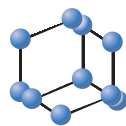
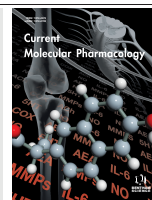


REVIEW ARTICLE

The CRF Family of Neuropeptides and their Receptors - Mediators of the Central Stress Response

**BENTHAM
SCIENCE**Nina Dedic¹, Alon Chen^{1,2,*} and Jan M. Deussing¹¹Department of Stress Neurobiology and Neurogenetics, Max Planck Institute of Psychiatry, Kraepelinstr, 2-10, 80804 Munich, Germany; ²Department of Neurobiology, Weizmann Institute of Science, 76100 Rehovot, Israel

Abstract: Background: Dysregulated stress neurocircuits, caused by genetic and/or environmental changes, underlie the development of many neuropsychiatric disorders. Corticotropin-releasing factor (CRF) is the major physiological activator of the hypothalamic-pituitary-adrenal (HPA) axis and consequently a primary regulator of the mammalian stress response. Together with its three family members, urocortins (UCNs) 1, 2, and 3, CRF integrates the neuroendocrine, autonomic, metabolic and behavioral responses to stress by activating its cognate receptors CRFR1 and CRFR2.

Objective: Here we review the past and current state of the CRF/CRFR field, ranging from pharmacological studies to genetic mouse models and virus-mediated manipulations.

Results: Although it is well established that CRF/CRFR1 signaling mediates aversive responses, including anxiety and depression-like behaviors, a number of recent studies have challenged this viewpoint by revealing anxiolytic and appetitive properties of specific CRF/CRFR1 circuits. In contrast, the UCN/CRFR2 system is less well understood and may possibly also exert divergent functions on physiology and behavior depending on the brain region, underlying circuit, and/or experienced stress conditions.

Conclusion: A plethora of available genetic tools, including conventional and conditional mouse mutants targeting CRF system components, has greatly advanced our understanding about the endogenous mechanisms underlying HPA system regulation and CRF/UCN-related neuronal circuits involved in stress-related behaviors. Yet, the detailed pathways and molecular mechanisms by which the CRF/UCN-system translates negative or positive stimuli into the final, integrated biological response are not completely understood. The utilization of future complementary methodologies, such as cell-type specific Cre-driver lines, viral and optogenetic tools will help to further dissect the function of genetically defined CRF/UCN neurocircuits in the context of adaptive and maladaptive stress responses.

Keywords: Corticotropin-releasing factor, urocortin, stress, mouse genetic tools, hypothalamic-pituitary-adrenal (HPA), neuropsychiatric disorders.

1. INTRODUCTION

“It’s not stress that kills us, it is our reaction to it” Hans Selye (1907-1982).

Over the few past decades, growing evidence has linked life stress to various pathologies, including cardiovascular disease, inflammation, metabolic dysfunctions, and most prominently neurodegenerative and psychiatric disorders. In particular, conditions of severe prolonged stress are considered to be the most devastating because they tend to induce long-term or permanent changes in the physiological, emotional and behavioral responses that influence susceptibility to disease. Persistent stress, such as prolonged exposure to war, physical abuse, devastating socioeconomic status or

social/psychological surroundings, has been shown to increase the likelihood of developing depression and anxiety disorders, cognitive dysfunction, metabolic conditions such as obesity and diabetes, as well as sleep and cardiovascular disorders, to name just a few [1-7]. However, we tend to forget that stress *per se* is not a bad thing, but rather the reaction and/or inability to adapt to it that constitutes health or disease. Importantly, acute stress can exert a wide range of positive effects, as it primes the brain towards optimal alertness, behavioral and cognitive performance [8-13]. The reaction to stress represents an adaptive mechanism, triggering the so-called “fight-or-flight” response in order to cope with a dangerous situation, be it a predator, an accident, or a natural disaster. Stress can be discriminated on the one hand into eustress, or “positive” stress, meaning that the succeeding adaptive response is able to re-instate homeostasis, and on the other hand into distress, or “negative” stress resulting in pathological outcomes [14]. In general, we are not equipped to withstand chronic activation of specific stress-pathways,

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which is increasingly occurring in today's urbanized social environments partially due to disparities in income, education, occupation and other dimensions of socioeconomic status [2]. But when does stress, or more precisely the response to stress, cross the line from being adaptive to maladaptive? This question is extremely difficult to answer, considering that the threshold of stress-resistance is different for each individual and is influenced to a variable degree by genetic predisposition [15, 16].

Two closely interplaying systems are primarily responsible for orchestrating the stress response: the sympathetic nervous system (SNS) and the hypothalamic-pituitary-adrenal (HPA) axis. The SNS is largely responsible for initiating the "flight-or-flight" reaction by stimulating, amongst others, the release of adrenaline and noradrenaline from the adrenal medulla. The latter exerts its commands at multiple sites, including the spinal cord, medulla, pons and higher order centers such as the hypothalamus [17, 18]. The HPA axis is characterized by the release of different neuropeptides and hormones, and is believed to mediate the immediate, as well as the long-lasting effects of stress. As a result of the two interplaying systems, various substances are released in response to stress, which are then orchestrated into a coordinated physiological and behavioral response [19]. These so-called stress-mediators are broadly classified into three groups; the monoamines, neuropeptides and steroids. Importantly, different stressors are processed by distinct circuits and/or in specific brain areas. The non-specific effects of stress are mirrored by the rapid activation of the SNS and the neuroendocrine arm of the stress response, *i.e.* the HPA axis, which in turn regulates the synthesis and release of glucocorticoids (GCs) from the adrenal glands [19]. Conversely, psychological and anticipatory stressors (stress due to hypothetical events that may/may not occur) are primarily processed by the limbic system, including the hippocampus, amygdala and prefrontal cortex [19, 20]. The complex interaction between these different stress-mediators and pathways has gained substantial interest in the past [21]. However, some play very specific and potentially even opposing roles than others and are especially relevant in dysfunctional stress circuits, which can result in various neuropsychopathologies [19, 22]. Already in 1955, Hans Selye postulated that "through some unknown pathway, the "first mediator" travels directly from the injured target area to the anterior pituitary. It notifies the latter that a condition of stress exists and thus induces it to discharge adrenocorticotrophic hormone (ACTH)" [23]. It took another 26 years until Wylie Vale's group discovered this central stress mediator - the neuropeptide corticotropin-releasing factor (CRF). This major breakthrough contributed significantly to our understanding of the neurobiological mechanisms underlying the stress response [24].

2. CRF MODULATES THE NEUROENDOCRINE STRESS RESPONSE VIA THE HPA AXIS

CRF (also referred to as corticotropin-releasing hormone – CRH) is the major physiological activator of the HPA axis, and coordinates the neuroendocrine response to stress. Perception of physical or psychological stress by an organism is followed by a series of events, including the release of CRF from parvocellular neuroendocrine neurons of the paraven-

tricular nucleus of the hypothalamus (PVN). These neurons project *via* the external zone of the median eminence and release CRF into the hypophysial portal vasculature, which transports the neuropeptide to secretory corticotrope cells of the anterior pituitary, which express the CRF receptor type 1 (CRFR1; Fig. 1). The activation of CRFR1 stimulates the release of ACTH and other pro-opiomelanocortin (POMC) - derived peptides [25]. ACTH, in turn, triggers the synthesis and release of GCs from the adrenal cortex (cortisol in humans, corticosterone in rodents), which mediate numerous physiological and metabolic reactions and ultimately prepare the organism to deal with the stressful situation. These responses to GCs include cardiovascular activation, energy mobilization, anti-inflammatory effects and suppression of reproductive and digestive functions, [26-32]. In order to restore the HPA axis to its normal state and to protect it from overshooting, GCs signal back *via* glucocorticoid (GR) and mineralocorticoid receptors (MR) at various feedback levels (*e.g.* pituitary, hippocampus, PVN and amygdala), which ultimately inhibit the secretion of CRF and consequently ACTH (Fig. 1). Similar to many physiological processes in the body, GCs exhibit a circadian rhythm, with increased levels toward the active phase of the light/dark cycle, which is regulated by the main circadian pacemaker in the suprachiasmatic nucleus [33, 34]. At this point it is also important to note that the HPA axis is not exclusively activated during aversive stressful situations. In fact, the physiological stress-response to appetitive, rewarding stimuli (which are generally not considered as stressors) can be as large as the response to a negative stimulus. For example, positive experiences such as sexual encounter, wheel running and social victory in rats induce a similar degree of HPA axis activation as an aversive footshock, social defeat and restraint stress [35, 36]. Consequently, HPA axis activation and enhanced cortisol levels are not solely indicative of an aversive stressful state.

3. THE FAMILY OF CRF-RELATED NEUROPEPTIDES AND THEIR RECEPTORS

The actions of CRF are not confined to the neuroendocrine HPA system. The anatomical distribution of CRF in the brain suggests that this peptide not only acts as a key neuroendocrine stress mediator, but is also able to regulate neuronal activity in a neuromodulatory fashion. In fact, CRF is expressed throughout the central nervous system (CNS) including in most limbic and cortical structures (Fig. 2), where it has been shown to regulate the emotional and cognitive components of the stress response. The mature and biologically active form of CRF is a 41-amino acid peptide generated by proteolytic cleavage from a 196-amino acid precursor. To date, the mammalian CRF family comprises three additional peptides (Fig. 3). Urocortin (UCN) 1 was initially described in 1995 by Vaughan and colleagues [37] followed by the discovery of UCN2 (or stresscopin-related peptide) and UCN3 (or stresscopin) shortly afterwards [38-40]. CRF is most closely related to UCN1, sharing 43% amino acid homology, whereas CRF sequence identity with UCN2 and UCN3 is 34% and 26% respectively [41]. In comparison to CRF, urocortin-expressing neurons are found in more discrete regions and nuclei of the CNS (Fig. 2). In rodents, UCN1 is mainly expressed in the Edinger-Westphal nucleus

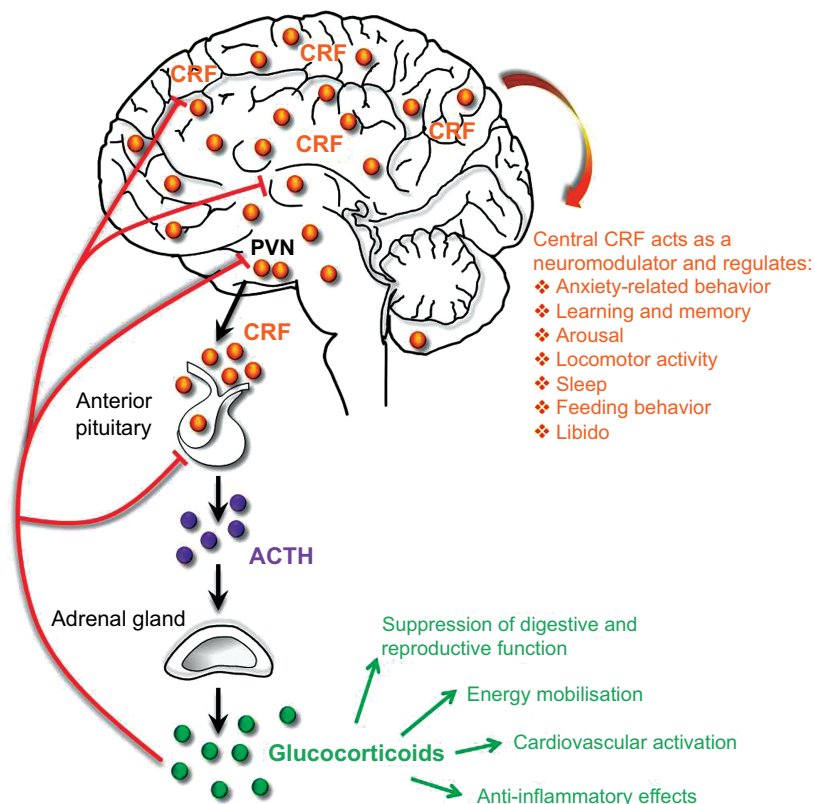


Fig. (1). CRF regulates neuroendocrine and behavioral responses to stress. CRF integrates neuroendocrine and higher-order behavioral responses by regulating peripheral HPA axis function and modulating synaptic transmission in the CNS. ACTH: Adrenocorticotropic hormone; CRF: corticotropin-releasing factor, PVN: hypothalamic paraventricular nucleus.

and sparsely distributed in the lateral superior olive and supraoptic nucleus [37]. UCN2 is found in the rodent PVN, supraoptic nucleus, arcuate nucleus, locus coeruleus and brainstem, whereas UCN3 is mainly expressed in the medial amygdala, rostral perifornical area of the hypothalamus, the bed nucleus of the stria terminalis (BNST), superior paraolivary nucleus, nucleus parabrachialis and the premammillary nucleus (Fig. 2); [38-40, 42]. All four neuropeptides have also been detected in the periphery, in particular UCN2 and UCN3, which have recently been recognized as novel modulators of centrally- and peripherally-controlled metabolic function [43-45].

In contrast to the detailed expression profiles in rodents, data is more limited with respect to CRF and UCN1-3 distribution within the human CNS. However, a few studies have examined CRF/UCN1 expression in human postmortem brains. In accordance with the observations in mice and rats, CRF immunoreactivity and mRNA levels in humans were demonstrated in the PVN, pituitary stalk, hypothalamus, and cortex [46-50]. As in rodents, the most abundant expression of UCN1 in humans is found in the Edinger-Westphal nucleus [51, 52]. However UCN1 expression was also detected in the human anterior pituitary and Purkinje cells of the cerebellar cortex, which has not been observed in the mouse brain [52, 53].

CRF and the urocortins signal through the activation of two, class B1, membrane-bound, G-protein-coupled receptors (GPCRs), CRFR1 and CRFR2, which share 70% amino

acid identity [54-58]. The lowest degree of homology exists in their N-terminal extracellular domains (40% identity). In contrast, the transmembrane domains of CRFR1 and CRFR2 are highly homologous (80-85% amino acid identity) [41]. CRF shows a much higher affinity for the CRFR1 than for the CRFR2 while UCN1 displays equal affinities for both receptors (Fig. 3). UCN2 and UCN3, on the other hand, appear to be selective ligands of CRFR2 [38, 41, 56]. Similarly to its main ligand CRF, CRFR1 mRNA is found throughout the rodent CNS including the cortex, cerebellum and limbic forebrain (Fig. 2). It is also highly expressed in anterior pituitary corticotropes where it initiates HPA axis activity in response to CRF binding [59, 60]. CRFR1 has only one known functional splice variant (α) expressed in the CNS [61, 62]. Particularly in the skin, numerous additional splice variants have been identified resulting in soluble or membrane bound isoforms that might potentially affect receptor activity *via* dimerization [62-64]. In contrast, CRFR2 has three functional splice variants in humans (α , β and γ) and two in rodents (α and β) [56, 61, 65-68]. CRFR2 α (the major splice variant in rodents and from here on referred to as CRFR2), displays a more confined but partially overlapping expression with CRFR1, with high densities in the olfactory bulb, BNST, lateral septum (LS), ventromedial hypothalamic nucleus, and dorsal raphe nucleus (Fig. 2); [60, 69-71]. In mice, CRFR2 β is primarily expressed in peripheral tissue (with the highest levels of expression in skeletal muscle, heart and skin), as well as in the choroid plexus of the brain [56, 69].

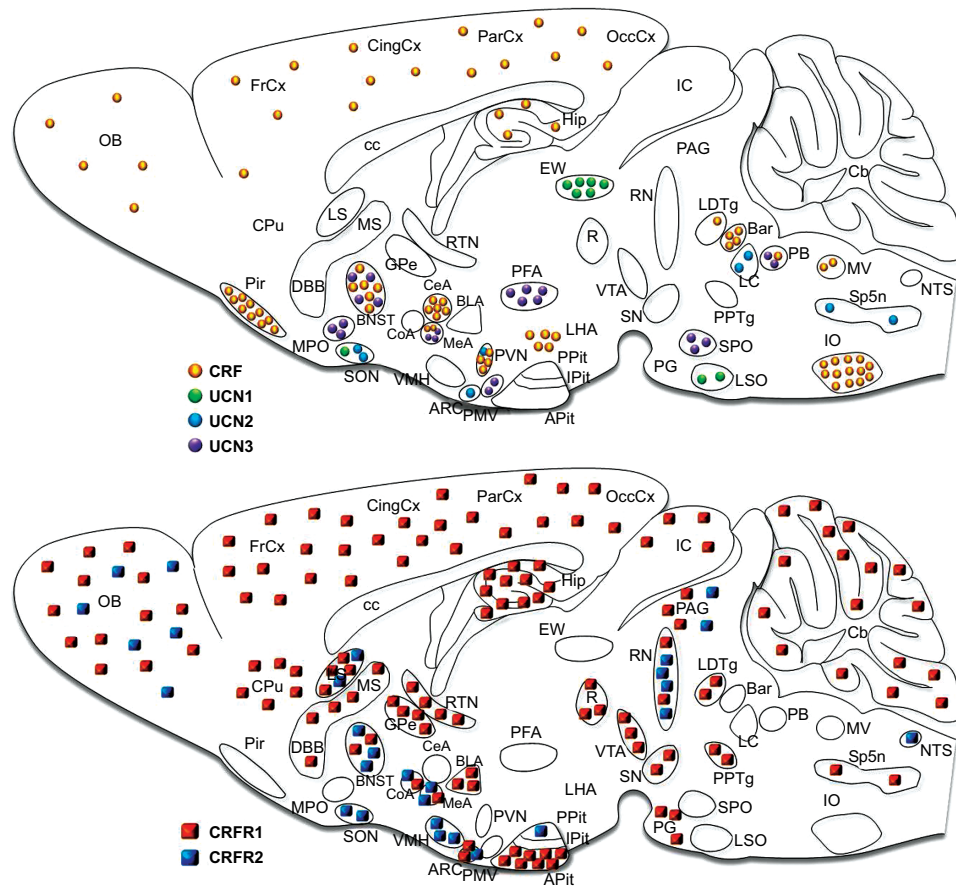


Fig. (2). Schematic illustrations of the spatial distribution and relative expression of CRF family peptides and their receptors in the mouse brain. Abbreviations: Anterior pituitary (APit), arcuate nucleus (ARC), basolateral nucleus of the amygdala (BLA), bed nucleus of the stria terminalis (BNST), caudate putamen (CPu), central nucleus of the amygdala (CeA), cerebellum (Cb), cingulate cortex (CingCx), corpus callosum (cc), cortical nucleus of the amygdala (CoA), Barrington's nucleus (Bar), diagonal band of Broca (DBB), Etinger Westphal nucleus (EW), frontal cortex (FrCx), globus pallidus (GPe), inferior colliculi (IC), inferior olive (IO), intermediate lobe of the pituitary (IPit), locus coeruleus (LC), lateral septum (LS), laterodorsal tegmental nucleus (LDTg), lateral hypothalamic area (LHA), lateral superior olive (LSO), medial nucleus of the amygdala (MeA), medial preoptic area (MPO), medial septum (MS), medial vestibular nucleus (MV), nucleus tractus solitarius (NTS), olfactory bulb (OB), occipital cortex (OccCx), parietal cortex (ParCx), parabrachial nucleus (PB), periaqueductal gray (PAG), perifornical area (PFA), piriform cortex (Pir), pontine gray (PG), posterior pituitary (Ppit), pedunculopontine tegmental nucleus (PPTg), premammillary nucleus (PMV), paraventricular nucleus of the hypothalamus (PVN), red nucleus (R), raphe nuclei (RN), reticular thalamic nucleus (RTN), superior colliculi (SC), substantia nigra (SN), supraoptic nucleus (SON), spinal trigeminal nucleus (Sp5n), superior paraolivary nucleus (SPO), ventral medial hypothalamus (VMH), ventral tegmental area (VTA). Modified from [88].

CRFR1 and CRFR2 are also highly expressed in the human brain, although their expression patterns deviate to some extent from the rodent brain [72-75]. In general, CRFR1 represents the major receptor in the brain of humans and non-human primates, while CRFR2 shows a predominant expression in peripheral tissues [72]. Both receptors are abundantly expressed in the pituitary, which is in contrast to the distribution in rodents, where mainly CRFR1 is found in the pituitary [72, 73]. CRFR1 and CRFR2 are also present in the amygdala, thalamus, and hippocampus. In addition, significant levels of CRFR1 have been detected in the cerebellum and cortex of humans and non-human primates [73, 75]. Another discrepancy between rodents and primates is the presence of CRFR2 in different cortical regions [73]. The absence of CRFR2 from the cerebellum, and a strong expression in the choroid plexus of the human brain is again in line with observations in the rodent brain [73].

The activity of CRF and UCN1 can be regulated additionally by the CRF binding protein (CRF-BP) [41, 76]. Past research has largely ignored the presence of the CRF-BP, which is thought to act as an endogenous buffer, possibly by regulating the availability of active CRF and UCN1 [76-78]. The complexity of the CRF-system is further increased by the recently discovered soluble splice variant of CRFR2 (sCRFR2 α), which encodes the extracellular receptor ligand-binding domain, but terminates before the first transmembrane domain [79]. sCRFR2 α has been proposed to act as a decoy receptor, mimicking the ability of the CRF-BP to sequester free CRF [79]. Alternatively, Evens and colleagues suggested that the unproductive splicing of CRFR2 pre-mRNA to sCRFR2 may selectively alter the cellular levels of full-length CRFR2 mRNA and consequently affect the number of functional CRFR2 receptors, a mechanism common to GPCRs of the secretin family [80, 81]. The diverse and

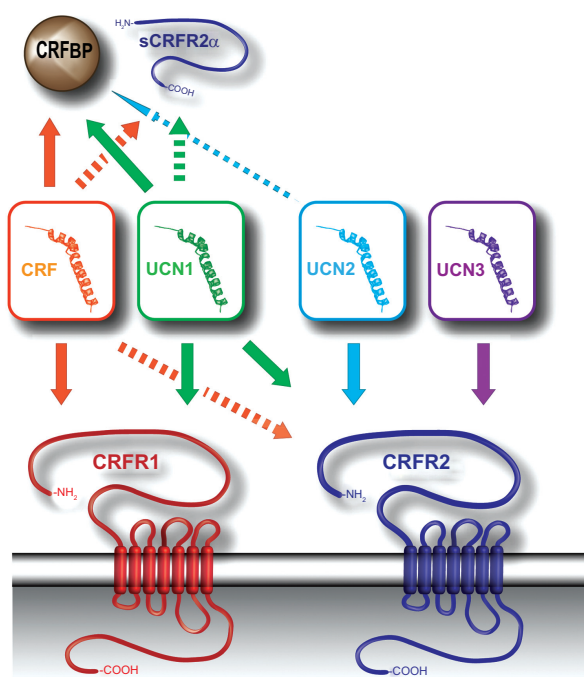


Fig. (3). CRF family members, their receptors and binding proteins. CRF and the UCNs signal through one of two CRF receptors (CRFR1 and CRFR2). The arrows represent ligand-receptor or ligand-binding protein interactions. Dashed arrows indicate low-affinity binding, compared to solid arrow-lines. CRF displays a relatively high affinity for CRFR1 and a low affinity for CRFR2, while UCN1 binds to both receptors with equal affinity. UCN2 and UCN3 are selective ligands for CRFR2. CRFBP and sCRFR2 α are able to sequester both CRF and UCN1, while CRFBP exerts a low affinity for UCN2. Abbreviations: corticotropin-releasing factor (CRF), CRF receptor 2 (CRFR2), CRF binding protein (CRFBP), soluble variant of CRFR2 (sCRFR2 α).

broad expression patterns of CRF-related peptides and receptors, as well as the high level of signaling complexity, enable this circuitry to effectively integrate neuroendocrine, autonomic and behavioral responses of stress.

4. NEUROMODULATORY EFFECTS OF CENTRAL CRF-CRFR1 SIGNALING

As previously introduced, CRF coordinates the physiological/neuroendocrine responses to stress *via* the HPA axis. In addition, CRF and its high affinity receptor, CRFR1, are widely distributed throughout the brain, which allows them to orchestrate autonomic and behavioral stress responses. Consequently, hyperactivity of the CRF/CRFR1 system has been linked to stress-related psychiatric disorders that involve a strong emotional component such as depression and anxiety [6, 82-87]. Alterations in HPA axis function, such as impaired negative feedback, which results in hypercortisolemia, have been reported repeatedly in a subset of depressed patients, and attributed to centrally elevated CRF levels [82, 88]. Increased CRF levels have also been detected in the CSF of untreated depressed patients [87]. Furthermore, reduced CRF binding sites have been identified in the frontal cortex of suicide victims, which was attributed to an adaptive downregulation in response to CRF hypersecretion [89].

Shortly after its isolation in 1981, a number of rodent studies demonstrated that intracerebroventricular (i.c.v.) administration of CRF results in behavioral responses that are similar to those observed in stressed animals. These include increased arousal and anxiety-related behavior, altered locomotor activity and social behavior, diminished sexual behavior and food consumption as well as sleep disturbances [90-99]. Importantly, many of these effects were independent of downstream GC effects, defining the ability of central CRF to coordinate behavioral responses independent of, or in synergism with peripheral HPA axis function [87, 100, 101]. Evidently, all of these studies support a role of CRF hyperactivity in stress-related neuropathophysiology. In order to further elucidate the brain regions responsible for mediating the effects of CRF on behavior, a large body of research focused on site-specific CRF administration and pharmacological manipulations [99]. Naturally the involvement of the limbic system was investigated given its modulatory role in emotion, motivation and cognition.

4.1. Limbic System: Hippocampus and Extended Amygdala

The effects of CRF on hippocampal function and integrity with respect to learning and memory were repeatedly investigated in the past. The hippocampus contains scattered CRF-expressing GABAergic interneurons and numerous CRFR1-expressing excitatory pyramidal neurons [59, 102-106]. It is generally proposed that a short-lived increase in CRF facilitates hippocampus-dependent learning and memory (similarly to acute stress), whereas prolonged exposure to elevated CRF impairs cognitive performance [107-109]. However, this is largely based on electrophysiological studies in slice cultures and requires more validation *in vivo*. CRF exerts its effect by potentiating excitatory neurotransmission in the hippocampus, providing direct evidence for an interaction with the glutamatergic system [106, 110-112]. Yet it remains largely unknown how CRF is able to modulate excitatory neurotransmission in the hippocampus considering that it is primarily released from GABAergic interneurons, in which co-release of GABA would presumably induce inhibition. Local diffusion, or long-distance signaling *via* volume transmission has been proposed as a mode of action by which CRF is able to target receptors in the vicinity of its release site [107]. Knowledge about the precise cellular localization of the receptor would advance our understanding in this regard, but has so far been hampered by the lack of specific CRFR1 antibodies [106]. At present, CRFR1 has been found on cell bodies, dendritic shafts and dendritic spines of hippocampal neurons [105, 113, 114]. The adverse effects of chronic CRF release (as they occur during persistent stress) are proposed to result from CRF-induced dendritic spine loss on CRFR1-expressing neurons in the hippocampus [107, 115, 116]. Interestingly, only a subpopulation of co-called "thin spines" seem to be lost following excessive CRF release, but their absence results in profound memory impairments [117]. Along these lines, stress-induced dendritic atrophy has also been linked to excessive release of CRF [114, 117]. Accordingly, CRFR1-antagonists were shown to reduce hippocampus-dependent deficits in memory and synaptic long-term potentiation [117]. At the same time, CRF is required for fear memory formation given that acute injections into the dorsal hippocampus enhance contextual and auditory fear memory [118]. This is also supported by previous ex-

periments with CRF receptor antagonists, which have resulted in fear memory impairments [119, 120]. In addition, CRF injections into the ventral hippocampus were shown to increase anxiety-related behavior in rats [121].

The amygdala plays a prominent role in fear memory acquisition and expression, and modulates aspects of anxiety-related behavior. In rodents, CRF is highly expressed in the central nucleus of the amygdala (CeA), whereas CRFR1 is primarily located in the basolateral amygdala (BLA) [59, 102, 103, 106, 122, 123]. CRF application into the BLA enhances anxiety-related behavior and reduces social interaction in rats [124]. CeA-infusion of CRF receptor antagonists ameliorates stress-induced anxiety and freezing behavior [91, 125], which is likely due to blockage of receptors in the BLA caused by diffusion of the antagonists. Similarly, intra-BLA administration of antalarmin, a CRFR1-antagonist, counteracts social defeat-induced defensive behavior in mice [126]. Along these lines, viral-mediated knockdown of CRFR1 in the BLA of mice mimicked the anxiolytic effect of environmental enrichment [127]. Another CRF-expressing brain region which has attracted increasing attention in the recent years is the BNST. Often referred to as the extended amygdala, the BNST is heavily innervated by the amygdala and projects to the PVN and brainstem monoaminergic nuclei including the locus coeruleus and ventral tegmental area [128-133]. Recent optogenetic studies in mice have clearly implicated the BNST in the modulation of anxiety [134, 135]. Consequently, some of these effects might be modulated *via* CRF, but so far only few studies have investigated the role of BNST-CRF neurons in emotional behavior. Microinfusion of CRF into the BNST enhanced the startle amplitude, and retention in an inhibitory avoidance task in rats [136, 137]. Similarly, intra-BNST administration of CRF in rats elicited a dose-dependent increase in anxiety-related behavior, which could be reversed upon CRFR1 antagonist treatment [138]. In addition, CRF in the BNST has repeatedly been related to addiction, more specifically stress-induced relapse [139, 140]. This is interesting considering that 30-40% of individuals suffering from addictive disorders have a comorbid mood or anxiety disorder [141]. Intra-BNST injection of CRF in rats can induce reinstatement (relapse to drug-seeking behavior in animals), which can be blocked by application of CRF antagonists [142, 143]. Numerous studies have consequently linked the CRF-pathway to dopaminergic signaling, which is primarily involved in addiction-related processes [139, 140, 144-147]. Direct and indirect mechanisms were proposed by which CRF is able to enhance dopaminergic firing in order to drive stress- or cue-induced drug- and alcohol seeking behaviors.

4.2. Modulation of Catecholamines: Ventral Tegmental Area (VTA) and the Locus Coeruleus (LC)

CRF-dopamine interactions in the context of stress and addiction have been extensively examined [144, 148, 149]. The CRF/CRFR1 system has been repeatedly associated with stress-induced drug reinforcement, where it acts to facilitate relapse and increase anxiety during acute and chronic withdrawal [139]. Accordingly, CRF was shown to increase dopamine neuron firing and dopamine release [106, 150-155], although opposite results were also observed depending on the study-design and the investigated release sites [150, 151].

A recent study by Lemos and colleagues demonstrated that CRF acts in the nucleus accumbens (NAc) to increase dopamine release and promote appetitive behavior in mice; an effect which is lost following previous stress exposure [150]. This suggests that CRF might differentially affect the reward circuitry under basal and stress conditions. CRFR1 is expressed in dopaminergic neurons of the VTA and substantia nigra pars compacta [106] while the source of CRF is believed to derive from axons that originate in the forebrain [147, 150, 156-160]. However, Grieder and colleagues recently identified a subpopulation of VTA dopaminergic neurons that express CRF, which is upregulated following chronic nicotine exposure [161]. The ability of CRF to modulate dopaminergic neurotransmission has recently prompted investigations in other behavioral domains, revealing an important role for CRF/CRFR1-dopamine interactions in the context of anxiety and social behavior [106, 162].

CRF is also viewed as a potential mediator of stress-elicited locus coeruleus (LC) activation, the brain's major noradrenergic nucleus [163, 164]. The LC sends noradrenergic projections throughout the brain, including the brainstem, cortical, limbic and hypothalamic structures, and is consequently able to modulate various behavioral, endocrine and autonomic responses [164-167]. Dysregulated noradrenergic circuits *via* excessive CRF have been proposed to underlie pathological hyperarousal observed in numerous stress-related psychiatric disorders [168-170]. CRF is able to induce LC neuronal firing, which is believed to modulate behavioral arousal and attention during stressful situations [163, 164, 171]. Hence, CRF not only facilitates activation of glutamatergic neurotransmission in the hippocampus, and dopaminergic firing in the VTA, but also noradrenergic firing in the LC. In addition, CRF can indirectly regulate endocrine responses *via* activation of the noradrenergic system, which in turn regulates components of the HPA axis [164, 169, 172, 173]. CRF-immunoreactive fibers innervate the LC [174], although unequivocal evidence for the presence of CRFR1 or CRFR2 in the LC is still lacking, in particular in mice [60, 164]. Moreover, the sources of CRF afferents to the LC which modulate specific behavioral effects have not been clearly identified.

5. NEUROMODULATORY EFFECTS OF CENTRAL UCN/CRFR2-SIGNALING

Whereas the role of CRF/CRFR1 in the modulation of HPA axis activity, stress-induced behavior and cognitive functions is well established, the role of CRFR2 and the urocortins still remains controversial. Although still debated, it is postulated that CRF/CRFR1 signaling mediates the initial reaction to stress, whereas UCN/CRFR2 activation controls the later adaptive phase [51, 88, 175]. UCN1 neurons are mainly localized in the Edinger Westphal (EW) nucleus where they constitute the centrally-projecting part of the nucleus (EWcp). UCN1 neurons are recruited following chronic stress exposure, and stay active for a prolonged period of time, suggesting that this peptide plays a prominent role in the later adaptive phase of the stress response [176-178]. Evidence for a possible role of UCN1 in the regulation of mood comes from enhanced UCN1 expression in the EWcp of depressed male suicide victims [53]. Similar to the *i.c.v.* administration of CRF, UCN1 application was also

shown to result in behavioral responses similar to those observed in stressed animals [99, 179], which include increased grooming, locomotion and anxiety-related behavior [180, 181]. However, it is often not clear whether these effects result from CRFR1 or CRFR2 activation. Contradictory is also the observation that UCN1 knockout mice display enhanced anxiety-related behavior (discussed later on). The behavioral outcomes of central UCN2 or UCN3 application have yielded contrasting results depending on the dose and site of application. Previous studies have demonstrated anxiolytic-like effects of CRFR2 activation following i.c.v. administration of UCN2 [182] or UCN3 [183-185] in mice and rats. Surprisingly, CRFR2 antagonism with antisauvagine-30 was also shown to produce anxiolytic responses in rats [186]. The interpretation of these results is further complicated by additional studies reporting no changes in behavioral arousal and anxiety, while others revealed anxiogenic effects of UCN2 administration or CRFR2 activation in the LS [187, 188] and dorsal raphe nucleus [189]. Consequently, CRFR2-activation might exert differential behavioral and neuroendocrine effects depending on the brain region and experienced stress conditions.

5.1. Modulation of Serotonergic System: Raphe Nuclei

A growing body of literature is implicating CRFR2 activation in the modulation of the serotonergic system, although some of the effects are not solely attributable to the UCNs, but also to CRF [190-196]. CRFR2 is abundant in the mid-brain raphe nuclei [60, 69], where it regulates firing rates of serotonin (5-HT) neurons and 5-HT release in efferent stress-related nuclei of the forebrain [191, 193-195]. I.c.v. administration of UCN2 in mice induces enhanced c-fos immunoreactivity in serotonergic neurons of the DRN [193, 197]. Site-specific injection of UCN2 into the DRN was shown to increase c-fos expression in a subpopulation of serotonergic neurons [198], induce 5-HT release in the BLA [190] and potentiate conditioned fear and escape deficits in a model of learned helplessness [189]. On the other hand, injection of the CRFR2 antagonist, antisauvagine-30, resulted in anxiolytic effects [186] including reversal of the potentiation of conditioned fear and the escape deficits following exposure to inescapable stress [189]. These effects were not observed upon CRFR1-antagonist application, suggesting that CRFR2 on serotonergic neurons in the DRN is conveying anxiety in response to uncontrollable stress. However, the identity and expression sites of CRF/UCN-producing neurons targeting CRFR2 in the DRN remain largely elusive.

5.2. Modulation of Energy Balance and Feeding Behavior

Chronic stress has repeatedly been associated with altered metabolic function, in both animal and human studies. Along these lines, growing evidence indicates that mediators of the stress response represent a key locus for gene-environment interactions in the shared biology of depression and obesity [199]. It is well documented that members of the CRF family, and more prominently the UCNs, are capable of suppressing food intake and altering energy expenditure following central or peripheral administration [183, 200-205]. Employing specific pharmacological and genetic tools, a number of studies have established a predominant role for

CRFR2 in mediating the anorexigenic effects of CRF and UCNs [43, 206]. The major CRFR2-expressing brain structures identified to play a role in CRFR2-mediated anorexigenic responses are the ventromedial hypothalamus (VMH), LS, PVN, medial amygdala and DRN [207-212]. CRFR2 signaling pathways in the VMH are starting to gain increasing attention, which is not surprising considering the significant role of this brain region in metabolic regulation. Microinjections of UCN3 into the PVN or VMH were shown to induce satiety [207, 213], and site-specific knockdown of CRFR2 in the VMH resulted in enhanced food intake under basal conditions and following food-deprivation [214]. These results are in part supported by CRFR2 knockout mice, which display increased nocturnal food intake of normal chow [215] and consumed more high fat food compared to littermate controls [216]. In support of this, mice lacking UCN3 showed elevated basal feeding and increased nocturnal food intake after overnight fasting [214], while mice overexpressing UCN3 show leaner body composition and are protected against diet-induced obesity and hyperglycaemia. On the other hand, UCN1 and UCN2 knockout mice display normal spontaneous food intake, which could be the result of a functional compensation by the other two family members, CRF and UCN3 [217, 218]. However, the mechanisms underlying CRFR2/UCN-induced anorexia are not fully understood. Several concepts have been proposed including suppression of gastric emptying and induction of hyperglycemia by CRF/UCNs, as well as direct effects on ghrelin and leptin, the main orexigenic and anorectic peptides in the brain [43, 206]. In addition, CRFR1, CRFR2, UCN1 and UCN3 were shown to be expressed in human adipose tissue, which might indicate a direct effect on fat cell function in addition to the central effects on weight regulation [219]. More recent work has additionally demonstrated that UCN2 and UCN3 also act as autocrine and/or paracrine regulators of glucose homeostasis in the periphery by modulating insulin sensitivity in skeletal muscles or by regulating glucose-induced insulin secretion in beta-cells of the pancreas, respectively [43, 206]. UCN3 is expressed by both beta and alpha cells in human islets [44, 220, 221] and was shown to be markedly depleted in human diabetic islets [222]. Together, CRFR2 and the UCNs help to maintain energy homeostasis in the presence of diverse stressors, through numerous adaptive responses in both the CNS and peripheral tissues [43].

6. GENETICALLY ENGINEERED MICE TARGETING CRF-FAMILY MEMBERS AND THEIR RECEPTORS

Although pharmacological studies have provided valuable insights into the function of the CRF/UCN system, they also face certain limitations. Comparability amongst many of the studies is difficult due to the use of mouse *versus* rat models. More importantly however, is the fact that they assessed effects of acute or repeated administration of exogenous peptides which might not necessarily mimic normal patterns of endogenous CRF/UCN1-3 release and signaling. Moreover, these experiments provide little insight into the outcomes of long-lasting CRF system dysregulations as they might occur in stress-related mood and anxiety disorders. In addition, it is not always clear whether the effects of centrally administered CRF/UCN1 were mediated by CRFR1 or CRFR2. Although some studies applied CRFR1 and CRFR2

antagonists to tackle this question, many of these compounds are not fully discriminative at the applied concentrations. The generation of transgenic mice, overexpressing or lacking different CRF-family members, has provided valuable insights into the involvement of the CRF/UCN system in stress-related physiology and behavior (Table 1).

6.1. CRF Overexpressing Mice

In order to study the role of chronic CRF hyperdrive in the context of mood and anxiety-disorders, independent models of CRF excess were developed (Table 1). The first CRF overexpressing mouse line was generated *via* classical pronuclear injection applying the broadly active metallothionein 1 (*Mt1*) promoter [223]. These mice (CRF-OE^{*Mt1*}) showed strong CRF overexpression in the brain and peripheral organs including lung, adrenal, heart, and testis. CRF overproduction resulted in elevated plasma corticosterone levels and Cushing's-like symptoms. Of note: the *Mt1* promoter harbors a GC response element and thus might react in a feed-forward manner upon elevated corticosterone levels. CRF-OE^{*Mt1*} mice showed increased anxiety-related behavior, which was reversible with the CRF receptor antagonist α -helical CRF [224]. Moreover, these mice displayed deficits in learning, decreased immobility in the forced swim test (FST) and reduced attention [108, 225]. Another CRF overexpressing mouse line was developed using the *Thy1.2* promoter driving CRF expression in postnatal and adult neurons of the brain [226]. However, CRH-OE^{*Thy1.2*} mice did not show an altered stress response or phenotype indicative of changes in anxiety behaviors [226, 227]. Instead, CRH-OE^{*Thy1.2*} mice displayed reduced startle reactivity as well as reduced freezing following fear conditioning [226, 227]. With some delay CRH-OE^{*Thy1.2*} mice also developed a mild cushingoid phenotype [228]. Finally, different conditional CRF-overexpressing mouse lines have been established in recent years. Two studies applied the “*tet-on/tet-off*” system, which allows for reversible and inducible overexpression of CRF [229, 230]. Although both made use of the forebrain-specific *Camk2a* promoter combined with a tet-operator driven CRF-construct, the behavioral and neuroendocrine consequences of CRF excess were rather specific for each mouse strain (Table 1). Taken together, these examples illustrate the difficulties of comparing results from different transgenic mouse lines even if they are based on similar constructs. A mouse model which permits conditional CRF overexpression avoiding common uncertainties of classical transgenesis, such as unpredictable influences of the site of transgene insertion and the number of inserted transgene copies, is the CRH-COE mouse line [231]. This mouse line was generated by introducing a CRF expression unit into the ubiquitously expressed *Rosa26* (*R26*) locus. Expression of exogenous CRF driven by the *R26* promoter is prevented unless a *loxP* flanked transcriptional terminator is deleted *via* a Cre recombinase, which determines the spatial and temporal pattern of CRF overexpression. A similar strategy was recently applied to generate a model which enables conditional overexpression of UCN2 [232]. This model of CRF overexpression demonstrated that CNS-restricted CRF overexpression in CRH-COE^{*CNS*} mice, achieved by breeding to *Nestin-Cre* mice, leads to HPA axis hyperactivity, increased active stress-coping behavior in the FST and altered sleep regulation [231, 233]. Importantly, reduced immobility in the FST was not observed in mice overex-

pressing CRF specifically in forebrain *Camk2a*-positive or forebrain GABAergic neurons, suggesting an involvement of hindbrain-regions in CRF-induced active-stress coping behavior. The same model was consequently used to induce CRF overexpression specifically in the forebrain during adulthood by breeding CRH-COE mice to the tamoxifen-inducible *Camk2a-CreERT2* driver line (CRH-COE^{*CamkCreERT2*}). Behavioral assessment of these animals revealed an anxiogenic phenotype, supporting earlier findings in conditional CRFR1 knockout mice that anxiety-related behavior is regulated by forebrain CRF/CRFR1 during adulthood [106]. In order to discriminate the direct effects of centrally hypersecreted CRF from those resulting from HPA axis activation, two additional conditional CRF-overexpressing mouse lines were created [234]. CRH-COE^{*Del*} mice overexpress CRF in a ubiquitous manner, while CRH-COE^{*Ant*} mice selectively overexpress CRF in the anterior and intermediate lobes of the pituitary. Both mouse lines displayed increased basal plasma corticosterone levels and consequently signs of Cushing's syndrome. However, alterations in anxiety were only observed upon ubiquitous CRF overexpression, suggesting that chronic hypercorticosteroidism alone is not sufficient to alter emotional behavior. The study implies that central CRF hyperdrive on its own or in combination with elevated GCs is responsible for the observed behavioral alterations in CRH-COE^{*Del*} mice [234]. With respect to clinical findings and in order to fully understand the effects of CRF hyperdrive in the context of stress-related neuropathologies, the generation of mice overexpressing CRF under its endogenous promoter represents a matter of particular importance. As discussed later on, this goal could ultimately be achieved by breeding conditional CRF overexpressing mice with the recently generated CRF-Cre driver lines. Along these lines, a recent study utilized a mouse N-ethyl-N-nitrosourea (ENU)-screen, and identified a point mutation in the CRF promoter region, which results in a gain-of-function mutation, *i.e.* CRF overexpression and consequently development of Cushing's syndrome [235].

6.2. CRF Knockout Mice

Although CRF overexpressing mice represent valuable disease models with respect to chronic CRF and HPA axis hyperactivity, they are confounded by ectopic expression in non-endogenous brain regions/neurons and peripheral organs. Consequently a loss-of-function approach is more likely to reveal physiologically relevant effects of CRF on behavior. The development of constitutive CRF knockout mice (CRH-KO) by Muglia and colleagues has been important in addressing this issue [236]. CRH-KO mice displayed severely reduced plasma corticosterone levels indicative of blunted HPA axis activity (Table 1). Importantly, this study revealed fetal GC requirement for lung maturation, which was severely impaired in CRF-deficient mice obtained from homozygous breedings [236]. Surprisingly, CRH-KO mice displayed no gross alterations in emotional behavior and CRFR1 antagonists were still able to exert an anxiolytic effect in these animals [237]. The discrepancy between constitutive CRF and CRFR1 null mutants (discussed below) with respect to behavioral outcomes could be due to a number of reasons: 1) UCN1, the only other CRFR1 ligand, might compensate for the loss of CRF; 2) early deletion of CRF might trigger general compensatory processes; 3) corticosis-

terone deficiency might mask potential phenotypes; 4) CRF might exert its action primarily under conditions of chronic or severe stress; 5) CRFR1 might comprise ligand-independent activity, e.g. due to constitutive activity or heteromerization with other receptors; 6) and last but not least, it might suggest the presence of a yet unidentified CRFR1-ligand. The generation of conditional CRF knockout mice would significantly help to shed light on some of these issues.

6.3. CRFR1 Knockout Mice

The contribution of CRFR1 to the modulation of stress-related behaviors was addressed by conventional and conditional CRFR1 knockout mice (Table 1). Expectedly, CRFR1 null mice exhibited a chronic GC deficiency due to disrupted HPA axis activity, which was observed in two independently generated CRFR1-deficient mouse lines; *CRHR1-KO* and *CRFR1-KO* [238, 239]. Furthermore, both mouse lines displayed reduced anxiety-related behavior [238-240]. In order to exclude the possibility that the GC deficit is mediating the observed behavioral effects, a conditional forebrain-specific CRFR1 knockout mouse line (*Crhr1^{FB-CKO}*) was generated [241]. In this mouse model, Cre-mediated deletion of CRFR1 is initiated in the second week of postnatal life, and is primarily restricted to cortical and limbic forebrain regions including the amygdala, hippocampus, BNST, but not the anterior pituitary. *Crhr1^{FB-CKO}* mice displayed reduced anxiety-related behavior and normal GC levels under basal conditions, supporting the notion that limbic CRFR1 can regulate emotional behavior independent of HPA axis alterations [241]. However, corticosterone levels were slightly elevated in *Crhr1^{FB-CKO}* mice 30 and 90 min following a 5 min restraint stress, suggesting that limbic CRFR1 itself is partially involved in HPA axis feedback regulation [241]. Both *CRHR1-KO* and *Crhr1^{FB-CKO}* mice displayed impairments in remote fear memory consolidation, suggesting that cognitive processes are also mediated by CRFR1 in forebrain cortical and limbic structures [120]. In addition, more recent work demonstrated that forebrain CRFR1 deficiency prevents cognitive deficits induced by early-life stress and chronic stress during adulthood [242-244]. Moreover, stress-induced dendritic remodeling and spine loss was attenuated in *Crhr1^{FB-CKO}* mice [242, 244]. Although all these studies clearly implicate CRFR1 in the modulation of emotional and cognitive responses, a fundamental question remained unsolved: Which are the underlying neurotransmitter circuits controlled by CRFR1 that modulate anxiety-related behavior? In order to address this question the neurochemical identity of CRFR1-expressing neurons was established by sensitive neurochemical methods and genetic tools [106]. CRFR1 expression was demonstrated in forebrain glutamatergic and GABAergic neurons as well as in midbrain dopaminergic neurons, and in a few serotonergic cells of the dorsal and median raphe nucleus [106]. In order to dissect the underlying neurotransmitter circuits, previously generated floxed *CRFR1* mutants [241] were bred to a set of neurotransmitter-specific Cre-driver lines resulting in selective deletion of the receptor from glutamatergic (*Crhr1^{Glu-CKO}*), GABAergic (*Crhr1^{GABA-CKO}*), dopaminergic (*Crhr1^{DA-CKO}*) and serotoner-

gic (*Crhr1^{5-HT-CKO}*) neurons, respectively (Fig. 4). Selective deletion of CRFR1 in forebrain glutamatergic circuits reduced anxiety, which is in agreement with the previously established phenotype of forebrain-specific CRFR1 knockout mice, and the anxiolytic properties of CRFR1 antagonists. Moreover, CRF-induced changes in glutamatergic neurotransmission in the amygdala and hippocampus were absent in *Crhr1^{Glu-CKO}* mice [106, 112]. Remarkably, specific deletion from midbrain dopaminergic neurons enhanced anxiety-related behavior, highlighting a previously unrecognized negative effect of CRFR1-ablation on emotional behavior. Importantly, this anxiety-inducing effect was associated with reduced dopamine release in the prefrontal cortex, establishing the relevance of dopamine-CRF interactions in behaviors other than addiction [106]. These results defined a novel bidirectional role of CRFR1 in anxiety, suggesting that CRF/CRFR1-controlled glutamatergic and dopaminergic systems might function in a concerted but antagonist manner to keep emotional responses to stressful situations in balance. This was further supported by the absence of an anxiety phenotype in *Crhr1^{CNS-CKO}* animals, in which the CRFR1 is absent from both neurotransmitter systems [106]. HPA axis activity, FST behavior and auditory fear condition were not differently affected in the investigated mouse lines, suggesting that the bidirectional role of CRFR1 might be specific for anxiety-related behavior. However, additional cognitive parameters and behavioral domains need to be investigated in the future to further substantiate the selectivity of the observed effects. An anxiolytic role for CRFR1 was also demonstrated shortly afterwards in the globus pallidus, a central component of the basal ganglia circuitry which is indirectly controlled by dopaminergic substantia nigra-striatal projections [245]. As previously mentioned, Lemos and colleagues recently demonstrated a novel appetitive effect of CRF in the nucleus accumbens under basal conditions resulting from CRF's ability to positively regulate dopamine release [150]. However, severe stress was shown to induce a persistent dysregulation of CRF-dopamine interactions that normally produce a positive affective state, resulting in an aversive phenotype. Whether stress is able to induce a switch in CRF-neurotransmitter interactions in the conditional CRFR1 mutants analyzed by Refojo and colleagues remains to be investigated. An important insight from all three studies is that we can no longer regard CRF as a generally "aversive" stress neuropeptide.

6.4. CRFR2 Knockout Mice

In contrast to the consistent and reproducible phenotype of CRFR1 knockout mice, a number of discrepancies have been observed in CRFR2 knockout mice (Table 1). Until now, three conventional CRFR2 knockout mouse models have been generated [246-248]. Two studies reported enhanced ACTH and corticosterone release in response to stress, but an early termination of ACTH release, suggesting that CRFR2 is involved in maintaining HPA drive [246, 247]. In addition, Coste and colleagues observed an overall blunted corticosterone recovery in *CRHR2-KO* mice, implying an involvement in HPA feedback function. The effects of

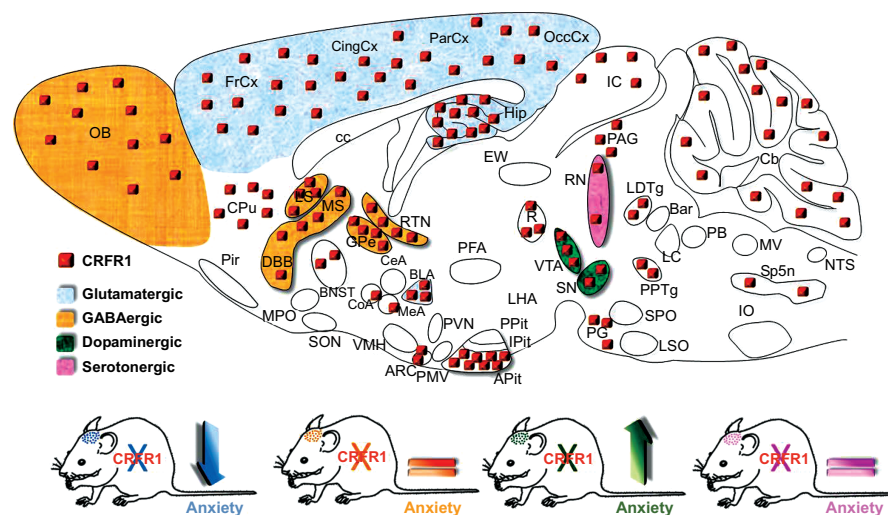


Fig. (4). CRFR1 modulates anxiety-related behavior in a bidirectional manner. CRFR1 is expressed in diverse neuronal subpopulations. Selective deletion of the receptor in glutamatergic neurons reduces anxiety-related behavior while deletion in dopaminergic neurons produces the opposite effect [106].

Table 1. Summary of genetic mouse models targeting CRF family members and their receptors.

Transgenic Line	Targeting Strategy	CNS-Related Phenotype	References
CRF overexpression			
<i>CRF-OE^{Mt}</i> Developmental / ubiquitous	Non-selective OE of rat CRF under murine metallothionein promoter	Cushing-like phenotype (↑ ACTH & CORT levels, = CORT stress response), adrenal hypertrophy, ↓ general locomotion, ↑ anxiety, ↓ immobility FST, deficits in learning and spatial memory	[108, 223-225]
<i>CRH-OE^{Thy1.2}</i> Developmental	OE of rat CRF under <i>Thy-1</i> promoter	Cushing-like phenotype at 6 months (↑ CORT levels, marginal increase in ACTH, = CORT stress response), adrenal hypertrophy, nonsuppression of DXM, ↓ locomotion, ↓ startle reactivity & habituation, ↓ PPI, = anxiety	[227, 228]
<i>CRH-COE^{CNS}</i> Developmental / CNS-specific	Conditional <i>Nestin-Cre</i> induced OE of murine CRF driven by the <i>Rosa26</i> -promoter	Stress-induced hypersecretion of CORT, = basal CORT levels, ↓ immobility FST/TST, ↑ REM sleep & slightly suppressed non-REM sleep	[231, 233]
<i>CRH-COE^{FB}</i> Postnatal / forebrain-specific	Conditional <i>Camk2α-Cre</i> induced OE of murine CRF driven by the <i>Rosa26</i> -promoter	= basal HPA axis & CORT stress response, = immobility FST, ↑ REM sleep	[231, 233]
<i>CRH-COE^{GABA}</i> Developmental / restricted to GABAergic neurons	Conditional <i>Dlx5/6-Cre</i> induced OE of murine CRF driven by the <i>Rosa26</i> -promoter	= basal HPA axis & CORT stress response, = immobility FST	[231, 233]
<i>FB-CRH</i> (<i>Camk2α-rTA/tetO-Crf</i>) Inducible forebrain-specific	<i>Camk2α-Cre</i> mediated COE of rat CRF driven by the CMV promoter (<i>tet</i> -on system; DOX induces expression)	DOX administration at P56 for 3 weeks, ↑ CORT at circadian nadir, = ACTH levels & CORT stress response, ↓ thymus size, = dexamethason suppression	[229]
<i>FBCRHOE</i> (<i>Camk2α-rTA/tetop-CRH</i>) Inducible forebrain-specific	<i>Camk2α-Cre</i> mediated COE of CRF driven by the CMV promoter (<i>tet</i> -off system; DOX represses expression)	Early life forebrain CRF OE (off DOX E15-P21) causes ↑ CORT levels only during development & long-lasting anxiogenic & despair-like alterations; Lifetime CRF OE induces Cushing-like phenotype at 8 weeks & ↑ CORT & ACTH levels only at circadian nadir	[230]

Transgenic Line	Targeting Strategy	CNS-Related Phenotype	References
<i>CRH-COE^{CamkCreERT2}</i> Inducible forebrain-specific	Conditional <i>Camk2α-CreERT2</i> induced OE of murine CRF driven by the <i>Rosa26</i> -promoter; tamoxifen induces expression	Tamoxifen administration at P 56, ↑ anxiety, = locomotion	[106]
<i>CRH-COE^{Del}</i> Developmental /ubiquitous	Conditional <i>Deleter-Cre</i> induced OE of murine CRF driven by the <i>Rosa26</i> -promoter	Cushing-like phenotype at 3 weeks, ↑ body weight, adrenal hypertrophy, ↑ CORT levels at circadian peak & trough, = CORT stress response, = locomotion, ↑ anxiety, ↓ immobility FST	[234]
<i>CRF-COE^{Api}</i> Developmental /anterior pituitary-specific	Conditional <i>Pomc-Cre</i> induced OE of murine CRF driven by the <i>Rosa26</i> -promoter	Mild Cushing-like phenotype at 5-6 months, ↓ body weight, adrenal hypertrophy, thymus atrophy, ↑ CORT levels at circadian trough, = CORT levels at circadian peak, = CORT stress response (↓ in females), = locomotion, = anxiety, slightly ↓ immobility FST, ↑ NREM sleep	[234]
<i>Crh^{-120/+}</i> Developmental / ubiquitous	ENU-induced gain-of-function mutation in the CRF promoter region	Cushing-like phenotype including obesity, muscle wasting, thin skin, hair loss, ↑ CORT levels, hyperglycemia, hyperinsulinemia	[235]
UCN2 overexpression			
<i>UCN2-COE</i> Developmental / conditional	<i>SF-1-Cre</i> induced OE of murine UCN2 driven by the <i>Rosa26</i> promoter	Down-regulation of adrenal and ovarian steroidogenesis	[232]
UCN3 overexpression			
<i>UCN3OE</i> Developmental / ubiquitous	Constitutive OE of mouse UCN3 under the <i>Rosa26</i> promoter	Leaner body composition, protected against diet-induced obesity and hypoglycaemia, = basal & response CORT levels, ↓ ACTH response to acute stress, ↑ anxiety, ↓ spatial memory following restraint stress, stress-induced alterations in tissue serotonin levels in the DRN	[196, 258]
CRF knockout			
<i>CRH-KO</i> Developmental / constitutive	Replacement of the CRF coding region with a neomycin cassette	Blunted HPA axis activity (↓ basal & stress-induced CORT levels), = anxiety, = anxiety post restraint stress, = locomotion, = ASR & learning	[236, 237]
CRFR1 knockout			
<i>CRHR1-KO</i> Developmental / constitutive	Replacement of exons 8-13 with a neomycin cassette	Blunted HPA axis activity (↓ basal & stress-induced CORT levels), ↓ anxiety, ↑ locomotion, ↑ & delayed stress-induced alcohol intake, ↓ remote fear memory consolidation	[120, 238]
<i>CRFR1-KO</i> Developmental / constitutive	Replacement of exons 5-8 with a neomycin cassette	Blunted HPA axis activity (↓ basal and stress-induced CORT levels), ↓ anxiety, ↓ spatial memory performance	[239, 240]
<i>Crhr1^{FB-CKO}</i> Postnatal / forebrain-specific inactivation	Conditional <i>Camk2α-Cre</i> mediated CRFR1 inactivation (floxed exons 9-13)	= basal HPA axis activity, slightly enhanced CORT after acute stress, ↓ anxiety, ↓ chronic-stress induced cognitive deficits, ↓ dendritic atrophy & spine loss, ↓ remote fear memory consolidation	[120, 241-244]

Table (1) contd....

Transgenic Line	Targeting Strategy	CNS-Related Phenotype	References
<i>Crhr1^{Glu-CKO}</i> Developmental / inactivation in glutamatergic neurons	Conditional <i>Nex</i> -Cre mediated CRFR1 inactivation (floxed exons 9-13)	= basal & stress-induced HPA activity, ↑ locomotion, ↓ anxiety, = immobility FST, = auditory fear conditioning, impaired glutamatergic neurotransmission in the amygdala and hippocampus	[106, 112]
<i>Crhr1^{GABA-CKO}</i> Developmental / inactivation in GABAergic neurons	Conditional <i>Dlx5/6</i> -Cre mediated CRFR1 inactivation (floxed exons 9-13)	= basal & stress-induced HPA activity, = locomotion, = anxiety, = immobility FST, = auditory fear conditioning, = CRF induced neuronal excitability in the hippocampus	[106, 112]
<i>Crhr1^{5HT-CKO}</i> Developmental / inactivation in serotonergic neurons	Conditional <i>ePet</i> -Cre mediated CRFR1 inactivation (floxed exons 9-13)	= basal & stress-induced HPA activity, = locomotion, = anxiety, = immobility FST, = auditory fear conditioning	[106]
<i>Crhr1^{DA-CKO}</i> Inducible / inactivation in dopaminergic neurons	Conditional <i>DAT</i> -CreERT2 mediated CRFR1 inactivation (floxed exons 9-13), tamoxifen induces knockout	Tamoxifen administration at P56, = basal & stress-induced HPA activity, ↓ locomotion, ↑ anxiety, = immobility FST, = auditory fear conditioning, ↓ footshock-induced dopamine release in the prefrontal cortex	[106]
<i>Crhr1^{CNS-CKO}</i> Developmental / CNS-specific	Conditional <i>Nes</i> -Cre mediated CRFR1 inactivation (floxed exons 9-13)	= locomotion & anxiety	[106]
<i>Crhr1^{ΔEgfp}</i> Developmental CRFR1 reporter allele	Knockin of EGFP into exon 2 resulting in a null allele	Designed to visualize 1CRFR1-expression, with the ability to conditionally restore expression of a GFP tagged full- length CRFR1, which can be deleted <i>via</i> Cre recombinase, blunted HPA axis activity (↓ basal & stress-induced CORT levels)	[106]
<i>Crhr1^{ΔZ}</i> Conditional multifunctional CRFR1 allele	Knockin of tau-LacZ (tZ) into endogenous CRFR1 locus resulting in a null allele	Designed to genetically label CRFR1-expressing cells with the ability to conditionally restore or delete CRFR1 with Flp and Cre recombinase, <i>Crhr1^{ΔZ}</i> reporter mice revealed novel aspects of <i>CRFR1</i> expression, blunted HPA axis activity (↓ basal & stress-induced CORT levels)	[59]
CRFR2 knockout			
<i>CRHR2-KO</i> Developmental / constitutive	Replacement of transmembrane domains 3-5 with a neomycin cassette	↓ ACTH & CORT response to stress & early termination of ACTH release, = anxiety, ↑ social discrimination	[42, 246]
<i>CRFR2-KO</i> Developmental / constitutive	Replacement of exons 10-12 with a neomycin cassette	↓ ACTH & CORT response to stress & early termination of ACTH release, ↑ anxiety & immobility FST, ↑ basal & high-fat diet food consumption	[215, 216, 247]
<i>CRHR2-null</i> Developmental / constitutive	Replacement of 3 rd cytoplasmic region with a neomycin cassette	= HPA axis activity, = locomotion, ↑ anxiety & immobility FST	[248]

Table (1) contd....

Transgenic Line	Targeting Strategy	CNS-Related Phenotype	References
CRFR1/CRFR2 double knockout			
<i>CRHR1/CRHR2 dKO</i> Developmental / constitutive	Crossbreeding	↓ HPA stress response	[251]
<i>CRFR1/CRFR2 dKO</i> Developmental / constitutive	Crossbreeding	↓ HPA stress response, ↓ anxiety only in females	[250]
UCN1 knockout			
<i>UCN1-KO</i> Developmental / constitutive	Replacement of coding region with neomycin cassette	= HPA axis and feeding, ↑ anxiety, impaired hearing	[217]
<i>UCN1-null</i> Developmental / constitutive	Replacement of exon 2 with eGFP-LacZ reporter cassette	= HPA axis, = locomotion & anxiety, ↓ impaired ASR	[252]
<i>UCN1^{-/-}</i> Developmental / constitutive	Replacement of exon 2 with a neomycin cassette	= basal HPA axis, ↓ HPA adaptation to repeated restraint stress	[253]
UCN2 knockout			
<i>UCN2-KO</i> Developmental / constitutive	Replacement of exon 2 with a neomycin cassette	↑ nocturnal ACTH & CORT levels, ↓ FST immobility only in females, = anxiety & locomotion & fear conditioning in males and females	[218]
<i>Ucn2^{tz/tz}</i> Developmental / constitutive	Replacement of open reading frame with <i>tau-LacZ</i> reporter cassette	= HPA axis, = anxiety, = immobility FST, = social discrimination, ↓ aggressiveness	[42, 254]
UCN3 knockout			
<i>UCN3-KO</i> Developmental / constitutive	Replacement of coding region with a neomycin cassette	↓ glucose-induced insulin secretion, ↓ basal glucose and insulin secretion under high-fat diet, ↑ basal feeding & following food deprivation, ↑ ethanol intake & preference	[256, 257]
<i>Ucn3^{tz/tz}</i> Developmental / constitutive	Replacement of open reading frame with <i>tau-LacZ</i> reporter cassette	= HPA axis, = anxiety & immobility FST, ↑ social discrimination	[42]
UCN1/UCN2 double knockout			
<i>UCN1-KO/UCN2-KO</i> Developmental / constitutive	Crossbreeding	↑ stress-induced HPA response only in males, ↓ anxiety	[192]
UCN1/UCN2/UCN3 triple knockout			
<i>UCN1-KO/UCN2-KO/UCN3-KO</i> Developmental / constitutive	Crossbreeding	= HPA activity, ↓ locomotion, ↑ anxiety 24h after acute stress but not under basal conditions	[255]

Overexpressing mouse lines were either generated *via* classical pronuclear DNA injection or *via* targeted insertion in embryonic stem cells. All knockout lines were generated by means of targeted deletion *via* homologous recombination in embryonic stem cells. Nomenclature of mouse lines is based on the first publication or laboratory of origin. ↑ indicates an increase, ↓ indicates a decrease, = indicates no difference compared to control animals. Abbreviations: Acoustic startle response (ASR), adrenocorticotropic hormone (ACTH), corticosterone (CORT), dexamethasone (DXE), overexpression (OE), conditional overexpression (COE), doxycycline (DOX), forced swim test (FST), hypothalamic-pituitary-adrenal (HPA) axis. Italics highlight the original publications.

CRFR2 deficiency on anxiety-related behavior are less clear. Whereas Coste and colleagues observed no changes in anxiety-related behavior, Bale *et al.* and Kishimoto *et al.* detected increased anxiety-related behavior. The absence of a baseline anxiety phenotype in *CRFR2-KO* mice was also confirmed in a later study in which it appeared that the constitutive deletion of CRFR2 enhances anxiety 24 h after restraint stress, but not immediately or following prolonged periods of stress [249]. Importantly, Bale and colleagues reported a compensatory upregulation of CRF in the central amygdala of *CRFR2-KO* mice, which might have influenced the anxiogenic phenotype. Two of the studies also reported increased immobility of CRFR2-deficient mice in the FST [247, 248]. In addition, enhanced social discrimination was observed in *CRHR2-KO* mice [42]. Based on the above studies, CRFR2 was initially proposed to exert opposite functions compared to CRFR1, but this simplistic view has been rejected by more recent research. Differences in strain background might underlie some of the discrepancies observed between the mouse lines. In addition different exons were targeted in the individual knockout lines, which leads to the speculation that different truncated splice variants might still be present and affect the phenotypic outcome of the genetic manipulation. Along these lines, the soluble variant of CRFR2 (sCRFR2 α) should be intact in all three *CRFR2-KO* mouse lines. Double CRFR1/CRFR2 knockout mice have also been generated, but only mild behavioral alterations were observed in one of the lines, which were specific for female mice [250, 251]. However, both lines displayed impaired stress-induced HPA axis activation. A more detailed analysis of both double knockout lines, as well as the utilization of respective conditional alleles, might provide additional insights with regards to the behavioral discrepancies. Importantly, assessment of feeding behavior in CRFR2 knockout mice supports a role for the receptor in anorexigenic responses and centrally controlled metabolic functions [215, 216]. The generation of conditional CRFR2 knockout mice will be mandatory in order to uncover the precise involvement of CRFR2 in stress-related behaviors, HPA axis activity, as well as metabolic function.

6.5. Genetically Modified Mice Targeting the Urocortins

Three different UCN1 knockout lines have been independently generated (Table 1); however their phenotypes remain controversial [217, 252, 253]. Vetter and colleagues reported increased anxiety-related behavior in their mouse model, which was not confirmed by the study of Wang *et al.* and so far not investigated in the UCN-deficient mice generated by Zalutskaya and colleagues. Consequently, UCN1 seems an unlikely candidate to compensate for CRF deficiency in *CRH-KO* mice. This is also supported by the restricted expression pattern of UCN1 compared to CRF (Fig. 3). Importantly, all UCN1 knockout lines exhibited normal basal and stress-induced GC levels, supporting the notion that UCN1 plays a minor role in control of HPA axis function. However, Zalutskaya and colleagues observed that corticosterone levels in male UCN1-deficient mice did not adapt to repeated restraint stress. The role of the other two urocortin members has been assessed in UCN2 and UCN3 knockout mice. Female UCN2 knockout mice display mild alterations in their basal circadian rhythm of ACTH and cor-

ticosterone secretion [218]; whereas no differences were observed in basal and stress-induced corticosterone levels in male mice [218, 254]. Male and female *UCN2-KO* mice generated by Chen *et al.* exhibited no alterations in locomotion, anxiety and contextual fear conditioning. However, only female *UCN2-KO* mice displayed reduced immobility in the FST [218]. Male UCN2-deficient mice developed by Deussing and colleagues (*Ucn2^{tz/tz}*) displayed reduced aggressiveness, but showed no changes in anxiety, immobility in the FST and social discrimination [42, 254]. Differences in HPA axis activity and anxiety were also not observed in UCN3-deficient mice (*Ucn3^{tz/tz}*) [42]. However, male and female *Ucn3^{tz/tz}* mice showed enhanced social discrimination abilities, which was also observed in *CRHR2-KO* mice, and attributed to UCN3-expressing neurons of the BNST and medial amygdala - nuclei functionally connected to the accessory olfactory system. These data suggest an involvement of the UCN3/CRFR2-system in social memory.

The relevance of the UCN family members in stress-related behavior was additionally assessed in recently generated *UCN1/UCN2* double and *UCN1/UCN2/UCN3* (*tKO*) triple knockout mice. *UCN1/UCN2* knockout mice displayed no changes in basal HPA axis activity, but exhibited elevated corticosterone levels following acute stress exposure [192]. On the other hand, HPA axis function was indistinguishable in *tKO* mice compared to controls [255]. *UCN1/UCN2* knockout mice displayed decreased anxiety under basal and acute-stress conditions, which was accompanied by elevated serotonin concentrations in a number of brain regions including the DRN, hippocampus, BLA and subiculum [192]. In contrast, *tKO* mice exhibited increased anxiety-like behavior, but only 24 h following restraint-stress. Moreover, *tKO* mice displayed an increased stress-induced startle response [255]. As opposed to *UCN1/UCN2* knockout mice, the behavioral phenotype in *tKO* mice was associated with decreased serotonergic metabolism in regions such as the septum, central and basolateral amygdala [255]. Again, the effect of compensatory changes in CRF expression on emotional behavior cannot be excluded in many of the urocortin mouse models, as shown in *UCN1/UCN2* knockout mice [192]. Overall the majority of data suggests that the urocortins and CRFR2 are able to regulate specific aspects of stress-related emotional behavior complementing effects of CRF and CRFR1. As previously mentioned, more recent studies are starting to implicate the UCNs, especially UCN3, in the modulation of glucose homeostasis and metabolic function. Both, UCN3 knockout mice [256, 257] and UCN3 overexpressing mice [258] display alterations in glucose metabolism, although the results of the former have to be interpreted with caution considering the uncertainties that arise with ectopic gene expression.

7. VIRAL-BASED GAIN- AND LOSS-OF-FUNCTION STUDIES

In addition to gene targeting in mice, viral-mediated gene transfer is another popular method used to study neuronal function in the rodent brain. Adeno-associated viruses (AAVs) and lentiviruses (LVs) are the most commonly used viral vectors to infect adult neurons *in vivo*, which can be achieved *via* stereotactic injections in any given brain-region. AAVs and LVs can be used to *e.g.* express fluores-

cent markers, genes of interest, fusion proteins, Cre recombinases or short-hairpin RNAs (shRNAs) [259-268]. Spatial restriction can be achieved by utilizing cell type-specific promoters to drive gene expression. Consequently, AAVs or LVs represent a versatile tool that can be used for a number of applications including gain- or loss-of-function approaches, tracing studies, or simply to label neurons. For example, delivery of AAVs, expressing the Cre recombinase, into a specific brain region of mice with a floxed gene of interest results in Cre-mediated deletion of the floxed gene. Another possibility is to generate floxed viral constructs (Cre-dependent viral vectors), which are only active upon the presence of Cre recombinase, for example in region- or cell type-specific Cre-driver mouse lines. Tracing and mapping studies of neuronal circuits have also heavily relied on viral-mediated delivery of fluorescent proteins [269-273]. For instance, AAVs can be designed to harbor synaptic proteins fused to fluorescent markers, which will be actively transported to the synapses enabling the visualization of axonal projections. Specific neuronal circuits can be targeted by expressing these “tracers” in a Cre-dependent manner. One important issue is variable transduction efficiencies between different viral serotypes [274-277]. In addition, serotypical variations are also observed with respect to toxicity and immune responses triggered by the viral capsule [278-280]. Nonetheless, the combination of mouse genetics and recombinant AAVs/LVs has greatly improved our ability to map, monitor and manipulate neurons and circuits. For instance optogenetics, one of the latest technological leaps in neuroscience, greatly benefits from these combinatorial approaches. As a result, an increasing number of studies have started to make use of viral-mediated gene transfer to study the CRF system. The most prominent examples are listed in Table 2. As opposed to conventional pharmacological administrations (i.c.v. or microinjection pumps), viral-based methods have greatly improved the spatial and temporal precision of neuropeptide delivery into the CNS. This was effectively demonstrated by a recently described viral-mediated genetic approach for i.c.v. delivery of CRF to the CSF. Applying the choroid plexus-specific CRFR2 promoter, a LV-based system was established which enables a doxycycline-inducible and hence reversible delivery of CRF into the CSF [267]. The induction of CRF overexpression in the choroid plexus resulted in enhanced anxiety-related behavior, which is in accordance with previous findings in mice treated with i.c.v. injections of CRF [reviewed in 99], CRF overexpressing mice [224], and depressed individuals that displayed enhanced CRF levels in the CSF [87]. A number of additional studies investigated the effects of LV-based CRF hyperdrive in the CeA. Keen-Rhinehart and colleagues reported increased despair-like behavior in the FST and an enhanced acoustic startle response, which is indicative of enhanced anxiety [281]. Along these lines, Regev *et al.* demonstrated that short-term inducible overexpression of CRF in the CeA enhances stress-induced effects on anxiety-like behavior [282] however, prolonged overexpression produced the opposite effect [283]. This possibly suggests that acute and chronic CRF hyperactivity might be able to differentially regulate basal and stress-induced emotional behaviors. Regev and colleagues additionally investigated the effects of site-specific CRF overexpression in the BNST, but observed no effects on basal and stress-induced anxiety

[283]. An important drawback of the above mentioned viral studies is the lack of overexpression specificity. In all cases, exogenous CRF expression is driven by ubiquitously active promoters, resulting in transgene expression that extends beyond the cell population of interest – neurons endogenously expressing CRF. Thus the observed effects may partially originate from unintentionally targeted cells. These problems can be circumvented by designing viral vectors with CRF-specific promoters, as demonstrated by two recent studies [284, 285]. Both made use of the same LV-construct which contains a 3.0 kb CRF promoter region, previously utilized for the generation of transgenic CRFp3.0Cre mice [286]. Applying this overexpression model, Sink and colleagues demonstrated that CRF neurons within the lateral BNST modulate conditioned anxiety-like behaviors, thus suggesting that enhanced CRF expression within these neurons may contribute to inappropriate regulation of emotional memories. However, the expression of the exploited CRF promoter does not fully recapitulate the endogenous CRF expression pattern, possibly due to missing regulatory elements, which might be located further up- or downstream of the promoter sequence. In fact, the size restriction of the DNA/RNA which can be incorporated into the desired viruses is a major hurdle when it comes to transgene expression driven from cell type-specific promoters. An alternative strategy is the delivery of Cre-dependent viral vectors into cell type-specific Cre-driver lines, which will be discussed later on.

In addition to the gain-of-function procedures, viral-mediated loss-of-function approaches have also been employed to study aspects of the CRF system (Table 2). LV-mediated knockdown of CRF in the CeA attenuated stress-induced anxiety-like behaviors and altered HPA axis activity, reinforcing a role for amygdalar CRF in the modulation of fear and anxiety [282]. In support, LV-based knockdown of CRFR1 in the BLA decreases anxiety-related behavior [127]. Elliott and colleagues targeted CRF in the PVN *via* stereotactic delivery of LV-constructs, carrying short hairpin RNAs (shRNA) against CRF. The knockdown attenuated chronic stress-induced social avoidance, which was shown to result from demethylation of the CRF promoter [287]. Similar viral-mediated loss-of-function tools were used to knockdown CRFR1 in the globus pallidus, revealing a previously unknown anxiolytic effect of the receptor in this brain region, which was further confirmed with site-specific CRFR1 antagonist administration [245]. The relevance of VTA CRFR1 receptors in addiction and reward processes was recently investigated *via* LV-mediated knockdown in this region, which attenuated cue- and acute food deprivation stress-induced cocaine seeking, but had no effect on self-administration behavior. Consequently, the authors propose that CRFR1 signaling in the VTA presents a target for convergent effects of both cue- and stress-induced cocaine-seeking pathways [288]. The previously mentioned, newly described population of CRF neurons within the VTA was investigated, amongst others, *via* AAV2-mediated knockdown, which prevented the aversive effects of nicotine withdrawal in these mice [161]. Similar to CRFR1 manipulations, virus-based tools have been used to investigate the CRFR2/UCN-system. LV-mediated knockdown and overexpression of CRFR2 in the BNST revealed an important role

Table 2. Viral-mediated gain- and loss-of-function approaches targeting CRF and its receptors.

Viral Construct	Target Site	Phenotype	References
Viral-based gain-of-function studies			
LV - tet-on based CRF OE in mice driven by a choroid plexus-specific promoter	Choroid plexus	↑ anxiety, = home cage activity	[267]
LV - CMV driven CRF OE, study in females rats	CeA	↑ CRF & AVP expression in PVN, ↓ HPA axis feedback, ↑ ASR, ↑ immobility in FST, ↓ GnRH expression in CeA, altered sexual behavior	[281]
LV - CMV driven CRF OE in mice, injection at week 7; tested 4 months later	CeA BNST	= basal anxiety, ↓ stress-induced anxiety, = immobility in FST, = HPA axis activity, = ASR, ↓ reaction time to startle stimuli = basal & stress induced anxiety, = ASR = immobility in FST, = HPA axis activity	[283]
LV - tet-on based, short-term CRF OE in mice driven by a CMV promoter; analyses 3 days post DOX induction	CeA	= basal anxiety, ↑ stress-induced anxiety, = immobility in TST, = fear conditioning, = HPA axis activity	[282]
LV - CRF OE driven by a 3kb CRF promoter fragment, analyses in rats 2 weeks post injection	CeA BNST	↑ CRF & AVP expression in PVN, HPA axis hyperactivity, ↑ anxiety = basal anxiety & ASR, = HPA axis activity, alterations in conditioned anxiety	[284] [285]
LV - CRFR2 OE in rats driven by a pCSC minimal promoter	BNST	↓ PTSD-like symptoms	[290]
Viral-based loss-of-function studies			
LV - shRNA based CRFR1 KD in mice	BIA	↓ anxiety, = general locomotion	[127]
LV - shRNA based CRF KD in mice	CeA	↓ stress-induced anxiety, ↑ basal CORT levels, = home cage activity, = immobility in TST, ↑ CRFR1 expression in BNST	[282]
LV - shRNA based CRFR1 KD in mice	Globus pallidus	↓ anxiety, = general locomotion	[245]
LV - siRNA based CRF KD in mice	PVN	= basal social avoidance, ↓ stress-induced social avoidance	[287]
LV - shRNA based CRFR2 KD in mice	BNST	↓ susceptibility to PTSD-like behavior	[289]
LV - shRNA based CRF KD in mice	VTA	Prevents aversive effects of nicotine withdrawal, ↓ escalation of nicotine intake	[161]
LV-shRNA based CRFR1 KD in mice		Blocks acute food deprivation stress-induced reinstatement of cocaine seeking, ↓ cue-induced cocaine seeking	[288]

↑ indicates an increase, ↓ indicates a decrease, = indicates no difference compared to control animals. Abbreviations: Acoustic startle response (ASR), arginine vasopressin (Avp), basolateral nucleus of the amygdala (BIA), bed nucleus of the stria terminalis (BNST), central nucleus of the amygdala (CeA), corticosterone (CORT), doxycycline (DOX), forced swim test (FST), gonadotropin-releasing hormone (GnRH), hypothalamic-pituitary-adrenal (HPA) axis, knockdown (KD), lentivirus (LV), overexpression (OE), paraventricular nucleus of the hypothalamus (PVN), post-traumatic stress disorder (PTSD), tail-suspension test (TST), ventral tegmental area (VTA).

of this receptor in PTSD-like behavior [289, 290], while knockdown in the VMH further emphasized CRFR2 as a critical mediator in regulating feeding and lipid metabolism [214].

LVs and AAVs have clearly become indispensable tools in neuroscience research, and will help to further elucidate the involvement of the CRF system in healthy and pathological stress-responses. Yet it will be important to enable more precise comparability between the different studies, which is currently problematic considering the diversity of employed viruses and serotypes, behavioral paradigms, test-

ing conditions, and rodent models (mouse versus rat). In addition, the utilized promoters driving transgene expression, as well as the duration of viral expression, varied across studies, which might have resulted in different intensities of gene expression depending on the respective construct.

8. INDISPENSABLE TOOLS IN CRF RESEARCH – PAST, PRESENT AND FUTURE

The pharmacological, genetic and virus-based studies reviewed above clearly support a role for the CRF/CRFR system in stress-related neuroendocrine, autonomic and be-

havioral alterations, which is of relevance to a number of psychiatric disorders. In particular, the involvement of CRF/CRFR1 in the regulation of HPA axis function and emotional behavior is well established. On the other hand, the role of the central UCN/CRFR2-system in the context of emotional behavior is less clearly defined, while the impact of this circuit on metabolic function is just starting to evolve. The field of stress research has advanced considerably, yet little is known about specific CRF/UCN-neurotransmitter interactions during normal and pathological conditions. Although the neurochemical identity of CRFR1-expressing neurons in the mouse brain has largely been dissected, the same cannot be stated for CRF, CRFR2 and the UCNs. In addition, the projection sites of neurons expressing CRF and the UCNs have scarcely been investigated, as well as the precise subcellular localization of CRFR1 and CRFR2 in different neuronal subpopulations. This gap of knowledge can be ascribed to the lack of specific antibodies and genetic tools targeting CRF system components. The necessity to further dissect the CRF system at the circuit level becomes evident in view of the fact that CRFR1 was shown to regulate anxiety-related behavior in opposite directions in glutamatergic and dopaminergic neurons [106]. Based on this and other recent studies, it can be implied that the CRF/CRFR1-system does not solely modulate aversive responses similar to those observed after stress. However, a detailed analysis of “aversive” vs “appetitive” CRF pathways is still missing. Consequently, researchers have to increase their focus on the physiological role of CRF and the UCNs, and assess behavioral effects not only in response to site/region-specific manipulations of the system, but also at the circuit level. The future generation of conditional CRF, UCN1-3 and CRFR2 knockout mice, as well as specific Cre-driver lines reflecting the expression of the CRF ligands and their receptors, is mandatory for such an endeavor. The latter is of utter importance, considering the limited availability of specific and high-quality antibodies, and the low expression level of CRF-related peptides and receptors in certain brain regions, which also challenges their detection at the mRNA level by in situ or double in situ hybridization. In addition, the combination of Cre-dependent optogenetic tools with specific Cre-driver lines (e.g. CRF- or CRFR1-Cre) would permit selective, bidirectional control of neural activity in genetically defined cell populations [291-293], enabling direct analysis of endogenous circuits related to emotion, cognition and addiction. With respect to the CRF/UCN system, this was elegantly demonstrated by Anthony and colleagues, who made use of optogenetics to specifically activate or inhibit CRFR2-expressing neurons in the LS. This was achieved by generating a transgenic CRFR2-Cre drive line and subsequently injecting Cre-dependent AAVs expressing either activating channelrhodopsin or inhibiting halorhodopsin into the LS [294]. The study revealed that CRFR2-positive cells of the LS include GABAergic projection neurons that innervate the anterior hypothalamus. Surprisingly enough, these CRFR2 outputs were shown to promote, rather than suppress, stress-induced behavioral measures of anxiety and corticosterone levels [294]. However, the utility of specific CRF family-related Cre-lines is much more versatile, especially when bred to Cre-dependent reporter strains. This makes the cells of interest easily accessible for different pur-

poses: 1) visualization and morphological characterization, 2) co-expression analyses, 3) mRNA isolation and RT-PCR, 4) laser capture dissection, 5) fluorescence-activated cell sorting, 6) and electrophysiological assessment. In addition, breeding the specific Cre-lines to mouse strains that contain “floxod” genes of interest or expression cassettes, facilitates gain- or loss-of-function of specific genes within Cre-expressing cells. Spatial and temporal restriction can further be enhanced by the use of Cre-dependent viruses to deliver and express a gene of interest, which is also of relevance for neuronal tracing approaches. An alternative to the Cre-line × reporter approach are transgenic mice expressing fluorescent proteins under the direct control of the CRF and CRFR1 promoters [103, 123]. In addition, CRFR1, UCN1, UCN2 and UCN3 knockout mice that have been generated *via* replacement of the coding region by EGFP, LacZ or tau-LacZ reporter cassettes, can also be exploited to visualize the desired neurons [42, 59, 252]. However, the specificity of transgene expression in Cre-, or reporter-lines is a major issue and strongly dependent on the utilized promoter fragment, site of transgene insertion and the number of inserted transgene copies. As a result, Cre recombinase activity might only be detected in a subset of the targeted cell population, or in the worst case, be expressed ectopically [295, 296]. Thus, it is of utter importance to confirm that the transgene expression accurately recapitulates that of the gene of interest. Besides the mentioned CRFR2-Cre mice, two UCN3-Cre and four CRF-Cre mouse lines have been generated to date (Table 3). Both UCN3-Cre lines (KF43 and KF31) are available from the Gene Expression Nervous System Atlas (GENSAT) project (<http://www.gensat.org>) but have not been fully characterized yet. In contrast, all four available CRF-Cre mice have partially been investigated (Table 3). A recent study convincingly demonstrated a partially divergent pattern of CRF-Cre mediated reporter expression in two of these lines (CRFp3.0Cre and *Crh-ires-Cre* mice), highlighting the importance of confirming the cell-type specificity of the utilized Cre-driver [297]. Cre-activated LacZ expression in transgenic CRFp3.0Cre mice, generated by Martin and colleagues, was demonstrated in the PVN, CeA and BNST, three major CRF-expression sites in the brain [286]. A small amount of LacZ expressing neurons were scattered within the cortex, consistent with the pattern of cortical CRF expression. However, the authors reported lack of LacZ expression throughout the remainder of the brain and brainstem, which is not in accordance with the endogenous CRF expression, which is additionally found in other brain structures (e.g. piriform cortex, hippocampus, Barrington’s nucleus and inferior olive). Thus, CRFp3.0Cre mice seem to reflect the expression of a subset of CRF neurons in the brain. Importantly, CRFp3.0Cre-dependent GFP expression colocalized to some extent with endogenous CRF in the CeA [286], but was not confirmed for other CRF-expressing brain regions. Two additional studies made use of the CRFp3.0Cre-driver to study the role GABA(A) α 1 and the Grin1 receptor (glutamate receptor, ionotropic, N-methyl D-aspartate 1) specifically in CRF neurons [298, 299]. Deletion of the GABA(A) α 1 in CRF neurons was shown to enhance anxiety and disrupt fear extinction [298], while deletion of Grin1 enhanced fear memory [299]. In addition to the CRFp3.0Cre-line, GENSAT has also produced a CRF-Cre

Table 3. Cre-driver mouse lines related to the CRF/CRFR system.

Cre-driver lines	Construct	Expression pattern	Reference
CRF-Cre lines			
CRFp3.0Cre	Transgenic - 3.0 kb of CRF promoter were amplified and cloned in front of the Cre coding sequence	Reflects a subset of the endogenous expression pattern; CRF-specific expression in the PVN, CeA, BNST, hypothalamus and scattered neurons of the cortex; lack of expression in other CRF-containing nuclei	[286, 298, 299]
CRH-Cre (KN282)	BAC transgenic - insertion of the Cre gene, followed by a polyA sequence, at the initiating ATG codon of the 2 nd CRF exon	Only partially reflects the endogenous expression pattern e.g. PVN; absence of Cre activity in the CeA and BNST, ectopic expression throughout the thalamus, olfactory bulb, & hippocampal pyramidal neurons	[300, 301] http://www.gensat.org/cre.jsp
<i>Crh-IRES-Cre</i>	Knockin - insertion of an ires-Cre targeting cassette immediately after the translational STOP codon of the endogenous <i>CRF</i> gene	Cre expression is regulated by the endogenous CRF promoter and thus largely recapitulates the endogenous CRF expression pattern	[147, 302-307]
<i>Chr-ires-Cre</i>	Knockin - insertion of an ires-Cre targeting cassette after the coding region of the endogenous CRF gene	Cre expression is regulated by the endogenous CRF promoter and thus largely recapitulates the endogenous CRF expression pattern	[308]
CRH-creERT2	BAC transgenic - insertion of CreERT2 into the human CRF gene; BAC was inserted into the Hprt locus	Cre expression is induced by tamoxifen and regulated by CRF promoter elements; very few scattered Cre expressing cells in regions of endogenous CRF expression	http://www.informatics.jax.org/allele/MGI:5568222 Simpson & Deussing unpublished data
UCN3-Cre lines			
<i>Ucn3-Cre (KF31)</i>	BAC transgenic - insertion of the Cre gene, followed by a polyA sequence, at the initiating ATG codon of the first coding exon of <i>Ucn3</i>	Expression in specific hypothalamic nuclei, moderate to strong expression in the medial amygdala; scattered probably ectopic expression in the caudate putamen and ventral striatum	http://www.gensat.org/cre.jsp
<i>Ucn3-Cre (KF43)</i>	BAC transgenic - insertion of Cre gene, followed by a polyA sequence, at the initiating ATG codon of the first coding exon of <i>Ucn3</i>	Expression largely reflects endogenous <i>Ucn3</i> expression pattern	http://www.gensat.org/cre.jsp
CRFR2-Cre lines			
<i>Crfr2a-eGFP-Cre</i>	BAC transgenic - eGFP-Cre fusion protein driven by the CRFR2 α promoter	~ 92 % of Cre-expressing cells co-localize with endogenous CRFR2 mRNA in the lateral septum; no detailed analysis of other expression domains	[294]
<i>Crhr2-Cre (RT30-Cre)</i>	BAC transgenic - insertion of Cre gene, followed by a polyA sequence, at the initiating ATG codon of the first coding exon of CRFR2	Not fully characterized, little specificity for CRFR2-neurons	http://www.gensat.org/cre.jsp

strain (KN282), which was previously used by Sarkar and colleagues to investigate the effect of the stress-derived neurosteroid tetrahydrocorticosterone (THDOC) on CRF neurons in the PVN, in the context of HPA axis regulation [300]. Although Cre-expression in the PVN was verified to recapitulate that of endogenous CRF [300, 301], it is absent in the CeA and BNST, and ectopically expressed throughout the thalamus, olfactory bulb, and pyramidal neurons of the hippocampus (<http://www.gensat.org>). Cre expression in unintentionally targeted cells in the KN282 CRF-Cre transgenic mouse line could confound the interpretation of cell-type specific experiments, and should be used with caution. In order to

circumvent the uncertainties that arise due to random integration of DNA constructs in transgenic Cre-lines, targeted knockin strategies in embryonic stem cells have been used. The recently generated CRH-ires-Cre mouse line relies on this principle [302]. In this case, a targeting cassette containing an internal ribosome entry site (ires), followed by the Cre recombinase coding region, was inserted immediately after the translational STOP codon of the endogenous CRF gene [302]. As a result, Cre expression is regulated and directed by the endogenous CRF promoter/enhancer elements, without compromising CRF expression. CRH-ires-Cre recombinase activity is reported as highly specific and efficient, and largely recapitu-

lates the endogenous CRF expression pattern [302], including the PVN, BNST, CeA, piriform cortex, inferior olive and scattered neurons of the cortex and hippocampus (images of Cre-dependent reporter expression are available at the Allen Institute for Brain Science website - <http://brain-map.org/>). Consequently, an increasing number of studies has started to employ the *Crh-ires-Cre* driver line to analyze different populations of CRF neurons [147, 303-307]. For example, Cusulin and colleagues functionally characterized CRF-expressing PVN neurons [303], while Silberman *et al.* examined the electrophysiological properties of CRF neurons in the BNST and CeA in the context of addiction [147]. The latter demonstrate that CRF-expressing BNST neurons regulate ethanol-seeking behavior by potentiating glutamatergic transmission in a population of VTA-projecting BNST neurons [147]. Recently, a similar strategy was applied to generate an independent *Crh-ires-Cre* knockin line, which also strongly reflects the endogenous CRF expression pattern [308]. Attempts to generate an inducible CreERT2-based CRF Cre driver line, undertaken within the Pleiades Promoter Project and CanEuCre [309, 310], were of limited success, *i.e.* a CRF-specific BAC construct introduced into the *Hprt* locus showed only limited and partially ectopic Cre activity (E.M. Simpson & J.M. Deussing, unpublished results). However, the generation of an adequate *CRF-Cre* counterpart, a *CRFR1-Cre* driver, has not yet been realized. A *CRFR1-Cre* line would significantly contribute to our understanding of CRF/CRFR1-pathways in the brain, as it would enable more precise circuit analysis *via* retrograde tracing approaches and optogenetic manipulations. With the current development and expansion of CRISPR/Cas technology, this goal might be achieved in the near future.

Although a number transgenic Cre-driver lines have been generated for a subset of CRF family members and their receptors, most still lack extensive anatomical, molecular, and functional characterization. The necessity for this was recently demonstrated for Cre-driver lines that target dopaminergic neurons, some of which exhibited highly non-specific expression patterns [295, 296]. In the end, complementary methodologies such as pharmacological manipulations, immunohistochemical and *in situ* hybridization studies, will considerably strengthen the conclusions drawn from the use of transgenic Cre-driver mouse lines.

CONCLUDING REMARKS

Stress-related mental disorders including depression, anxiety-disorders, as well as alcohol and substance abuse, represent some of the most common and escalating health problems in today's society. Depression tops the estimated financial burden of all psychiatric disorders, and represents the most prevalent mental illness and third leading contributor to the global disease burden [311-313]. The lack of effective treatments or preventive interventions for most mental disorders partially reflects our limited understanding of the underlying brain-circuitries. This is largely owed to the substantial amount of overlapping symptoms of many psychiatric illnesses, which makes accurate diagnosis often very difficult [314]. This overlap in disease etiology partially arises from shared genetic susceptibility factors [315]; however environmental perturbations, such as trauma and chronic stress, are additional, well-established risk factors [6, 316, 317]. Chronic stress represents a strong proximal predictor of major depres-

sive disorder onset and can also induce recurrence of depressive symptoms [318-321]. Similarly, stressful life events are associated with substance and drug abuse and are frequently reported to trigger relapse [141, 322]. Even adverse experiences in utero or during early childhood are increasingly associated with lifelong health disparities [323, 324]. But why does stress cause disease in some individuals but not in others? A common perception is that adverse environments might trigger disease onset in genetically predisposed individuals. Evidence for such gene-environment interactions have been provided by a number of studies [325-327]. Consequently, altered stress-neurocircuits, either caused by genetic, and/or environmental changes, constitute a common domain of many mental disorders, and highlight the necessity to functionally dissect the brains' most prominent stress system – the CRF/UCN/CRFR system. As summarized above, a large body of evidence has implicated CRF-family members and their receptors in the neuroendocrine and behavioral responses to stress. The ability of the CRF system to interact and modulate monoaminergic circuits is of particular importance considering the involvement of serotonergic, dopaminergic and noradrenergic systems in essentially all aspects of emotion, motivation, reward and cognition. The interaction of the CRF/CRFR-system with different neurotransmitters possibly accounts for the generation of highly specific molecular, circuit and behavioral effects under both, basal and stress conditions. Furthermore, the identity and release sites of activated CRF/UCN neurons, as well as the identity and expression sites of the responding postsynaptic CRFR neurons are likely determining specific behavioral outcomes in response to stress. Although we have substantially advanced our understanding of functional CRF circuits and their effects on stress-related behaviors, we still lack detailed knowledge about the underlying molecular mechanisms and pathways by which the brain translates stressful stimuli into the final integrated biological response under physiological and pathological conditions. Based on recent genetic and pharmacological studies, it becomes evident that CRF is losing its reputation as an “all-aversive” peptide, which is not entirely surprising considering that CRF-mediated HPA axis activation also occurs during eustress - the positive stress-response. The CRF/CRFR1 system modulates very specific and partially opposing physiological and behavioral effects depending on the underlying neuronal circuits, brain regions and environmental conditions, which also most likely holds true for CRFR2. The generation of more specific mouse genetic models, viral and optogenetic tools will enhance our understanding of CRF/UCN-CRFR1/2 circuit dynamics in adaptive and maladaptive stress-related behaviors, and aid in the development of more effective treatment modalities in psychiatry.

CONSENT FOR PUBLICATION

Not applicable.

CONFLICT OF INTEREST

The authors declare no conflict of interest, financial or otherwise.

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