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Conservation of structural and interactional features of CD28 and CD80/86 molecules from Nile tilapia (*Oreochromis niloticus*)

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## ACCEPTED MANUSCRIPT

1	Conservation of Structural and Interactional Features of CD28 and CD80/86
2	Molecules from Nile Tilapia (Oreochromis niloticus)
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16	
17	Abstract
18	Interaction of CD28 with CD80 or CD86 molecules provides a costimulatory
19	signals required in T cell activation. In this study, we cloned and analyzed a CD28
20	gene (On-CD28) and a CD80/86 gene (On-CD80/86) from Nile tilapia (Oreochromis
21	niloticus). Sequence analysis revealed the typical characteristics of On-CD28 protein;
22	for instance, the proline-based motif ( <sup>117</sup> TYPPPL <sup>122</sup> ) is essential in binding of CD28
23	to CD80/86 ligands. Moreover, an extracellular Ig domain was found in On-CD80/86;
24	this domain is responsible in binding of CD28 to CD80/86 receptors. Subcellular
25	localization analysis showed that both On-CD28 and On-CD80/86 were distributed

predominantly in the cytomembrane. Yeast two-hybrid assay showed that On-CD28

directly interacted with On-CD80/86. On-CD28 and On-CD80/86 transcripts were

detected in all the examined tissues of healthy Nile tilapia, and the highest expression

29 levels of On-CD28 and On-CD80/86 were detected in the brain and heart, respectively.

30 Following a bacterial challenge using *Streptococcus agalactiae* in vivo, On-CD28 and

31 On-CD80/86 were upregulated in head kidney, spleen, intestines, and brain. However,

they showed different expression profiles in response to stimulation with inactivated *S*.

33 *agalactiae* in vitro. These findings indicated that the interaction of On-CD28 with

34 On-CD80/86 provides a costimulatory signals that possibly play an important role in

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