

SUPP Table 2: Assembly statistics for the 26 genes sequenced through target-enrichment – “depth of coverage” represents the number of reads that included in the reconstructed sequence against the mallard reference; “gene covered” refers to the total amount of the mallard reference that included any amount of coverage.

	<i>Speckled teal</i>		<i>Yellow-billed pintail</i>		<i>Cinnamon teal</i>		<i>All Three</i>	
<i>Gene Name</i>	<i>Depth of Coverage</i>	<i>Gene Covered</i>	<i>Depth of Coverage</i>	<i>Gene Covered</i>	<i>Depth of Coverage</i>	<i>Gene Covered</i>	<i>Depth of Coverage</i>	<i>Gene Covered</i>
<i>ACE</i>	147	97.70%	157	99.20%	165	99.20%	156.33	98.70%
<i>ANGPT1</i>	456	95.50%	463	96.10%	492	94.60%	470.33	95.40%
<i>ANGPT2</i>	514	99.40%	518	99.60%	570	99.60%	534.00	99.53%
<i>ARNT</i>	411	99.50%	428	99.80%	437	99.80%	425.33	99.70%
<i>ARNT2</i>	463	99.80%	478	99.90%	520	99.90%	487.00	99.87%
<i>ARNTL</i>	558	91.70%	567	92.10%	601	90.90%	575.33	91.57%
<i>CLOCK</i>	463	99.80%	471	100.00%	497	100.00%	477.00	99.93%
<i>CUL2</i>	589	99.90%	592	100.00%	572	98.00%	584.33	99.30%
<i>EGLN1</i>	388	99.30%	404	99.70%	398	99.60%	396.67	99.53%
<i>EGLN3</i>	421	99.30%	436	99.60%	460	99.70%	439.00	99.53%
<i>HIF2A</i>	456	99.70%	457	100.00%	485	100.00%	466.00	99.90%
<i>HIF1A</i>	512	99.90%	523	100.00%	546	100.00%	527.00	99.97%
<i>HSP90AA1</i>	476	99.90%	474	100.00%	405	100.00%	451.67	99.97%
<i>MTOR</i>	476	99.80%	485	100.00%	510	100.00%	490.33	99.93%
<i>NOS1</i>	140	95.20%	151	96.60%	152	96.10%	147.67	95.97%
<i>NOS2</i>	540	100.00%	549	100.00%	602	99.90%	563.67	99.97%
<i>P4HA1</i>	406	98.00%	413	98.50%	455	98.40%	424.67	98.30%
<i>P4HA2</i>	509	99.70%	514	99.90%	569	99.80%	530.67	99.80%
<i>P4HA3</i>	72	79.80%	78	81.80%	72	81.50%	74.00	81.03%
<i>PDHA1</i>	507	100.00%	506	99.90%	500	99.60%	504.33	99.83%
<i>PPARA</i>	532	98.40%	557	98.60%	596	97.70%	561.67	98.23%
<i>PPARD</i>	119	98.20%	132	100.00%	121	99.30%	124.00	99.17%
<i>PPARG</i>	516	99.70%	523	99.70%	565	99.80%	534.67	99.73%
<i>TCEB3</i>	127	95.70%	143	97.80%	128	96.60%	132.67	96.70%
<i>THRB</i>	430	99.50%	439	99.90%	492	99.80%	453.67	99.73%
<i>VEGF</i>	219	99.00%	229	99.80%	234	99.80%	227.33	99.53%