Table S4. Dock7 CRISPR primer pairs. Primer pairs were used to analyze the Dock7 gene by genotyping (G), cloning (C), and sequencing (S).

The cKO1 Δ primer pairs were used to assess deletion of exons 3 and 4 in the Dock7 gene. The cKO2 Δ primer pairs used to assess deletion of exons 3 through 7 in the Dock7 gene. The amplification of isolated DNA was performed according to the Jumpstart (J), Terra Taq (T), or Phusion polymerase (P) protocols described in Table S2. Forward (F) and reverse (R) primers are abbreviated accordingly.