

1 Tectal corticotropin-releasing factor (CRF) neurons respond to fasting and a reactive stressor in  
2 the African Clawed Frog, *Xenopus laevis*

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28      **Abstract**

29              It is well established that hypothalamic neurons producing the peptide corticotropin-  
30      releasing factor (CRF) play a key role in stress adaptation, including reduction of food intake  
31      when a threat or stressor is present. We have previously reported on the presence of an intrinsic  
32      CRF signaling system within the optic tectum (OT), a brain area that plays a key role in visually  
33      guided prey capture/predator avoidance decisions. To better understand the potential role of  
34      tectal CRF neurons in regulating adaptive behavior and energy balance during stress we  
35      examined evidence for modulation of tectal CRF neuronal activity after stressor exposure and  
36      food deprivation in the African clawed frog *Xenopus laevis*. We tested two predictions, 1) that  
37      exposure to categorically distinct stressors (ether vapors, a reactive stressor; and shaking, an  
38      anticipatory stressor) will reduce food intake and modulate the activity of tectal CRF cells, and  
39      2) that food deprivation will modulate the activity of tectal CRF cells. Exposure to ether  
40      increased tectal content of CRF and CRF transcript, but lowered CRFR1 transcript abundance.  
41      Two weeks of food deprivation reduced total fat stores in frogs and decreased tectal content of  
42      CRF content while having no effect on CRF and CRFR1 transcript abundance. Our data are  
43      consistent with a role for tectal CRF neurons in modulating food intake in response to certain  
44      stressors.

45      Keywords: Amphibian; stress; vision; fear; anxiety; foraging.

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52        **1. Introduction**

53            Corticotropin releasing factor (CRF) is a 41 amino acid peptide discovered in the search  
54        for a hypothalamic peptide that regulates corticotropin secretion from the pituitary gland (Spiess  
55        et al., 1981; Vale et al., 1981; Bale, 2014). Since its original discovery, roles for CRF in  
56        regulating processes as diverse as stress (Bale and Vale, 2004; Ulrich-Lai and Herman, 2009;  
57        Kormos and Gaszner, 2013; Bale, 2014; Carr and Lovejoy, 2015), depression (Waters et al.,  
58        2015), appetite (Stengel and Tache, 2014), synapse formation (Liao et al., 2014), integumental  
59        function (Slominski et al., 2013) and cryoprotection (Boorse et al., 2006) all have been studied.  
60        The best known role of CRF seems coordinating a myriad of physiological and behavioral  
61        responses during stress (Beurel and Nemeroff, 2014; Backström and Winberg, 2013). In fact,  
62        hypophysiotropic neurons in the paraventricular nucleus (PVN) of the hypothalamus contain a  
63        large population of CRF neurons in mammals (Makara et al., 1981; Bruhn et al., 1984). Whether  
64        this population of CRF cells contributes to every central nervous system (CNS) role attributed to  
65        CRF is unlikely, as neuroanatomical studies have shown across vertebrate species that CRF is  
66        produced in many other neuronal populations within the CNS (Merchanthaler et al., 1982; Vigh  
67        et al., 1982; Yao et al., 2004; Calle et al., 2005; Carr et al., 2010).

68            It is becoming increasingly clear that local brain circuits regulated by CRF, and other  
69        members of the CRF peptide family (urocortin 1, UCN1; urocortin 2, UCN2; and urocortin 3,  
70        UCN3), may mediate many adaptive behavioral and physiological responses (Ronan and  
71        Summers, 2011; Henckens et al. 2016). For example, the endogenous ligand for CRF receptor 2  
72        (CRF R2) appears to modulate adaptive features of social behavior in mice by acting on local  
73        circuitry in the medial amygdala (Shemesh et al., 2016). Hippocampal interneurons producing  
74        CRF play a role in mediating the adverse consequences of early life stress on learning and

75 memory (Chen et al., 2012). Knocking down CRF expression in the central nucleus of the  
76 amygdala does not influence anxiety but does impact grooming behavior and, in part, the  
77 endocrine response to a behavioral stressor (Callahan et al., 2013).

78 In amphibians one of the largest populations of CRF neurons occurs in the optic tectum  
79 (OT; Bhargava and Rao, 1993; Yao et al., 2004; Calle et al., 2005; Carr et al., 2010; Carr et al.,  
80 2013; Carr, 2015; Harris and Carr, 2016), a brain area involved in processing visual (Ewert et al.,  
81 1981; Vanegas, 1984) and lateral line sensory information (Hiramato and Cline, 2009) and  
82 integrating behavioral and endocrine responses (Carr, 2015; Liu et al., 2016). In particular, this  
83 brain area, anatomically and functionally homologous to the superior colliculus in mammals, is  
84 important in coordinating approach and avoidance behaviors and in adaptive responses to visual  
85 threats such as predators (Ewert et al., 1991; Vanegas, 1984) and it is likely that sensorimotor  
86 integration in the OT is coordinated and modulated by numerous peptides. Indeed, several  
87 bioactive peptides, in addition to CRF, are produced by neurons and retinal ganglion cells  
88 innervating the OT (e.g., somatostatin, Vandesande and Dierickx, 1980; substance P, Inagaki et  
89 al., 1981, Kuljis and Karten, 1982; leucine-enkephalin, cholecystokinin octapeptide, bombesin,  
90 avian pancreatic polypeptide, Kuljis and Karten, 1982; neuropeptide Y, Danger et al., 1985;  
91 GnRH, Jokura and Urano, 1986; TRH, Zoeller and Conway, 1989). However, there two lines of  
92 evidence making tectal CRF cells particularly interesting with respect to their potential for  
93 modulating the recognition and response to moving prey items. Firstly, tectal CRF-  
94 immunoreactive fibers and cells bodies. Tectal layer 9, which receives the majority of retinal  
95 afferents from retinal ganglion cells (Ten Donkelaar, 1998) is moderately to densely innervated  
96 by CRF fibers (Carr et al., 2010), and CRF producing cells located in tectal layers 6 and 8 (Carr  
97 et al., 2010), may play a critical role in sensorimotor integration (Baginskas and Kuras, 2008).

98 Secondly, CRF administration alters visually guided prey capture across anuran genera (Carr et  
99 al., 2002; Crespi and Denver, 2004; Morimoto et al., 2011).

100 The physiological role of tectal CRF neurons is unknown at present. It is unknown if tectal  
101 CRF neurons react to changes in food intake (Calle et al., 2006), and like CRF neurons in the  
102 PVN, to stressors (Bruhn et al., 1984; Chappell et al., 1986; Imaki et al., 1991). To address these  
103 issues we examined CRF content (by homologous radioimmunoassay (RIA) and mRNA  
104 abundance (by quantitative real-time PCR, qRT-PCR) in the OT in response to stressors and food  
105 deprivation. We hypothesized that if CRF plays a role in sensorimotor integration in response to  
106 a threat, stressor exposure should reveal evidence for modulation of tectal CRF cells. We also  
107 hypothesized if tectal CRF suppresses food intake then starvation should negatively modulate  
108 tectal CRF cell activity.

109 **2. Methods**

110 *2.1. Animals and care*

111 Juvenile (5-12 g sexually immature, for behavior and radioimmunoassay studies) and  
112 sub-adult (28-62 g males, for qRT-PCR studies) South African Clawed frogs (*Xenopus laevis*)  
113 were obtained from our in-house colony and were reared in deionized water containing 0.33 g/L  
114 Instant Ocean (Instant Ocean, Blacksburg, VA, USA) in a 300 L tank (178 cm L x 46 cm W x 51  
115 cm D) at a maximum stocking density of 30 frogs per tank. Frogs were maintained at a  
116 temperature of 22– 23° C on a 12L:12D light regimen. Juvenile frogs were fed Nasco (Ft.  
117 Atkinson, WI, USA) floating pellets and sub-adults fed Nasco sinking pellets three times per  
118 week. The water was changed and tanks were cleaned three times per week. All procedures were  
119 approved by the Texas Tech Animal Care and Use Committee.

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121 2.2. *Stressor effects on food intake.*

122 Frogs (n=32,  $7.90 \pm 0.26$  g) were exposed to a shaking stressor for 4 h (Yao and Denver,  
123 2004), ether vapors for 1 min (Olsen et al., 1999) or no treatment. Immediately after the shaking  
124 stressor or 60 min after ether vapor exposure frogs were placed into individual tanks (15 cm L X  
125 12 cm W X 13 cm D) filled with 0.5 L deionized water and 0.15 g Instant Ocean 24 h before  
126 food was added. Control animals were tested at the same time of day to eliminate any diurnal  
127 rhythm influence. For measurement of food intake, 1.2 g of chicken liver (Pilgrim's Pride  
128 Corporation, Greenly, CO) was dropped into the tank and, after 60 min, the remaining liver was  
129 weighed and food intake calculated as a percentage of body mass. Animals were returned to  
130 their home tank after the assay.

131 2.4. *Stressor effects on brain CRF content and transcript abundance.*

132 To assess changes in brain CRF content a total of 18 juvenile frogs (6 per group, 5-12 g)  
133 were subjected to the shaking or ether stressor as described above, while another group served as  
134 controls. Immediately after the shaking stressor and 60 min after ether vapor exposure frogs were  
135 euthanized for tissue collection.

136 For qRT-PCR analysis, one group ( $47.0 \pm 5.04$  g) of eight male frogs was exposed to ether  
137 vapors as described above while a second group of males ( $44.8 \pm 5.17$  g) served as controls.  
138 Immediately after ether vapor exposure frogs were euthanized for tissue collection.

139 2.3. *Food deprivation effects on brain CRF content and transcript abundance*

140 It was necessary to conduct two food deprivation experiments, one to collect tissue for  
141 RIA analysis (juvenile frogs, n = 16) and another to collect tissue for qRT-PCR analysis (subadult  
142 frogs, n = 14, to increase RNA yield). For each experiment all 16 or 14 animals were ranked by  
143 body mass immediately before group assignment. Frogs were then alternatively assigned to one

144 of the two groups so that starting bodyweight was exactly the same between groups. Body mass  
145 was recorded again after 1 wk and 2 wk. All frogs were housed in individual 8 L tanks filled with  
146 4 L of deionized water until they were euthanized.

147 We measured body composition (fat mass, lean mass, total body water) before and after  
148 the 2 wk food deprivation study using a quantitative magnetic resonance body composition  
149 analyzer (EchoMRI, Houston, TX). Although the use quantitative magnetic resonance for body  
150 composition analysis has been most widely documented in mammals (McGuire and Guglielmo,  
151 2010), recent studies have shown the utility of this approach for measuring body composition in  
152 poikilotherms (Fowler et al., 2016; Warner et al., 2016). The instrument was calibrated using  
153 both large (485 g) and small (15.5 g) canola oil standards and body mass ( $\pm 0.01$  g) was recorded  
154 prior to each measurement.

155 *2.5. Tissue Collection*

156 Frogs were euthanized with 0.1% MS222 buffered with equal parts NaHCO<sub>3</sub>, decapitated,  
157 and brain areas (telencephalon, Tel; hypothalamus/thalamus, H/T; optic tectum, OT; brainstem,  
158 BS) dissected in ice cold Earl's Balanced Salt Solution. For RNA extraction, brain areas were  
159 frozen in 10 vol RNALater (Thermofisher) in RNase free 1.5 mL microcentrifuge tubes and  
160 stored at -80° C. CRF was extracted from brain areas for RIA as described previously (Carr et  
161 al., 2010). Protein content was determined using a modification (Markwell et al., 1981) of the  
162 Lowry method (Lowry et al., 1951) for Tel and BS regions, and using the micro BSA protein  
163 assay kit (Thermo Scientific) for H/T and OT.

164 *2.6. xCRF iodination and RIA*

165 xCRF (courtesy of Dr. R. Denver, Univ. Michigan) was iodinated and purified as  
166 previously described (Carr et al., 2010). Brain extracts were reconstituted in CRF assay buffer

167 (0.1M sodium phosphate, 0.05M sodium chloride, 0.01% (w/v) sodium azide, 0.1% (w/v) BSA,  
168 and 0.1% (v/v) Triton X-100) prior to RIA. Briefly, reconstituted media samples were incubated  
169 with rabbit anti-xCRF (1:6000, courtesy of Dr. R. Denver, Univ. Michigan) overnight before  
170 addition of 10,000 cpm  $^{125}\text{xCRF}$  in a final assay volume of 400 $\mu\text{l}$ . After 48 h incubation at 4°C,  
171 bound hormone was separated by addition of goat anti-rabbit IgG (Sigma-Aldrich) in 5%  
172 polyethylene glycol and after a 30 min incubation at 4°C centrifugation at 2,500 x g. xCRF  
173 content was determined by comparison with authentic xCRF standards run in duplicate in the  
174 same RIA.

175 *2.7 RNA Extraction*

176 Tissues were removed from RNALater and RNA was extracted using the RNAqueous-  
177 4PCR DNA-free <sup>TM</sup> RNA Isolation for RT-PCR kits (Thermofisher), which includes a DNase  
178 step, according to the manufacturer's instructions and treated with murine RNase inhibitor (1  
179 unit/ $\mu\text{l}$ , New England Biolabs) prior to freezing at -80° C. RNA quality and concentration were  
180 determined using an Experion<sup>TM</sup> Automated Electrophoresis System (Bio-Rad Laboratories,  
181 Inc.) and Experion<sup>TM</sup> RNA StdSens Reagents and RNA StdSens Chips (Bio-Rad Laboratories,  
182 Inc.). RNA samples with an RNA quality indicator (RQI)  $\geq 7$  were then reverse transcribed to  
183 cDNA using a High Capacity cDNA Reverse Transcription Kit (Applied Biosystems). RNA  
184 samples with RQI  $< 7$  were eliminated from analysis. The thermal cycler was set at 25°C for 10  
185 min, then 37°C for 120 min, and was completed at 85°C for 5 min. A control RNA group treated  
186 with the High Capacity cDNA Reverse Transcription Kit but not reverse transcriptase was  
187 checked by qRT-PCR for a 10-fold difference. cDNA was stored at -20°C.

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190 2.8. *qRT-PCR*

191 For qRT-PCR assays, primer and template DNA concentrations were determined using a  
192 Nanodrop spectrophotometer. cDNA was then diluted to 200 ng/µl with DNase-free water.  
193 Primers for the *X. laevis* CRF and CRFR1 (Boorse and Denver, 2006), and *rpl8* (housekeeping  
194 gene, Carr et al., 2008) were used (GenBank Accession Numbers S50096, Y14036 and U00920,  
195 respectively). The identity of PCR products was confirmed using gel electrophoresis after RT-  
196 PCR and sequencing (Macrogen, Rockville, MD). qRT-PCR reactions were performed in  
197 duplicate in 96-well optical reaction plates (Applied Biosystems, Grand Island, NY) in a total  
198 volume of 25 µl containing 1 µl diluted cDNA template, 1 µl each of forward and reverse primer  
199 (final concentration of 200 nM each), 12.5 µl of SYBR green PCR master mix (Applied  
200 Biosystems), and nuclease free water. Non-template controls contained nuclease free water  
201 substituted for the cDNA template. Plates were centrifuged before loading onto an ABI Prism  
202 7000 detector. Efficiency was determined for amplification of primer pairs using 10-fold serial  
203 dilutions of template in duplicate and calculating the slope of the regression plotting Ct value  
204 against the log of the template amount. A standard curve was made from serial dilutions (300,  
205 30, 3, 0.3, and 0.03 ng) to determine primer amplification efficiency. Cycle threshold values  
206 were normalized compared to the reference gene *rpl8* and expressed as a percentage of the  
207 control values using the  $\Delta\Delta Ct$  method (Livak and Schmittgen, 2001). There were no changes in  
208 *rpl8* as a function of either ether stress or 2-wk food deprivation.

209 2.9 *Statistical analysis*

210 Data were analyzed using parametric statistics using Student's two-tailed *t*-test or  
211 ANOVA. Data were tested for homogeneity of variance using Bartlett's test. If data failed to  
212 meet the criteria for parametric tests nonparametric tests were used as needed. All statistical

213 analyses will be performed using InStat (v. 2.05a, GraphPad Software, San Diego, CA) or SPSS  
214 (v. 11, SPSS Inc., Chicago, IL).

215 **3. Results**

216 *3.1. Effects of stressors on food intake and regional brain CRF content.*

217 Food intake was reduced when exposed to the ether stressor, but not the shaking stressor  
218 (FIG.1;  $F_{2,29} = 6.16$ ;  $p < 0.05$ ). To avoid any effect of differences in food intake on brain CRF  
219 content, we examined stressor effects on CRF content in a separate experiment. The greatest  
220 concentration of CRF per unit protein was found in the H/T of control animals followed by the  
221 OT, Tel, and BS (FIG.2A). One-way ANOVA revealed no statistically significant effect of  
222 either stressor on CRF content in the Tel (FIG.2B;  $F_{3,15} = 3.18$ ,  $p = 0.08$ ). Bartlett's test for  
223 homogeneity of variance revealed that the CRF measurements in the H/T did not meet the  
224 requirements for parametric ANOVA, thus differences were assessed by non-parametric  
225 Kruskal-Wallis test followed by Dunn's test. This revealed an effect of stressor treatment (Fig  
226 2b;  $KW = 8.03$ ,  $p < 0.01$ ), with the difference primarily between the ether stressor and the  
227 shaking stressor, but not between either stressor and controls. Exposure to ether vapors elevated  
228 OT CRF content nearly four-fold relative controls (FIG.2b;  $F_{2,15} = 5.8$ ,  $p < 0.05$ ), while there was  
229 no effect of shaking on CRF content in this brain area. The same pattern was apparent in the BS  
230 where ether vapors, but not shaking, increased CRF content relative to controls (FIG.2B;  $F_{2,12} =$   
231  $4.23$ ,  $p < 0.05$ ).

232 *3.2. Effects of an ether stressor on regional brain crf and crfr1 transcript abundance.*

233 In a separate experiment we next investigated whether a reactive stressor, ether vapors,  
234 altered the abundance of transcripts encoding *crf* and *crfr1*. We did not evaluate the shaking  
235 stressor since it did not alter food intake or tectal CRF in our hands. Exposure to ether had no

236 effect on *crf* transcript abundance in Tel, H/T, or BS. However, ether exposure elevated (p <  
237 0.01, Student's two-tailed *t*-test) *crf* transcript abundance in the OT (FIG.3a). The abundance of  
238 *crfr1* transcripts were dramatically reduced (p < 0.05, Student's two-tailed *t*-test) in Tel and OT  
239 (p= 0.04 and p=0.03, respectively, Student's two-tailed *t*-test) in response to ether, while levels in  
240 H/T and BS were unaffected.

241 *3.3. Effects of food deprivation on regional brain CRF content and crf and crfr1 transcript*  
242 *abundance*

243 There was no difference in body mass between the control group ( $11.8 \pm 1.02$  g, n = 8)  
244 and food deprivation group ( $11.9 \pm 0.87$  g, n = 8) at the start of the experiment. After 1 wk of  
245 food deprivation, mean body mass tended to be lower ( $11.0 \pm 0.60$  g) but was not statistically  
246 different than in controls ( $12.4 \pm 1.06$  g, p = 0.27). This trend persisted 2 wk after the start of  
247 food deprivation ( $10.8 \pm 0.58$  g) although the difference relative to controls ( $12.4 \pm 1.06$ ) still  
248 was not statistically different (p= 0.14). As with the other experiment (Fig. 2A), the largest  
249 concentration of CRF per unit protein was found in the H/T of control animals followed by the  
250 OT, Tel, and BS (FIG. 4A). Following 2 wk of food deprivation, CRF content was lower in the  
251 OT (p > 0.05) two-tailed Student's *t*-test), higher in the BS (p = 0.03), but unchanged in the TEL  
252 (p > 0.05) or H/T (p > 0.05) (FIG. 4b).

253 In the second food deprivation experiment for qRT-PCR analysis, starting body mass was  
254  $39.9 \pm 4.05$  g (n = 7) for the controls and  $39.9 \pm 4.58$  g (n = 7) for the frogs in the deprivation  
255 group (p = 0.99). After one week of food deprivation body mass in controls remained unchanged  
256 at  $39.2 \pm 3.98$  g in the control group and  $37.8 \pm 3.64$  g in the food deprived animals (p = 0.79).  
257 By the end of the two-week experiment body mass in controls ( $38.4 \pm 4.08$  g) remained  
258 unchanged from the food deprived animals ( $37.2 \pm 3.99$  g, p = 0.84). Whole body fat analysis

259 using the noninvasive EchoMRI method revealed a decrease in body fat after 2 wk of food  
260 deprivation (Table 2,  $p < 0.05$  Student's one-tailed  $t$ -test). Two weeks of food deprivation did  
261 not statistically alter *crf* and *crfr1* transcript abundance in the OT or any other brain area (FIG.5).

#### 262 **4. Discussion**

263 The presence of a predator reduces food intake in *X. laevis* (Duggan et al., 2016), consistent with  
264 a large volume of work in other animals that under predation threat animals stop foraging to  
265 reduce their vulnerability to a predator (Lima, 1998; Caro, 2005; Cresswell, 2008; Ferrari et al,  
266 2009; Harris and Carr, 2016). Our data indicate that noxious stimuli known to activate the HPA  
267 axis (ether vapors, Olsen et al., 1999; shaking, Boorse and Denver, 2004) but which are  
268 otherwise ecologically irrelevant, have different effects on food intake in *X. laevis*, with ether  
269 vapors significantly reducing food intake and shaking having no effect (FIG.1). Considering that  
270 the animals tested were never previously exposed to ether vapors, our data suggest that the  
271 reduction in food intake caused by a reactive stressor is an innate response. In laboratory rodents  
272 ether vapors have been characterized as a reactive stressor (called initially a systemic stressor by  
273 Emmert and Herman, 1999) as this noxious stimulus vagal afferent sensory pathways are  
274 activated, in turn activating ascending A2 noradrenergic neurons in the nucleus of the solitary  
275 tract that project to the paraventricular nucleus and stimulate CRF release synthesis and release.  
276 Whether A2 noradrenergic neurons innervate the OT is unknown at this time. The fact that an  
277 ecologically relevant and completely novel stimulus may elicit this reduction in food intake is  
278 significant, as it speaks to the evolutionary selection pressures sculpting threat response/prey  
279 capture tradeoffs. The pressure to respond physiologically to an entirely novel threat overrides  
280 the pressure to maintain energy balance through prey capture. Whether a reactive stressor causes  
281 reductions in food intake in the same mechanisms involved in predator-induced food satiety in *X.*

282 *X. laevis* (Duggan et al., 2016) is unknown at present but is doubtful, as detection of a predator  
283 likely involves olfactory, visual, and lateral line sensory modalities rather than inflammation and  
284 cytokine activation of vagal afferents that most likely characterize the sensory pathway involved  
285 in ether stressor. Obviously this does not rule out the possibility that there is a final common  
286 pathway target by multiple modalities for reduction of prey capture in response to a threat.

287       Although the search for a CRF initially began to identify hypophysiotropic factors  
288 stimulating corticotropin release from the anterior pituitary gland, almost 40 yrs of research since  
289 the peptide's discovery have revealed a much broader distribution in many extrahypothalamic  
290 brain areas. While our laboratory (Carr et al., 2010) was not the first to report on CRF-  
291 immunoreactive neurons in the anuran OT (Yao et al., 2004; Calle et al, 2006), our findings are  
292 the first to suggest that tectal CRF neurons may play a role in modulating sensorimotor signaling  
293 in the OT in response to a reactive stressor that inhibits food intake. As we have shown before in  
294 *Bufo marinus*, and as Boorse and Denver (2004) have reported in *X. laevis*, tectal CRF  
295 concentrations are second only to those in the hypothalamus with the rank order of tectal CRF  
296 being hypothalamus > OT > Tel > BS in our hands (Figs. 2 and 4). Our data indicate an increase  
297 in CRF peptide content in the OT and brainstem after ether vapor exposure. Although we have  
298 shown that ether vapor exposure elevates plasma corticosterone in toads (Olsen et al., 1999), and  
299 H/T CRF content was not elevated over controls in this study. Shaking, which elevates anuran  
300 hypothalamus-pituitary-interrenal activity through an as yet unidentified pathway and increases  
301 CRF immunoreactivity in the preoptic area (Yao and Denver, 2004), elevated CRF content in the  
302 H/T, and while ANOVA revealed an overall effect post-hoc testing revealed this difference was  
303 between the ether and shaking stress groups, not the shaking stress and controls. It may be that  
304 the immunohistochemical approach for gauging regional changes in the hypothalamus and

305 forebrain CRF content is more powerful than our approach because of the more detailed  
306 anatomical resolution that it provides. Unfortunately, our results are not directly comparable to  
307 those of Boorse and Denver (2004) as hypothalamus, preoptic area, and part of the OT were  
308 included in the same tissue pieces analyzed for CRF. The focus of our work is on the OT, which  
309 is why we analyzed the intact OT separately from other brain areas. Nonetheless the trend for  
310 increased CRF content in the H/T after 4 h of shaking is still apparent in our data set.

311         Elevated CRF peptide content in the OT after a reactive stressor was accompanied by  
312 increased CRF transcript abundance in the OT (FIG.3), suggesting that at least some of the  
313 increase in tectal CRF during stress is contributed by tectal CRF neurons. The dissection method  
314 employed in our study eliminates the possibility that the hypothalamic, forebrain, or tegmental  
315 CRF neuronal populations are included in the OT samples. Although there is general (but not  
316 unanimous; Yamano et al., 2004) agreement that a wide variety of anticipatory and reactive  
317 stressors increase CRF transcript abundance in the hypophysiotropic neurons of the PVN, the  
318 influence of various stressors on CRF in extrahypothalamic locations has not been well studied  
319 although reports of stressor induced elevations of CRF and cRF mRNA in extrahypothalamic  
320 areas can be found. Chen et al. (2004) reported the hippocampal neurons release CRF in  
321 response to an acute anticipatory stressor. Hatalski et al. (1998) found that repeated exposure to  
322 an acute reactive stressor (cold) elevated CRF transcript abundance in the central nucleus of the  
323 amygdala while Yamano et al. (2004) reported a similar finding after foot shock in rats.

324         Patterns in *crfr1* mRNA abundance after a stressor exposure are not as clear, and seem to  
325 depend upon the tissue type and brain area examined as well as the type of stressor employed. In  
326 an early study Rivest et al. (1995) reported that acute exposure to immobilization elevated *crfr*  
327 transcript expression in the PVN as well as several other brain areas as determined by *in situ*

328 hybridization. Similarly, Luo et al. (1995) reported that an acute immobilization stressor elevated  
329 *crfr1* mRNA abundance in the PVN. In contrast, Roseboom et al. (2007) found no change in  
330 *crfr1* mRNA abundance in the PVN of rats exposed to a predator (ferret). In the pituitary gland,  
331 acute stressor exposure results in a biphasic change in *crfr* mRNA in rats, with a decrease  
332 observed at 2 h followed by a return to basal levels or an increase after 4 h. While  
333 immobilization leads to an increase in *crfr1* mRNA, as it does in the PVN in some studies, some  
334 stressors such as lipopolysaccharide consistently decrease pituitary *crfr* (Rabadan-Diehl et al.,  
335 1996). Studies on the relationship between transcript and receptor abundance in rat pituitary  
336 show a remarkable disconnect, suggesting that *crfr* mRNA may not reflect in any way actual  
337 receptor availability (Aguilera et al., 2004). Thus, our observation of a decrease in *crfr1* mRNA  
338 in the OT after an acute reactive stressor must be considered with this last point in mind. It is  
339 interesting to speculate that this decrease in mRNA might reflect down regulation of *crfr1*  
340 transcript levels associated with increased CRF receptor activation as a result of stress, but such a  
341 conclusion awaits ligand binding studies.

342 To our knowledge our study is the first to measure tectal CRF content after a change in  
343 energy status. We measured a decrease in CRF peptide content (FIG.4B) in the OT after 2 wk of  
344 food deprivation. While 2 wk of food deprivation did not alter total body mass, it did  
345 significantly reduce whole body fat content as measured by the non-invasive EchoMRI method,  
346 which allows us to measure body composition in the same animal before and after the  
347 experimental manipulation. Two weeks of food deprivation lowered mean *crf* transcript  
348 abundance in the OT, but the difference from control levels was not statistically significant ( $p >$   
349 0.05). Mercer et al. (1996) reported an increase or no change in *crf* mRNA abundance in the  
350 PVN after food deprivation in hamsters while others (Hwang and Guntz, 1997) have reported a

351 decrease in *crf* mRNA in the PVN after food deprivation. Obviously we cannot conclude that  
352 the reduction in tectal CRF was due entirely to reduced *crf* transcript expression by tectal  
353 neurons, and may involve changes in peptide content in CRF neurons innervating the OT from  
354 elsewhere. Given the diverse number of CRF cell populations in the anuran brain there is the  
355 possibility the not all tectal CRF arises from tectal CRF neurons. It is possible that CRF  
356 producing neurons in the limbic nucleus, preoptic area, and perhaps the mesencephalic  
357 tegmentum and brainstem project to the OT (Carr et al., 2010), and that reductions in the  
358 activities of these neurons contribute to the lower tectal CRF content we observed after food  
359 deprivation. This is especially important to consider given that food deprivation is likely to  
360 impact many different neuronal populations in the brain. Neuronal tracing studies will be  
361 required to identify any extra-tectal CRF cell populations that may innervate the OT in *X. laevis*.

362 Overall our data are consistent with a potential role for tectal CRF in reducing prey  
363 capture, although our data cannot allow us to conclude that changes in tectal CRF cause reduced  
364 food intake. Current ongoing work in our laboratory is addressing this question directly with the  
365 use of CRF receptor agonists and antagonists microinjected into the OT. Our data add to an  
366 increasing literature pointing out the significance of extrahypothalamic CRF cell populations in  
367 adaptive behavior and homeostatic regulation.

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584 FIGURE LEGENDS

585 **Figure 1.** Mean + S.E.M. food intake in in juvenile *Xenopus laevis* exposed to shaking (n=15) or  
586 ether (n=5) stressors or left untreated (controls, n = 12). Bars with different superscripts are  
587 statistically different based upon one-way ANOVA (p < 0.05).

588

589 **Figure 2.** Regional differences in CRF content in control animals and in response to two  
590 stressors, ether vapors and shaking. **FIG.2A.** Mean (+ S.E.M.) CRF concentrations, expressed  
591 per unit protein, in the telencephalon (Tel), hypothalamus/thalamus (H/T), optic tectum (OT),  
592 and brainstem (BS) of untreated juvenile *Xenopus laevis* (n=6). **FIG.2B.** Fold change (mean +  
593 S.E.M, n=4-6 per group) in CRF content relative to controls after ether and shaking stressors.  
594 Superscripts indicate significant differences within brain regions.

595

596 **Figure 3.** Regional differences in relative *crf* (FIG.3A) and *crfr1* (FIG.3B) transcript abundance  
597 of sub-adult *Xenopus laevis* in response to a reactive stressor, ether vapors. Telencephalon,  
598 (Tel); hypothalamus/thalamus, H/T; optic tectum, OT; brainstem, BS. Bars represent the mean +  
599 S.E.M. of 4-7 animals per group. Asterisks indicate significant difference from control based  
600 upon Student's t-test.

601

602 **Figure 4.** Regional differences in CRF content in control and food deprived animals after 2 wk  
603 food deprivation. **FIG. 4A.** Mean (+ S.E.M.) CRF concentrations, expressed per unit protein, in  
604 the telencephalon (Tel), hypothalamus/thalamus (H/T), optic tectum (OT), and brainstem (BS) of  
605 untreated juvenile *Xenopus laevis* from this experiment. **FIG. 4B.** Fold change in CRF content  
606 relative to controls after 2 wk food deprivation. Bars represent the mean + S.E.M. of 4-7 animals

607 per group. Asterisks indicate significant difference from control based upon Student's two-tailed  
608 *t*-test.

609 **Figure 5.** Regional differences in relative *crlf* (FIG. 3A) and *crlf1* (FIG. 3B) transcript abundance  
610 of sub-adult *Xenopus laevis* in response to 2 wk food deprivation. Telencephalon, (Tel);  
611 hypothalamus/thalamus, H/T; optic tectum, OT; brainstem, BS. Bars represent the mean +  
612 S.E.M. of 4-6 animals per group. No significant differences were observed.

613

614

**Table 1. List of Abbreviations**

615	A2 noradrenergic neurons,	Cells staining fluorescently for norepinephrine in the caudal brainstem as identified by Dahlstrom and Fuxe, 1964.
616		
617		
618	BS	Brainstem
619	CNS	Central nervous system
620	CRF	Corticotropin-releasing factor
621	CRFR2	Corticotropin-releasing factor receptor, Type 2
622	Ct	Cycle threshold
623	HPA	Hypothalamus-pituitary-adrenal axis
624	H/T	Hypothalamus/thalamus
625	OT	Optic tectum
626	PVN	Paraventricular nucleus
627	qRT-PCR	Quantitative Real Time-Polymerase Chain Reaction
628	RIA	Radioimmunoassay
629	Rpl8	Ribosomal protein L8
630	RQI	RNA quality indicator
631	Tel	Telencephalon
632	UCN1	Urocortin 1
633	UCN2	Urocortin 2
634	UCN3	Urocortin 3
635		
636		
637		
638		

639

640

641 Table 2. Mean Fat Mass (g)  $\pm$  S.E.M Before and after Two Weeks of Food Deprivation in *X.*  
642 *laevis.*

Treatment	n	Initial Fat Mass (g)	Final Fat Mass (g)	Ratio (final/initial)
<b>Control</b>	7	3.46 $\pm$ 0.37	3.52 $\pm$ 0.28	1.06 $\pm$ 0.07
<b>Food deprived</b>	7	4.66 $\pm$ 0.54	4.01 $\pm$ 0.39	0.88 $\pm$ 0.05*

643 Asterisk indicates significant decrease relative to controls based on one-tailed Student's *t*-test.

644

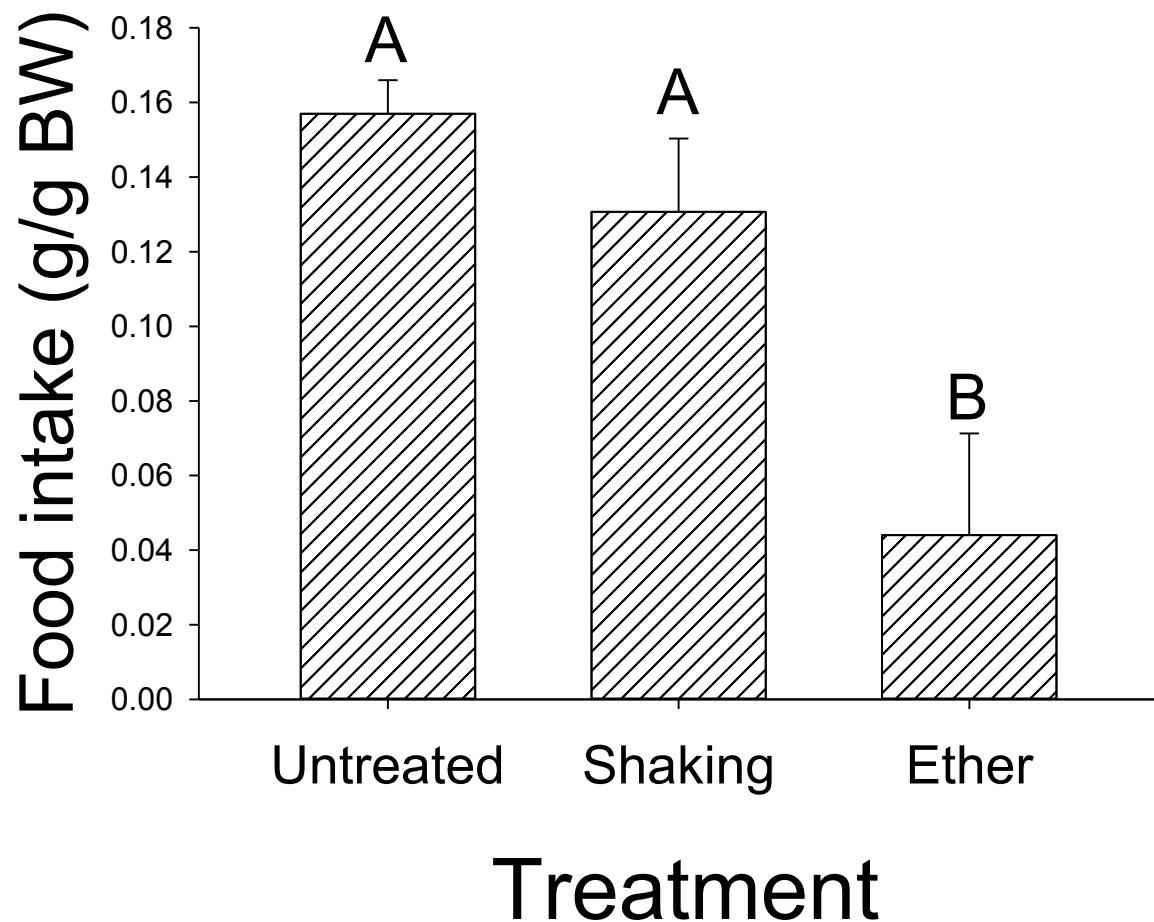


Fig. 2

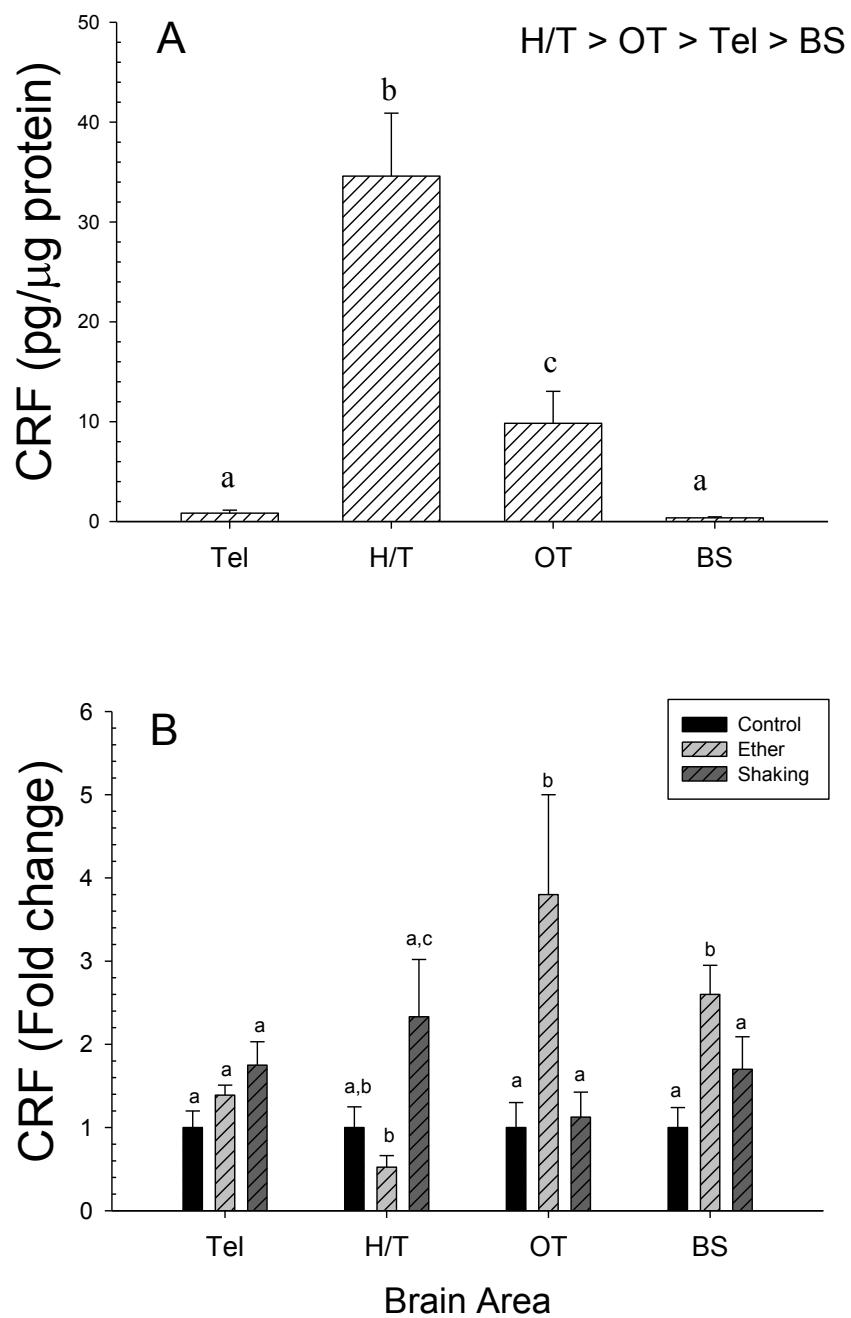


Fig. 3

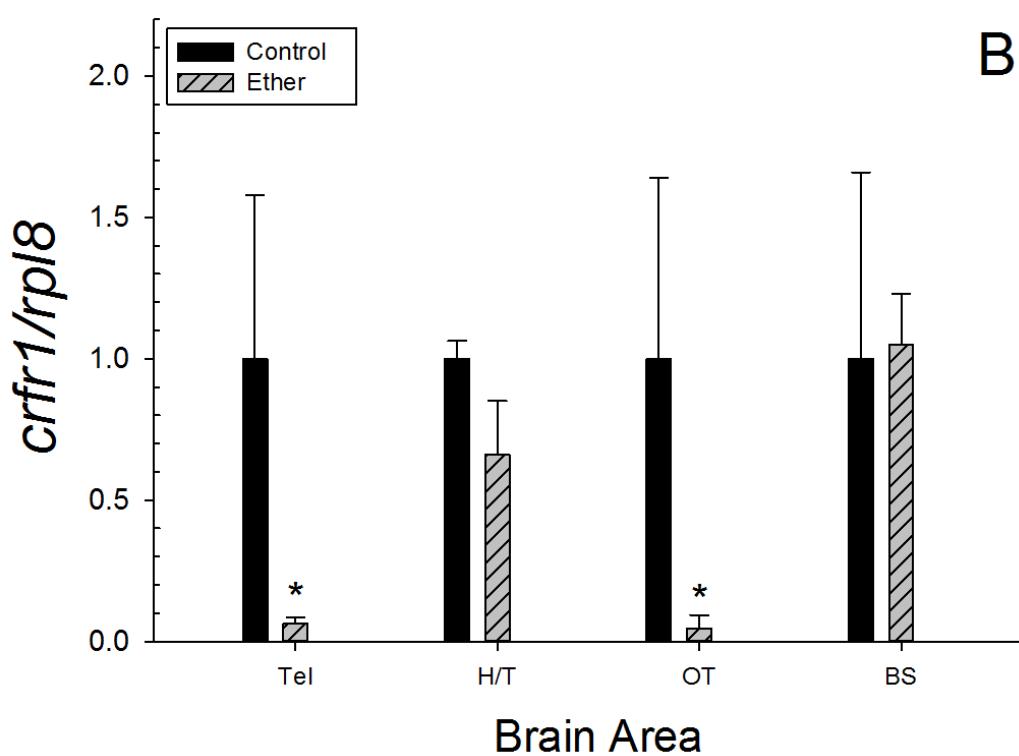
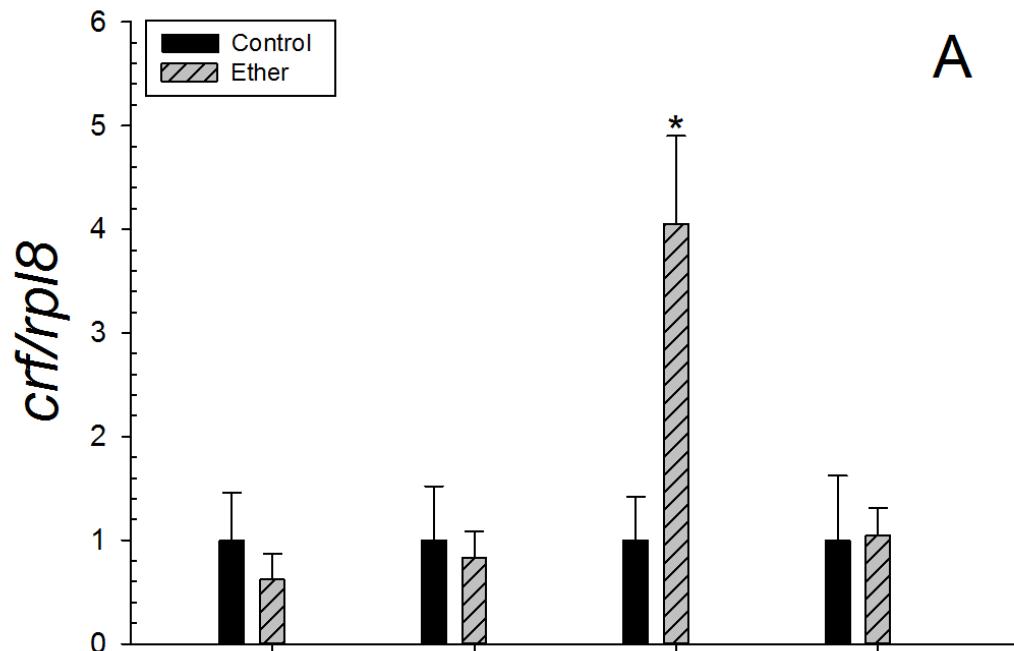


Fig. 4

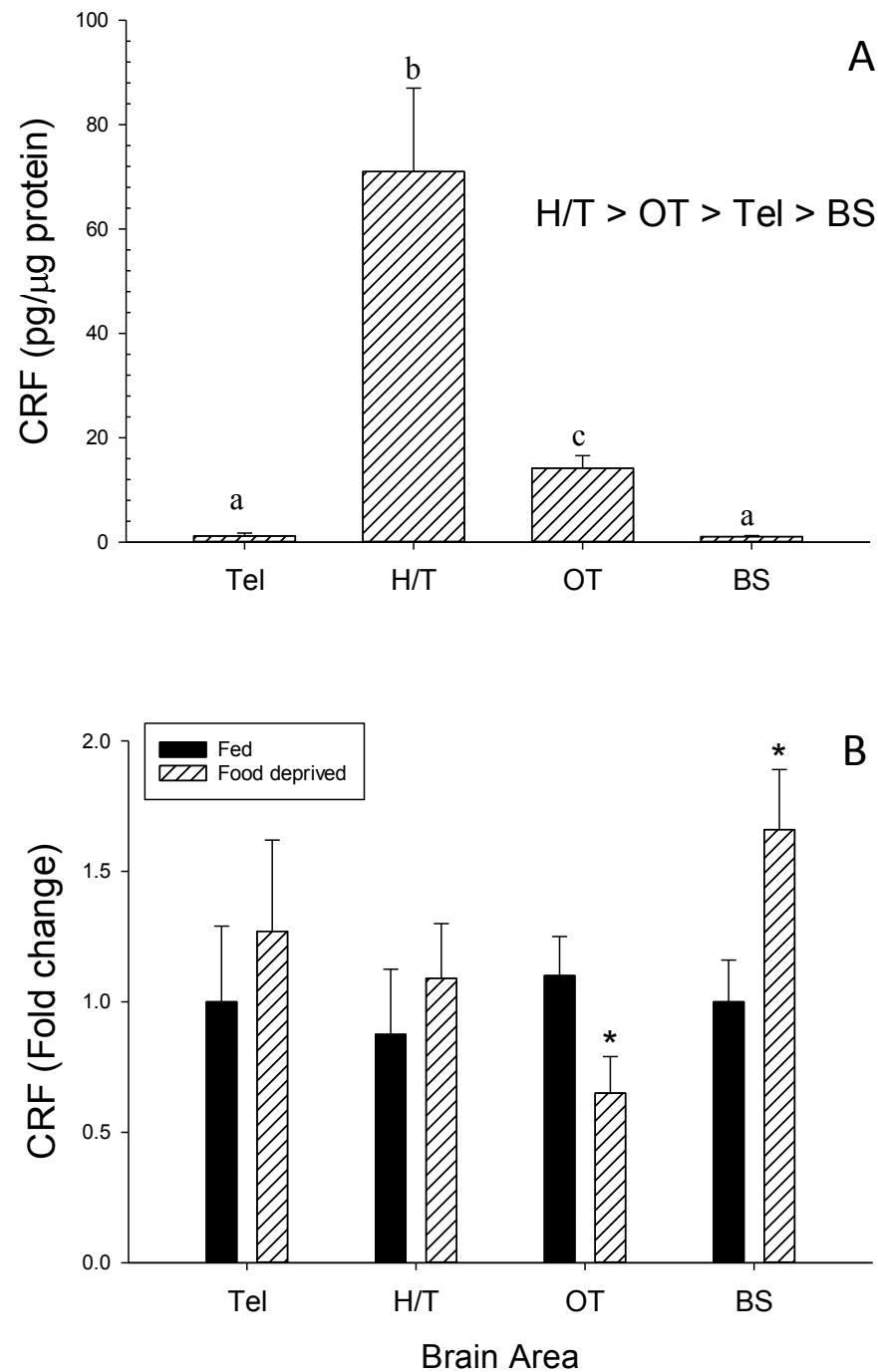
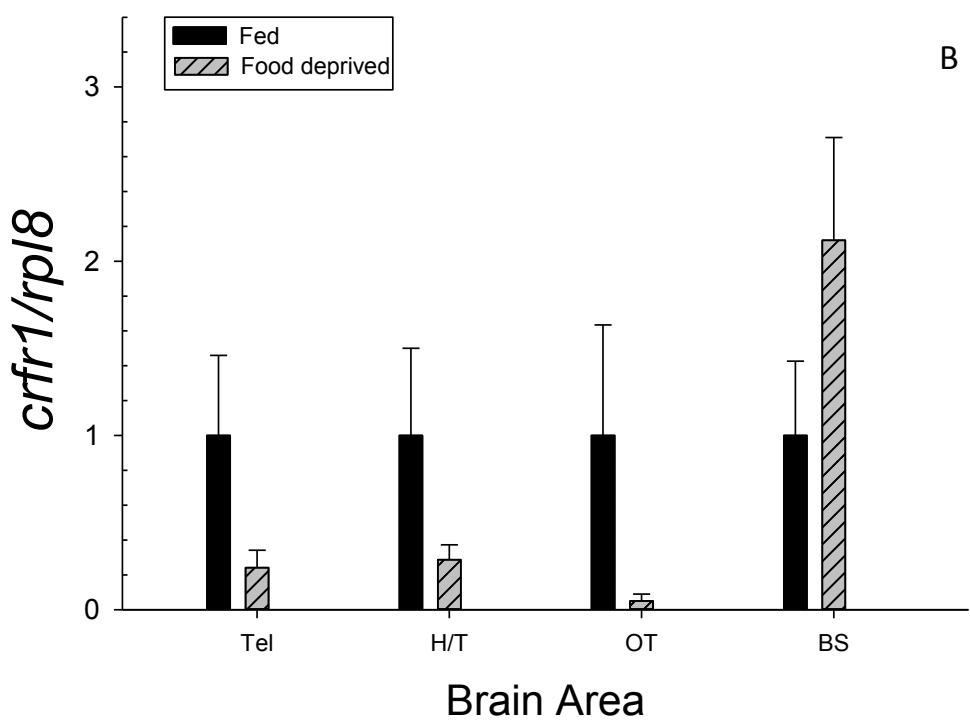
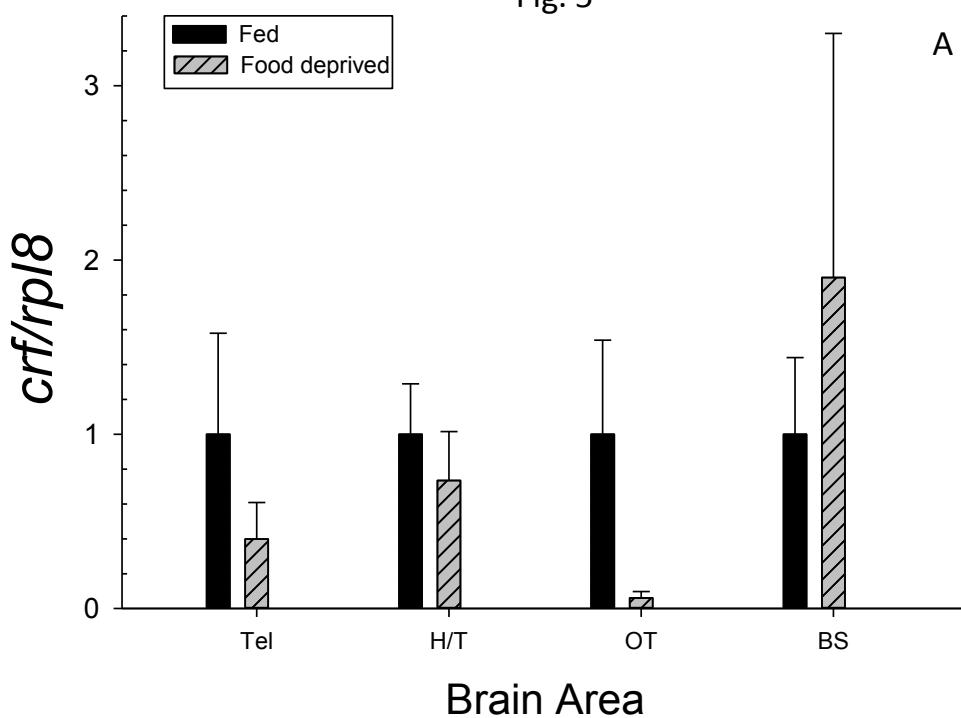


Fig. 5



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652