian wolf</b>	0.00113	0.00110	0.00112	0.00101		
<b>Chinese wolf</b>	0.00114	0.00111	0.00111	0.00106	0.00105	
<b>Golden jackal</b>	0.00211	0.00211	0.00212	0.00209	0.00209	0.00210