

**Table S4 Presence of putative CREA consensus binding sequence 5'-(C/G)YGGRG-3' in the 5' region of *A. flavus* genes assessed on this study. Positions relative to start codon of 5'-(C/G)YGGRG-3' were obtained by scanning 1kb upstream of the proposed start codon. Genes were from <http://www.ncbi.nlm.nih.gov/> and <http://www.aspgd.org>**

<b>Gene (accession number)</b>	<b>Position relative to start codon (5'-GCGGAG-3')</b>	<b>Position relative to start codon (5'-CTGGGG-3')</b>
<i>farA</i> (AFL2G_05109)	No motif found	-241
<i>farB</i> (AFLA_012010)	No motif found	No motif found
<i>pexK</i> (AFLA_036410)	-231	-985
<i>echA</i> (AFLA_043610)	No motif found	No motif found
<i>foxA</i> (AFLA_041590)	No motif found	-361
<i>acuJ</i> (AFLA_135240)	-298	-344
<i>eciA</i> (AFLA_055840)	No motif found	-982
<i>derA</i> (AFLA_061230)	-119, -229	No motif found
<i>afIR</i> (AFLA_139360)	No motif found	No motif found
<i>afID</i> (AFLA_139390)	No motif found	No motif found