Figure S3. Alignment shows that the degenerate *ctnna3* forward (a3-F) and reverse (a3-R) primers have no mismatches with chicken *ctnna3* genomic sequence. Both primers have one mismatch (yellow) with the corresponding region of chicken *ctnna1* and two mismatches with chicken *ctnna2*, suggesting that the primers preferentially amplify *ctnna3,* and *ctnna1* with a higher efficiency than *ctnna2* from chicken genomic DNA.