

Table S1: Genomic regions associated with Red Blood Cell Number (RBC)

<i>SNP</i>	<i>Chr</i>	<i>Position (bp)</i>	<i>Best fitting model</i>	<i>Nominal P-value</i>	<i>Effect Size</i>	<i>Other Significant Phenotypes</i>	<i>Genes within 100 kb on either side</i>
OAR15_7970400	15	8,237,224	allelic	4.2×10^{-6}	0.641	None	None