**Comparative genome analyses reveal distinct structure in the saltwater crocodile MHC**

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**Figure S3. Syntenic areas (red) between saltwater crocodile TAP2 (MHC gene cluster 3.2 or scaffold-7467 described in Table 1) and jungle fowl TAP2 (ID 2731, v2, unmasked and CDS).** Gray arrows indicate gene models; green arrows indicate protein coding sequences (CDS); blue arrows (on top of gray genes) indicate mRNA; and yellow arrows indicate approximately 50% GC content in codon wobble positions. Areas highlighted in orange show unsequenced positions