

Table S1. Summary statistics of clean transcriptome sequencing data

Samples	Clean reads	Clean bases	GC Content	%≥Q30
hybrid tilapia (NA)	20 851 305	6 218 510 436	49.15%	90.20%
<i>O.niloticus</i> (N)	21 076 229	6 302 527 684	50.02%	90.29%
<i>O.aureus</i> (A)	23 503 357	7 020 101 902	49.83%	90.42%