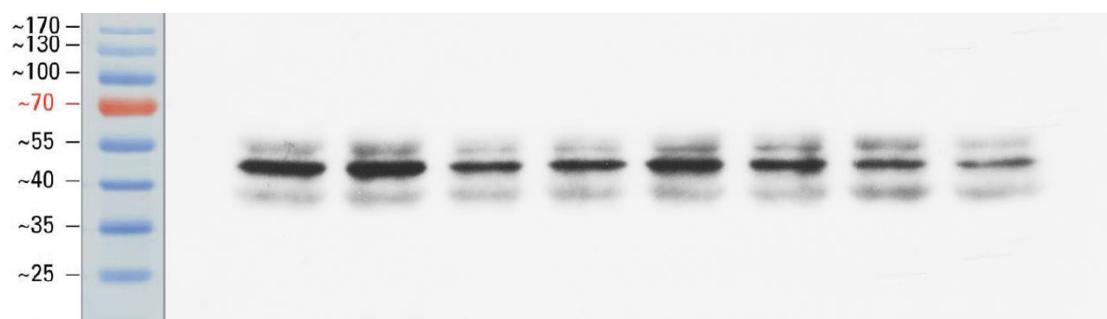


**A GenBank: XP\_005463932.1**

MEDLDALLADLESTSSPLPRCPVLLTSDPP**QNSDPTTQDSAQTRPPP**PAYTPQQ  
TVSSAMKTTQNSTPDKLYSTVCKPRSPRSADPPLAFSSSLGGGLSELDHLLQ  
ELNATQFNITDEILAQFPSSKKDERDNIKDKAPTSSSSAKPSATSATLELDKLM  
ASLSDFRVQSTPAAPVTPAPVTASPQQPAAAPPQPPSSGGSLDSMLGLLQSDL  
RQGVQTSSKGNCACQKPVVGQVVTALGKVWHPEHFVCTECETELGSRNFF  
EKDGRPYCEPDYFTLFSPHCAHCNKPILNKMTALDKNWHPQCFCCVKCSRA  
FGEEGFHDREGQQYCQQCFLTLFASRCQGCSQPILENYISALNSLWHPQCFVC  
RECYSFVNGSFFEHEGKPLCEAHYHQSRGSMCQACQQPILGRCVTAMGAKF  
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**TGF $\beta$ 1I1 protein sequence: QNSDPTTQDSAQTRP**

**B**



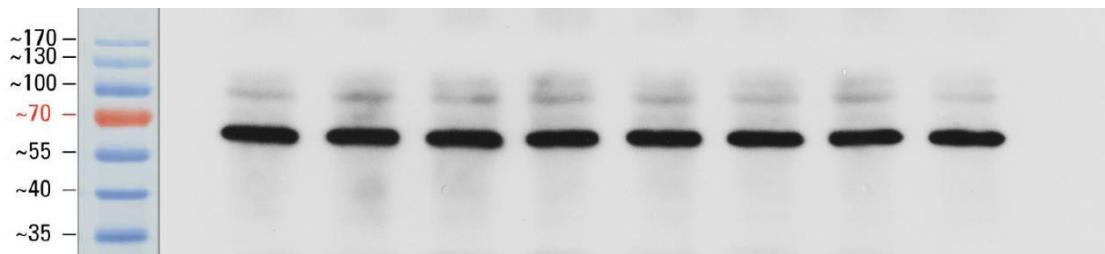
**Figure S1 TGF $\beta$ 1I1 protein sequence information and marker map of Nile tilapia *Oreochromis niloticus*.**

**A GenBank: AHN82370.1**

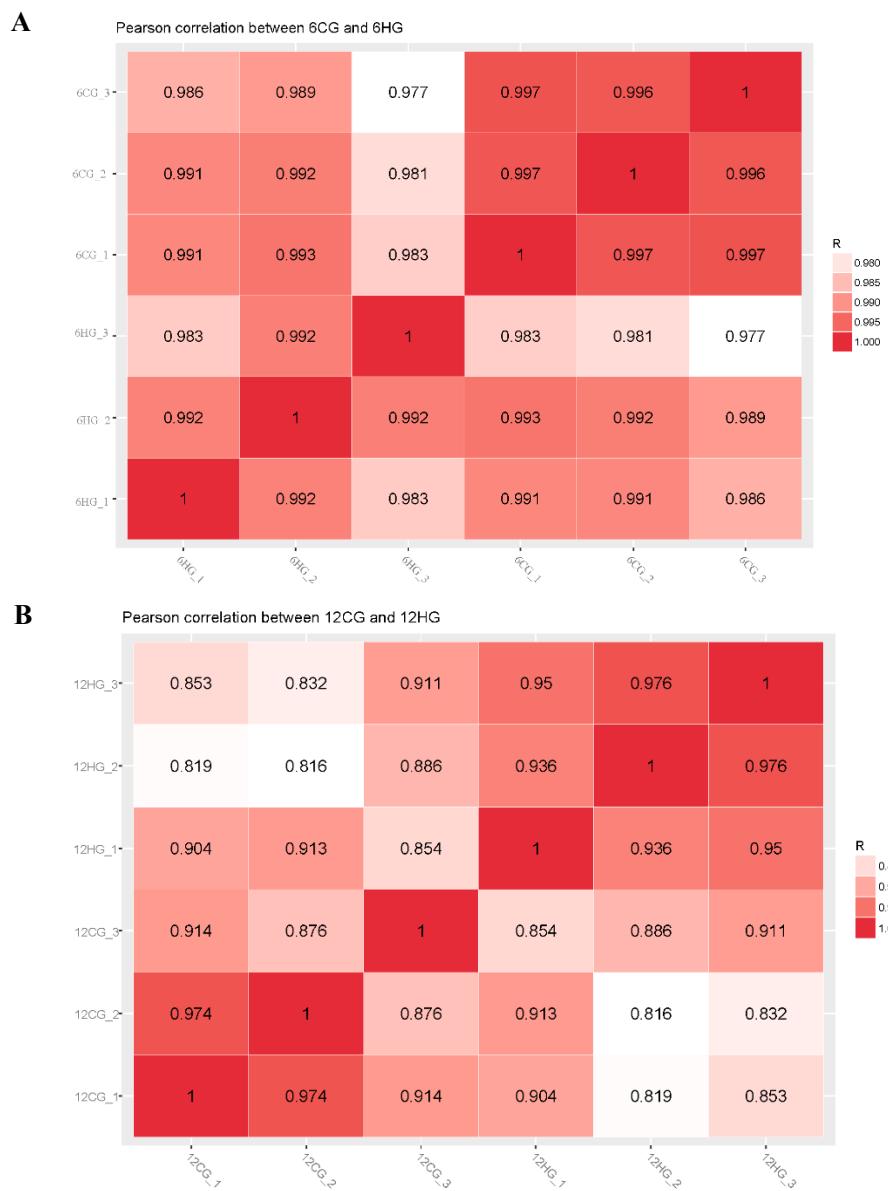
IAALVVDNGSGMCKAGFAGDDAPRAVFPSIVGRPRHQGVMVGMGQKDSYV  
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THTVPIYEGYALPHAILRLDLAGRDLTDYLMKILTERGYSFTTAEREIVRDIK  
EKLCYVALDFEQEMGTAASSSSLEKSYELPDGQVITIGNERFRCPEALSQPSFL  
GMESCGIHETTYNSIMKCDVDIRKDLYANTVLSGGTTMYPGIADRMQKEITA  
LAPSTMKIKIIAPPERKYSVWIGGSILASLSTFQQMWISKQEYDESGPSIVHRKF

**$\beta$ -actin protein sequence: ISKQEYDESGPSIV**

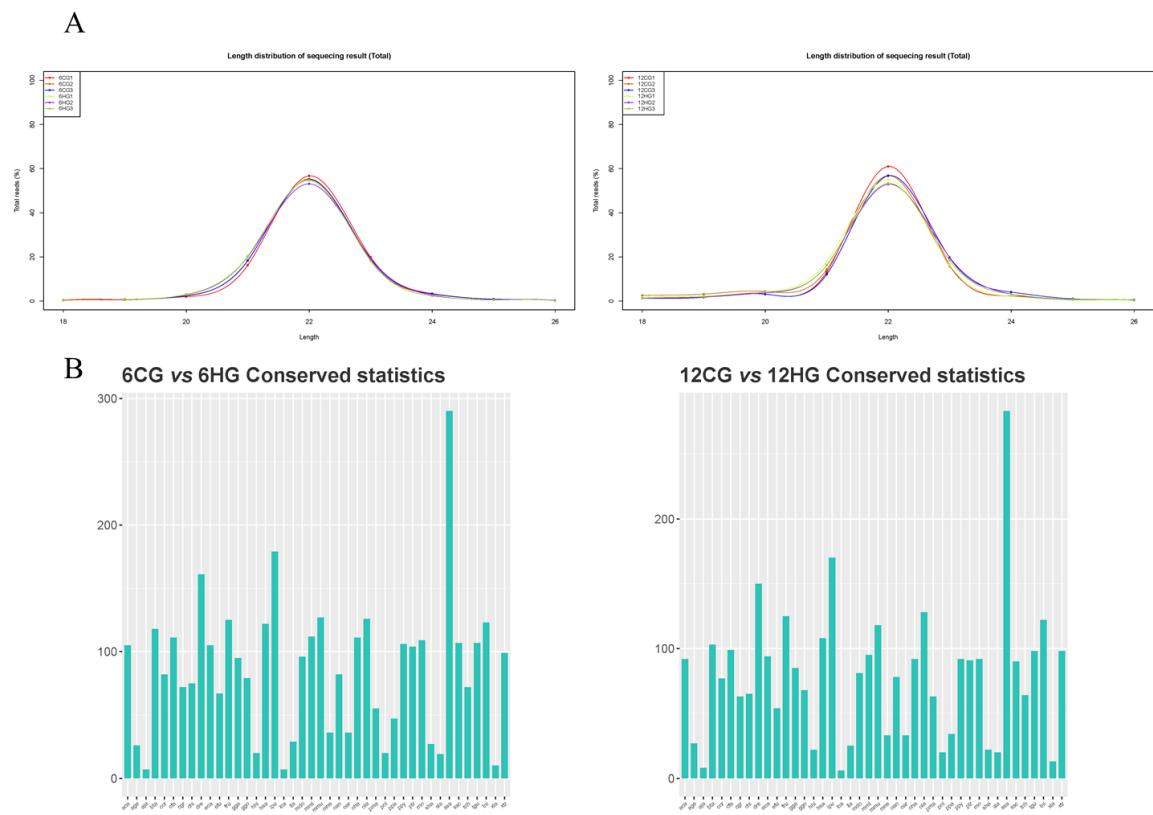
**B**



**Figure S2**  $\beta$ -actin protein sequence information and marker map of Nile tilapia *Oreochromis niloticus*.



**Figure S3 Quality assessment of miRNA sequencing results in 6CGs vs 6HGs and 12CGs vs 12HGs**



**Figure S4 Length distribution of counts of total and unique sRNAs (A) and conservation profile of the identified miRNAs (B)**