

# Oxytocin Reduces Alcohol Cue-Reactivity in Alcohol-Dependent Rats and Humans

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Approved pharmacological treatments for alcohol use disorder are limited in their effectiveness, and new drugs that can easily be translated into the clinic are warranted. One of those candidates is oxytocin because of its interaction with several alcohol-induced effects. Alcohol-dependent rats as well as post-mortem brains of human alcoholics and controls were analyzed for the expression of the oxytocin system by qRT-PCR, *in situ* hybridization, receptor autoradiography ([<sup>125</sup>I]OVTA binding), and immunohistochemistry. Alcohol self-administration and cue-induced reinstatement behavior was measured after intracerebroventricular injection of 10 nM oxytocin in dependent rats. Here we show a pronounced upregulation of oxytocin receptors in brain tissues of alcohol-dependent rats and deceased alcoholics, primarily in frontal and striatal areas. This upregulation stems most likely from reduced oxytocin expression in hypothalamic nuclei. Pharmacological validation showed that oxytocin reduced cue-induced reinstatement response in dependent rats—an effect that was not observed in non-dependent rats. Finally, a clinical pilot study (German clinical trial number DRKS00009253) using functional magnetic resonance imaging in heavy social male drinkers showed that intranasal oxytocin (24 IU) decreased neural cue-reactivity in brain networks similar to those detected in dependent rats and humans with increased oxytocin receptor expression. These studies suggest that oxytocin might be used as an anticraving medication and thus may positively affect treatment outcomes in alcoholics.

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## INTRODUCTION

Alcohol use and abuse account for a tremendous burden of disease worldwide. Excessive alcohol drinking is a leading risk factor for chronic non-communicable diseases and is linked to more than 60 diseases, including cancers, cardiovascular diseases, liver cirrhosis, neuropsychiatric disorders, and fetal alcohol syndrome. One further consequence of excessive alcohol use is that 76 million adults worldwide are alcohol dependent (Rehm *et al*, 2009). Alcohol dependence is characterized by cycles of excessive alcohol consumption, interspersed with intervals of abstinence, and

frequent relapses (Heilig *et al*, 2010). Relapse is a key element of this disease process and blocking relapse is therefore a key objective for the treatment of alcohol-dependent patients. Currently, there are four antirelapse medications approved by the US Food and Drug Administration to treat alcohol dependence: disulfiram, oral naltrexone, a long-lasting injectable naltrexone, and acamprosate. These drugs have small to medium effect sizes and only a minority of patients benefit from these treatments (Litten *et al*, 2012). Another problem is that there is a huge treatment gap—most alcoholics do not seem to obtain appropriate treatment (van Amsterdam and van den Brink, 2013). Given this situation, new drugs that can be easily translated into clinical application are warranted. One of those candidates is oxytocin (OXT), because this drug interacts with several effects induced by alcohol and has already been applied in many clinical settings.

OXT is a neuropeptide that has important and well-documented roles in mammalian social behavior and various other central and peripheral processes (Meyer-Lindenberg *et al*, 2011; Neumann and Landgraf, 2012; Grinevich *et al*, 2016). Numerous studies in rodents have found that OXT

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alters neuroadaptation to drugs of abuse including alcohol, and that OXT is a potential mediator and modulator of addictive behavior (Sarnyai, 2011; Sarnyai and Kovacs, 2014). Indeed, peripheral as well as central administration of OXT blocks the development of tolerance to the hypothermic and hypnotic effects of alcohol in rodents and modulates the severity of alcohol withdrawal, demonstrating that OXT alters neuroadaptations to alcohol (Bowen *et al*, 2015). Very recently, it has been shown that acute intranasal OXT treatment, which leads to a substantial increase of OXT in the cerebrospinal fluid (Striepens *et al*, 2013), also reduces withdrawal symptoms in alcohol-dependent patients (Pedersen *et al*, 2013). Several studies further found an OXT-dependent increase of functional connectivity between the amygdala and insular cortex, orbitofrontal cortex, anterior cingulate cortex (Cg), nucleus caudatus (NC), and hippocampus, which all are regions relevant in alcohol cue-reactivity and the development of craving (Bethlehem *et al*, 2013). These findings suggest that OXT might decrease cue-reactivity and craving in social drinkers.

Here we set out to systematically study neuroadaptive changes in the OXT system in alcohol-dependent individuals in brain sites that are relevant for mediating alcohol reinforcement and addictive behavior (Koob and Volkow, 2010). For this we quantified mRNA and protein levels of OXT and OXT receptors (OXTRs) in brain tissue from abstinent alcohol-dependent rats and deceased alcoholics. We then studied the effects of OXT on alcohol-seeking behavior in the reinstatement paradigm (Sanchis-Segura and Spanagel, 2006). For animal behavior experiments, we used the so-called 'post-dependent animal model' (Heilig and Koob, 2007) where dependence is induced through repeated intermittent cycles of alcohol vapor exposure. This model has already been proven valuable in the investigation of several neurotransmitter systems (Sommer *et al*, 2008; Hermann *et al*, 2012; Meinhardt *et al*, 2013; Hirth *et al*, 2016; Uhrig *et al*, 2017), and is used as a model for medication development in alcoholism (Meinhardt and Sommer, 2015). Additionally, in a pilot study in non-treatment seeking heavy social drinkers, we examined neural cue-reactivity, which was previously shown to be associated with subsequent craving and relapse in alcohol-dependent patients (Grüsser *et al*, 2004; Reinhard *et al*, 2015). Here we applied functional magnetic resonance imaging (fMRI) to measure neural reactivity to alcohol-related cues following intranasal administration of a single dose of OXT conducted in a crossover design with placebo control. The major hypothesis was a reduction of cue-reactivity following the presentation of alcohol cues by a single dose of OXT compared with placebo.

## MATERIALS AND METHODS

Human post-mortem brain samples of 27 male Caucasians (13 alcoholics and 14 controls) were obtained from the NSW Tissue Resource Center (University of Sydney, Australia; see Supplementary Table S1; Hermann *et al*, 2017), and analyzed for OXTR expression by quantitative RT-PCR (qRT-PCR) and receptor autoradiography ( $[^{125}\text{I}]\text{OVTA}$  binding) according to Olazabal and Young (2006b) and Uhrig *et al* (2016).

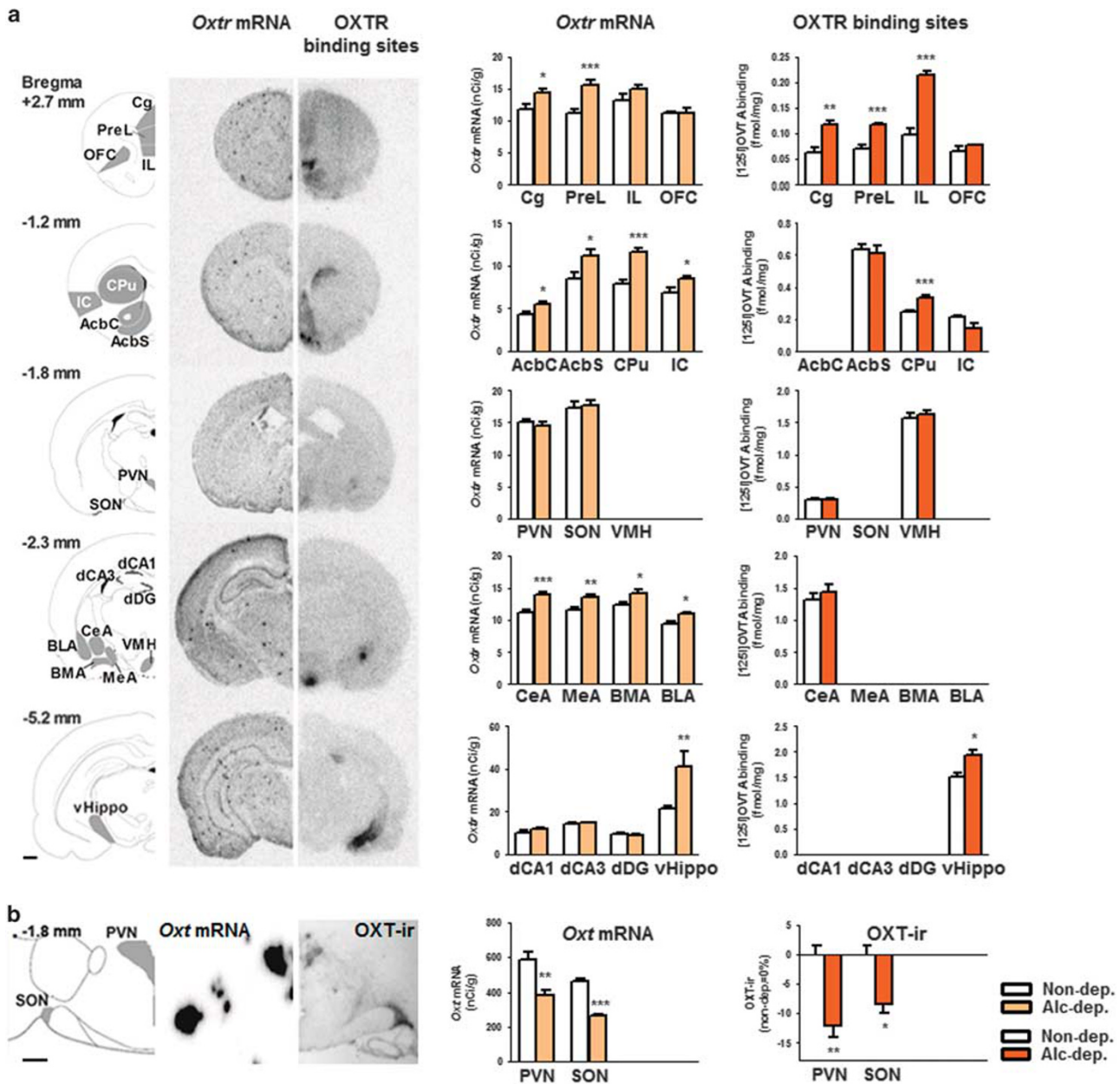
In male Wistar rats, alcohol dependence was induced by exposure to chronic intermittent cycles of alcohol inhalation and rats were killed at different time points of abstinence (0, 1, 3, 7, and 21 days after the last vapor exposure according to Hansson *et al*, 2010; Hermann *et al*, 2012; Hirth *et al*, 2016; Uhrig *et al*, 2017) for OXT and OXTR expression analysis by *in situ* hybridization, receptor autoradiography, western blotting, and OXT immunohistochemistry. Additionally, cue-induced reinstatement after intracerebroventricular injection of either 10 nM OXT or vehicle (aCSF) was performed in 3 weeks abstinent rats. The experiments were conducted in accordance with the European ethical guidelines for the care and use of laboratory animals, and were approved by the local animal care committee (Regierungspräsidium Karlsruhe, Germany, license number 35-9185.81/G-302/15 and AZ 35-9185.81/G-183/09).

For the clinical study, 12 heavy social drinkers were recruited at the Central Institute of Mental Health (Mannheim, Germany), for fMRI studies on alcohol cue-induced activation after a single application of intranasal OXT according to Sauer *et al* (2013) (German clinical trial number DRKS00009253; URL: [https://www.drks.de/drks\\_web](https://www.drks.de/drks_web)). All experimental procedures and statistics are described in detail in Supplementary Information.

## RESULTS

### Animal Studies

*Oxtr* is increased and the endogenous ligand OXT is decreased in abstinent rats. We used the post-dependent animal model—a well-established rat model for alcohol dependence (Rimondini *et al*, 2002; Meinhardt *et al*, 2013; Hirth *et al*, 2016; Uhrig *et al*, 2017). Rats were exposed to 7 weeks of intermittent cycles of alcohol vapor to induce dependence, reaching blood alcohol concentrations (BACs) of  $266.2 \pm 12.6$  mg/dl. Three weeks after the last vapor exposure, rats were killed in abstinence to investigate changes in the expression of OXT peptide and OXTRs at both the mRNA and protein levels. Forebrain regions related to reward circuits (Noori *et al*, 2012) were used for *in situ* hybridization, receptor autoradiography (Uhrig *et al*, 2016), western blotting, and immunohistochemistry experiments (see Supplementary Figure S1). Semiquantitative analysis of the *Oxtr* mRNA revealed an increase in alcohol-dependent animals in the medial prefrontal cortex (cingulate cortex (Cg): 22% increase; prelimbic region (PreL): 39% increase), striatum (nucleus accumbens core (AcbC): 30% increase; nucleus accumbens shell (AcbS): 33% increase; caudate putamen (CPu): 46% increase), insular cortex (IC: 23%), amygdala (central amygdaloid nucleus (CeA): 26% increase; medial amygdaloid nucleus (MeA): 17% increase; basomedial amygdaloid nucleus (BMA): 16% increase; basolateral amygdaloid nucleus (BLA): 17% increase), and ventral hippocampus (vHippo, 94% increase), including ventral hippocampal subregions (ventral cornu ammon (CA) regions vCA1 and vCA3, ventral dentate gyrus (vDG)). No changes of *Oxtr* mRNA expression were found in the infralimbic region (IL), orbitofrontal cortex (OFC), hypothalamic paraventricular nucleus (PVN), supraoptic nucleus (SON), and dorsal hippocampal subregions (dCA1, dCA3, dDG), see Figure 1a. F- and p-values are summarized in Supplementary Table S1.



**Figure 1** Expression of the oxytocin (OXT) system in 3 weeks abstinent rats. (a) OXT receptor (*Oxtr*) mRNA and OXTR binding sites are increased in several forebrain regions of dependent rats. (Left panels) Schematic outline of analyzed regions in the rat brain according to Paxinos and Watson (1998) (left). OXTR expression pattern for mRNA (*in situ* hybridization, middle) and binding sites ( $[^{125}\text{I}]\text{OVTA}$  receptor autoradiography, right) corresponding to the schematic outline. (Right panels) Bar graphs showing *Oxtr* mRNA (left) and OXTR binding sites (right) in analyzed regions of alcohol-dependent (Alc-dep., orange bars) and -non-dependent (Non-dep., white bars) rats. Data are expressed as mean  $\pm$  SEM (nCi/g for mRNA or fmol/mg for binding sites). Statistical analysis was performed by region-wise one-way analysis of variance (ANOVA),  $N=4-8/\text{group}$ ,  $p$ -values: \* $p < 0.05$ ; \*\* $p < 0.01$ ; \*\*\* $p < 0.001$ . For abbreviations and experimental procedures, see Materials and Methods. (b) Reduced hypothalamic OXT expression in dependent rats. (Left panels) Schematic outline of the hypothalamic rat brain region at Bregma level  $-1.8$  mm according to Paxinos and Watson (1998) (left). OXT mRNA pattern (middle) and OXT immunoreactivity (ir, right) corresponding to the schematic outline in the PVN and the SON. (Right panels) Bar graphs showing *Oxt* mRNA (left) and OXT-ir cells (right) in the PVN and SON of alcohol-dependent (Alc-dep., orange bars) and -non-dependent rats (Non-dep., white bars), displayed as mean  $\pm$  SEM (*Oxt* mRNA: nCi/g, OXT-ir: mean grey value). OXT-ir mean grey values of non-dependent rats (PVN: mean grey value =  $74.2 \pm 1.1$ ; SON: mean grey value =  $63.0 \pm 1.0$ ) are defined as 0% and changes in density show decreased density in OXT-ir in alcohol-dependent rats (dark-orange bars). Statistical analysis was performed by region-wise one-way ANOVA,  $N=7-8$  rats/group for *in situ* hybridization and  $n=3-4$  rats/group for immunoreactivity measurements,  $p$  values: \* $p < 0.05$ ; \*\* $p < 0.01$ ; \*\*\* $p < 0.001$ . AcbC, nucleus accumbens core; AcbS, nucleus accumbens shell; BLA, basolateral amygdaloid nucleus; BMA, basomedial amygdaloid nucleus; CeA, central amygdaloid nucleus; Cg, cingulate cortex; CPu, caudate putamen; dCA1, dorsal hippocampal cornu ammon 1 subregion; dCA3, dorsal hippocampal cornu ammon 3 subregion; dDG, dorsal dentate gyrus; IC, insular cortex; IL, infralimbic region; MeA, medial amygdaloid nucleus; OFC, orbitofrontal cortex; PreL, prelimbic cortex; PVN, hypothalamic paraventricular nucleus; SON, supraoptic nucleus; vHippo, ventral hippocampus; VMH, ventromedial hypothalamic nucleus. Scale bars 1 mm.



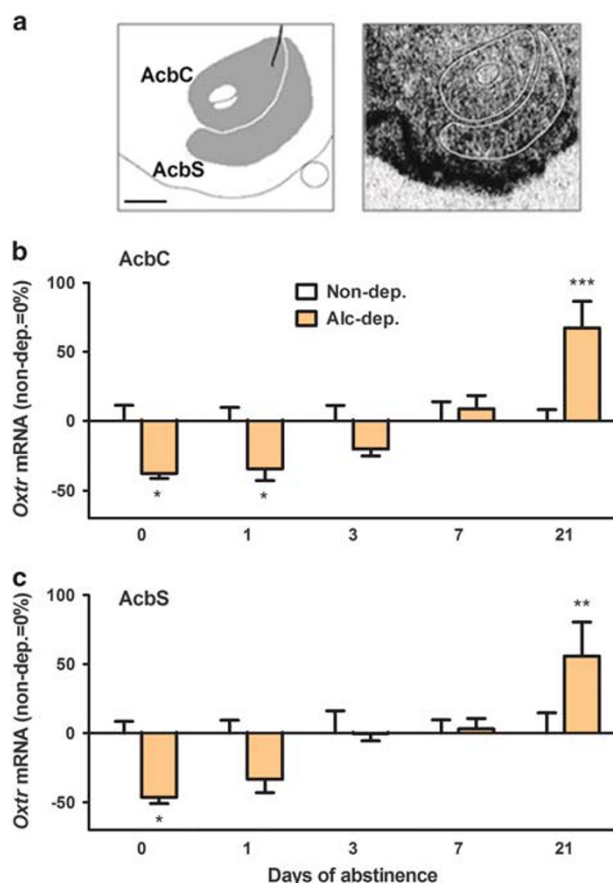
Changes in mRNA levels translated into an increase in OXTR binding sites in the Cg (86% increase), the PreL (65% increase), the CPu (38% increase), and the vHippo (28% increase) (Figure 1a). In the IL, there was also an increase of 117%. In all other investigated regions, no changes in OXTR binding sites were found. F and *p*-values are summarized in Supplementary Table S2.

The receptor binding results were validated in the CPu and the PVN by western blotting. There was a 2.8-fold increase of OXTR/ $\beta$ -actin ratio in the CPu ( $F[1,5]=6.7$ ,  $p<0.05$ ; Supplementary Figure S2) and no change of OXTR/ $\beta$ -actin ratio in the PVN ( $F[1,5]=0.2$ ,  $p>0.7$ ; Supplementary Figure S2) compared with controls. These data show a similar regulation as described for the receptor binding sites above. Thus, using two different methodological approaches, we found an increase of both receptor binding sites and protein in protracted abstinence.

Given the increased OXTR expression in alcohol-dependent rats, we next asked if the endogenous ligand OXT was also altered in dependence. *Oxt* mRNA expression and OXT-immunoreactive (-ir) cells were measured in the PVN and SON (Figure 1b), the principal brain regions in which OXT is produced and transported to secondary locations within and outside the brain. The *Oxt* mRNA and OXT-ir patterns were consistent with previous publications (Ben-Barak *et al*, 1985; Calcagnoli *et al*, 2014). A decrease in *Oxt* mRNA was found in both regions (PVN: 35% decrease,  $F[1,13]=11.9$ ,  $p=0.004$ ; SON: 44% decrease,  $F[1,13]=193.6$ ,  $p<0.001$ ; Figure 1b). This decrease was also supported by semiquantitative analysis of OXT, as evidenced by densitometry of immunoreactive cells (PVN: 12% decrease,  $F[1,5]=25.1$ ,  $p=0.004$ ; SON: 8% decrease,  $F[1,5]=13.2$ ,  $p=0.015$ ; Figure 1b).

Thus, after 3 weeks of abstinence, *Oxt* mRNA and OXT-ir were decreased, whereas *Oxtr* mRNA and OXTR binding sites were increased in several brain regions.

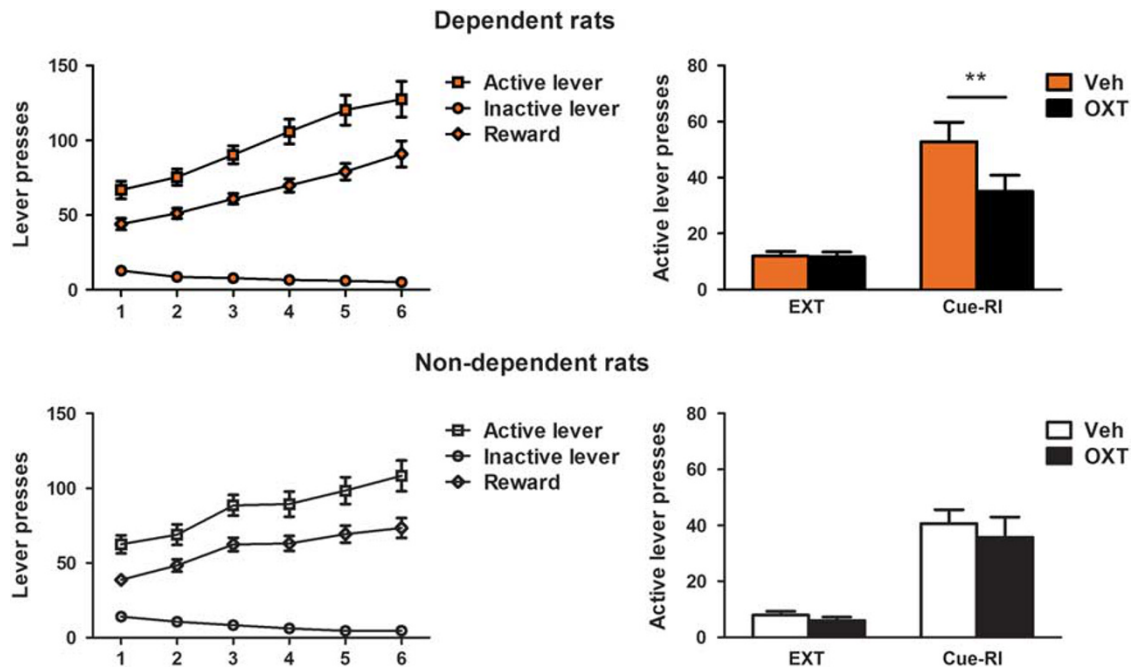
We recently described dynamic changes of the DA system during abstinence with the same animal model (Hirth *et al*, 2016). Since OXTRs are located on DAergic neurons of the nucleus accumbens (Acb) (Peris *et al*, 2017) and OXT can modulate alcohol-induced DA release in the same region (Peters *et al*, 2016), we chose the Acb region to investigate the changes in *Oxtr* mRNA expression over time during abstinence. Rats were exposed to 7 weeks of intermittent cycles of alcohol vapor and killed at several time points afterward (0, 1, 3, 7, or 21 d after vapor exposure) (Hansson *et al*, 2010; Hirth *et al*, 2016; Uhrig *et al*, 2017). There was a significant effect of Time ( $F[1,66]=7.7$ ,  $p<0.001$ ) and Treatment  $\times$  Time ( $F[1,66]=7.7$ ,  $p<0.001$ ) in the AcbC as well as in the AcbS (Time:  $F[1,65]=5.1$ ,  $p=0.001$ ; Treatment  $\times$  Time:  $F[1,65]=5.1$ ,  $p=0.001$ ). In both the AcbC and AcbS, *Oxtr* mRNA expression was significantly decreased directly after vapor exposure (0 day), when the rats were still intoxicated (BAC:  $273 \pm 52$  mg/dl; AcbC: 38% decrease,  $p=0.016$ ; AcbS: 46% decrease,  $p=0.011$ ). After 1 day of withdrawal, *Oxtr* mRNA was still decreased in the AcbC (34% decrease,  $p=0.033$ ) and there was a trend toward a decrease in the AcbS (33% decrease,  $p=0.055$ ). No effect was observed for both regions after 3 and 7 days of abstinence; however, after 21 days there was a significant increase in *Oxtr* mRNA expression (AcbC: 67% increase,  $p<0.001$ ;



**Figure 2** Oxytocin mRNA (*Oxtr* mRNA) is reduced in the nucleus accumbens by acute alcohol and during early withdrawal, while increased after prolonged abstinence. (a, left) Schematic outline of the nucleus accumbens core (AcbC) and the nucleus accumbens shell (AcbS) according to Paxinos and Watson (1998), and *Oxtr* mRNA pattern in the Acb region (right). (b) Transcriptional changes in *Oxtr* mRNA level during various stages of abstinence in the AcbC and (c) in the AcbS of dependent rats (Alc-dep., light-orange bars) and non-dependent rats (Non-dep., white bars). Bar graphs show the percentage of regulation relative to non-dependent rats. Absolute values are listed in Supplementary Table S3. Statistical analysis was performed by two-way analysis of variance (ANOVA), followed by Fisher's least significant difference (LSD) test,  $N=6-8$ /group,  $p$ -values: \* $p<0.05$ , \*\* $p<0.01$ , \*\*\* $p<0.001$ . Scale bar 1 mm.

AcbS: 56% increase,  $p=0.002$ ; Figure 2). Absolute values are listed in Supplementary Table S3.

We also investigated changes of *Oxt* mRNA during abstinence in the PVN and SON. No differences were found between the control groups in both regions. Therefore controls were pooled, and data were analyzed by one-way ANOVA and the respective alcohol-dependent group was compared to the pooled control group using Fisher's LSD post-hoc test. There was a significant overall treatment effect in the PVN ( $F[1,5]=3.1$ ,  $p<0.05$ ) and SON ( $F[1,5]=3.0$ ,  $p<0.05$ ). Post-hoc analysis revealed a similar regulation of *Oxt* mRNA in both regions. At the end of alcohol exposure (at day 0) transcripts were decreased in the PVN by 13% ( $p<0.01$ ) and in the SON by 50% ( $p<0.01$ ). Transcripts remained decreased during abstinence in both regions and reached significance in the PVN on day 3 (14%,  $p<0.01$ ), 7 (12%,  $p<0.05$ ), 21 (10%,  $p<0.05$ ) and in the SON on day 21 (56%,  $p<0.01$ ) as compared to controls (see



**Figure 3** Cue-induced reinstatement of alcohol-seeking is decreased after intracerebroventricular (icv) administration of oxytocin (OXT) in alcohol-dependent rats. (Left) Training of alcohol self-administration of rats later made dependent by alcohol vapor exposure (upper left) and non-dependent rats (lower left); displayed are active and inactive lever presses as well as number of received rewards of dependent (dark-orange color) and non-dependent rats (white color) (mean  $\pm$  SEM). (Right) Extinction (EXT) and cue-reinstatement (Cue-RI) of alcohol seeking after icv injection of 10 nM OXT (black bar) or vehicle (Veh) in alcohol-dependent (upper right, Veh = aCSF, dark-orange bar) and -non-dependent rats (lower right, Veh = aCSF, white bar). Data are expressed as mean  $\pm$  SEM. Statistical analysis was performed by general linear model variant analysis, followed by Newman-Keuls test if appropriate,  $p$ -values:  $**p < 0.01$ ,  $N = 7-11$ /group. Cue-reinstatement values were significantly different compared with extinction values in all tested groups (not shown in graph),  $p$ -values between groups:  $**p < 0.01$ .

Supplementary Figure S3, absolute values are summarized in Supplementary Table S3).

Thus, there was a dynamic regulation of *Oxtr* mRNA during abstinence, decreased by acute alcohol and during acute withdrawal, and increased transcripts after prolonged abstinence. In contrast, hypothalamic *Oxt* mRNA remained decreased during abstinence.

**Brain OXT decreases cue reinstatement of alcohol-seeking in alcohol dependent rats.** Our expression data shows decreased OXT and increased OXTR binding sites and expression during prolonged abstinence. These expression data predict an involvement of OXT in relapse-like behavior. Therefore, we aimed to pharmacologically validate our expression data by testing the effects of OXT on cue-induced reinstatement of alcohol-seeking behavior in 21 d abstinent rats.

Alcohol-dependent and non-dependent rats displayed a similar acquisition of alcohol self-administration during the six weeks of training sessions (see Figure 3). During the extinction phase, both groups reduced their active lever pressing to  $<10\%$  of the previous baseline in a similar manner (Figure 3). Cue-induced reinstatement of alcohol-seeking behavior was then tested after intracerebroventricular injection of OXT. The effect of the OXT dose (10 nM intracerebroventricular) was tested during reinstatement testing. The drug was injected directly before the start of the session. In alcohol-dependent rats, both groups displayed a significant reinstatement behavior ( $F[1,16] = 42.45$ ,  $p < 0.0001$ ), but rats treated with OXT showed a significantly

decreased active lever pressing compared to the control group ( $p = 0.002$ , Figure 3). Non-dependent rats equally showed significant reinstatement behavior ( $F[1,17] = 56.85$ ,  $p < 0.001$ ). Contrary to alcohol-dependent rats, OXT administration did not have an effect on the reinstatement response in non-dependent rats (Figure 3).

## Human Studies

**Increased OXTR expression and binding sites in human alcoholics.** To translationally validate these data in rats, we next examined OXTR expression analysis in post-mortem brain samples of human alcoholics, and performed a clinical pilot study on the effects of OXT on alcohol cue-reactivity in heavy social drinkers.

We analyzed changes on OXTR expression at both transcriptional (by qRT-PCR) and receptor binding (by OXTR autoradiography) levels, in a well-characterized brain tissue sample set of deceased alcoholics (Meinhardt *et al*, 2013; Hirth *et al*, 2016; Hermann *et al*, 2017). Alcoholics consumed more than 80 g alcohol per day, whereas control subjects had an average of  $<20$  g. Subjects were free of detectable blood alcohol at their time of death and there were no significant difference in age, post-mortem interval, or brain pH between the groups (see Supplementary Tables S4 and S5). These factors were included into region-wise general linear models (GLMs) for OXTR mRNA or binding (summarized in Supplementary Table S6). The analyses revealed significantly increased OXTR binding sites as assessed by [ $^{125}$ I]OVTA binding in all investigated regions (anterior cingulate cortex (ACC): 241% increase; dorsolateral

prefrontal cortex, Brodmann area 9 (BA9): 190% increase; ventral striatum (VS): 67% increase; NC: 94% increase; Figure 4 and Supplementary Table S6). Correspondingly, GLMs for qRT-PCR showed increased *OXTR* mRNA levels in BA9, VS, and NC, and no change in *OXTR* mRNA in ACC (Figure 4 and Supplementary Table S6). Thus in deceased human alcoholics and prolonged abstinent dependent rats, *OXTR* expression increases at both the mRNA and protein levels.

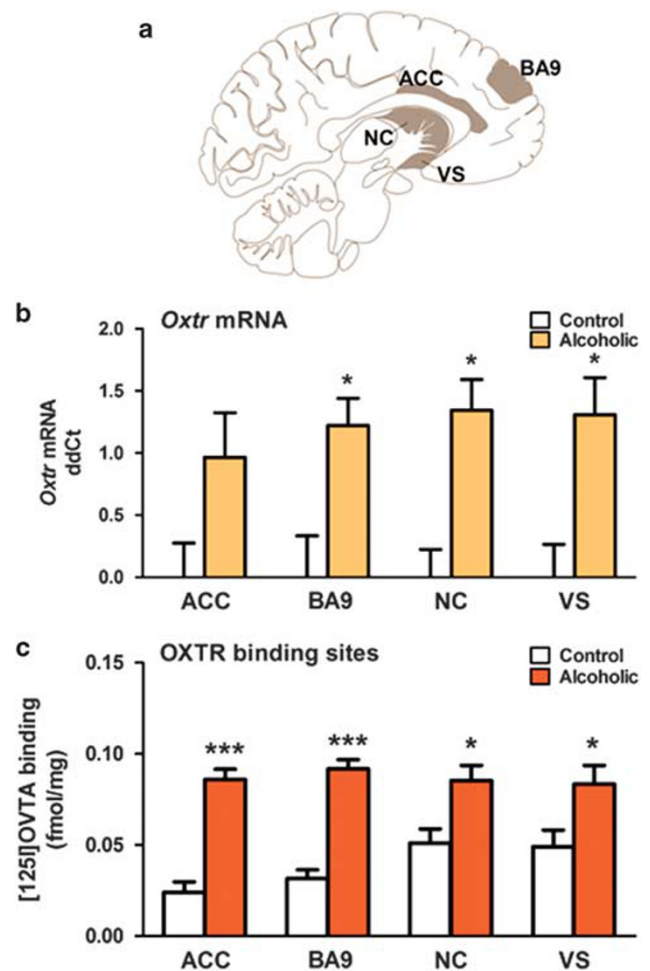
It has previously been shown that the commonly used radioligand [ $^{125}$ I]OVTA does not bind selectively to the *OXTR*, but also to the vasopressin V1a receptor and possibly other non-peptide receptors (Toloczko *et al*, 1997; Freeman *et al*, 2014a, b). We therefore investigated *AVPR1A* and *AVPR1B* mRNA expression in the BA9 region of deceased alcoholics and controls by qRT-PCR (see Supplementary Tables S4). We did not find any significant changes in alcoholic patients (*AVPR1A*:  $F[1,23] = 0.04$ ;  $p = 0.85$ ; *AVPR1B*:  $F[1,20] = 2.74$ ;  $p = 0.11$ ). This supports the specificity of increased *OXTR* binding in deceased alcoholics.

*OXT* reduces neural reactivity to alcohol-related cues in heavy social drinkers. Finally, we performed a pilot clinical trial (German clinical trial number DRKS00009253; URL: [https://drks-neu.uniklinik-freiburg.de/drks\\_web](https://drks-neu.uniklinik-freiburg.de/drks_web)) to examine whether IN OXT (24 IU) could reduce neural reactivity to alcohol-related cues in human subjects. We recruited 15 male subjects who were non-treatment-seeking heavy social drinkers (Supplementary Figure S4; for sample description and information on severity of dependence and drinking-related psychometric variables, see Supplementary Information). On average these participants were 33 ( $\pm 16$ ) years old. They consumed 2.86 ( $\pm 1.80$ ) standard drinks of alcohol ( $\pm 12$  g) per day, that is, an average of 34 ( $\pm 22$ ) g of alcohol per day. We performed fMRI on these subjects while they participated in a neural cue-reactivity test, which is a sensitive proxy measure of alcohol-seeking behavior (Jorde *et al*, 2014; Courtney *et al*, 2016; Noori *et al*, 2016).

Figure 5a displays examples of the pictorial stimuli and illustrates the fMRI block design. fMRI data of 12/15 heavy social drinkers were available for both conditions (OXT and placebo). The main effects of the contrast alcohol *vs* neutral cues for the OXT and placebo conditions are listed in Supplementary Tables S8 and S9. Comparing the neural activation during the OXT with the placebo conditions suggested that cue-reactivity was significantly decreased by OXT in the IC, the hippocampal/parahippocampal formation, the Cg, the inferior and the medial frontal gyrus, and in visual and motor regions (Figure 5 and Table 1). There was no significant increase in cue-reactivity in any brain region in the OXT condition.

## DISCUSSION

Here we provide evidence that OXT can act as a therapeutic candidate for alcohol use disorder, with promising results from three studies: analysis of an animal model of alcoholism, human post-mortem brain tissue, and supportive pilot data in heavy social drinkers. We found increased *OXTR* mRNA and *OXTR* protein levels in frontal and striatal brain areas in 3 weeks abstinent alcohol-dependent rats. Additionally, these rats showed a reduction of *Oxt* mRNA

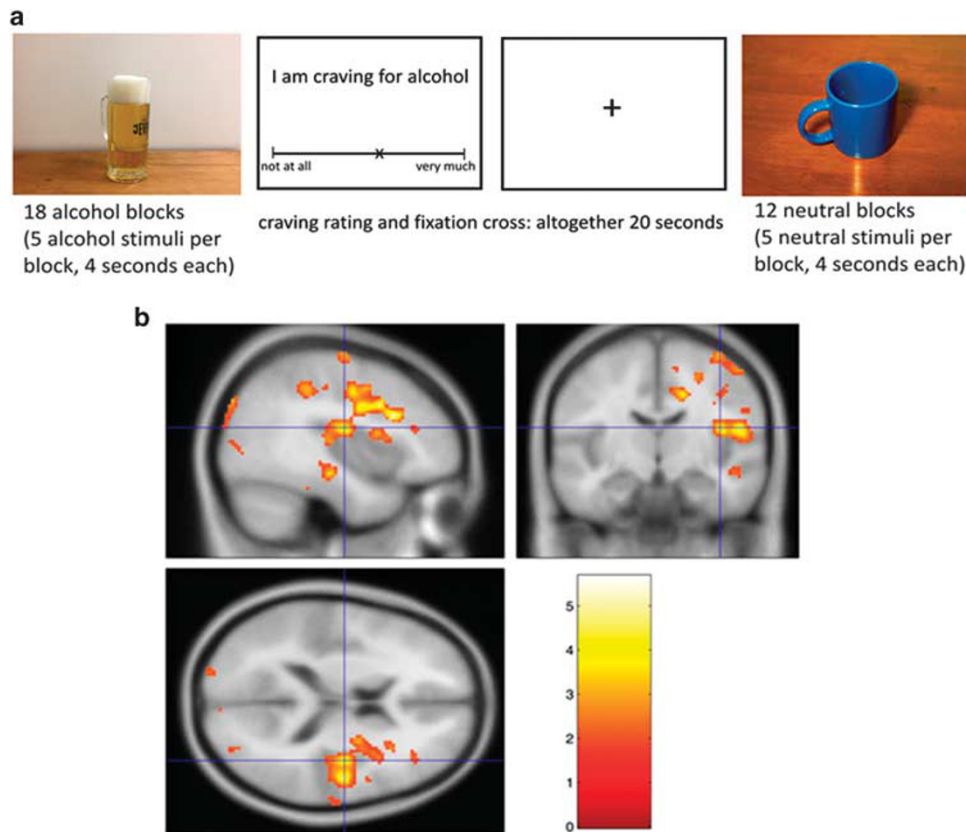


**Figure 4** Increased oxytocin mRNA (*OXTR* mRNA) and *OXTR* binding sites in post-mortem samples of alcoholics compared with control samples. (a) Schematic outline of analyzed regions in the human brain according to von Hagens *et al* (1990). (b) *OXTR* mRNA expression and (c) [ $^{125}$ I]OVTA binding; bar graphs illustrate quantitative analysis in post-mortem samples of alcoholics (*OXTR* mRNA: light orange bars; *OXTR* binding sites: dark-orange bars) and age-matched controls (white bars). Data are shown as mean  $\pm$  SEM (ddCt for mRNA expression, dCt values for controls are as follows: anterior cingulate cortex (ACC):  $11.13 \pm 0.2$ ; Brodmann area 9 (BA9):  $11.14 \pm 0.3$ ; nucleus caudatus (NC):  $9.80 \pm 0.2$ ; ventral striatum (VS):  $9.28 \pm 0.3$ ; fmol/mg for binding sites). Statistical analysis was performed by region-wise general linear models (GLMs), accounting for the possible influence of age, brain pH, post-mortem interval (PMI), and smoking;  $N = 9-15$ /group,  $p$ -values: \* $p < 0.05$ ; \*\* $p < 0.01$ ; \*\*\* $p < 0.001$ , see also Supplementary Table S6.

and OXT protein expression in hypothalamic nuclei, which provide the major source of OXT. Pharmacological validation showed that central administration of OXT reduced cue-induced reinstatement of alcohol-seeking in dependent rats, but not in non-dependent rats. We found increased *OXTR* mRNA and *OXTR* protein levels in frontal and striatal brain areas of human alcoholics, demonstrating the translational value of our results in rats. Finally, a clinical pilot fMRI study in heavy social drinkers revealed that OXT decreased neural cue-reactivity in similar brain networks as those implicated in dependent humans and rats.

OXT has been in the focus of preclinical addiction research, with findings of exogenous OXT decreasing the





**Figure 5** Oxytocin (OXT) reduces neural reactivity to alcohol-related cues in heavy social drinkers. (a) Cue-reactivity task (alcohol, neutral pictures). (b) Decrease in blood oxygenation level-dependent (BOLD) response to alcohol compared with neutral cues by oxytocin compared to placebo in heavy social drinkers,  $N = 12$ ,  $p < 0.05$  family-wise error rate (FWE)-corrected (Monte Carlo-based simulations),  $(x, y, z) = (40, -12, 16)$ , effect size in sphere of 1 cm diameter: 0.37. The neutral pictures were taken from the International Affective Picture System (IAPS; Lang *et al*, 2008).

**Table 1** Decrease in BOLD Response of Alcohol Cue-Reactivity by Oxytocin

Side	Lobe	Brain areas	Brodmann area	Cluster size	MNI coordinates	$t_{\text{maximum}}$
Right	Frontal lobe	Precentral gyrus/middle frontal gyrus/inferior frontal gyrus	6	3231	30 -4 34	5.66
Right	Limbic lobe	Cingulate gyrus/hippocampus/parahippocampal gyrus				
Right	Insular lobe	Insular				
Right	Occipital lobe	Cuneus/superior occipital gyrus/middle occipital gyrus	19	1557	32 -78 22	4.28
Left	Occipital lobe	Cuneus/superior occipital gyrus	19			
Right	Frontal lobe	Supplementary motor area/medial frontal gyrus	6	1484	16 -58 72	3.77
Right	Parietal lobe	Precuneus/inferior parietal lobule	7/40			

Abbreviations: BOLD, blood oxygenation level-dependent; FWE, family-wise error rate; MNI, Montreal Neurological Institute.

Brain areas with decreased BOLD response (alcohol compared with neutral cues) by oxytocin compared to placebo in heavy social drinkers ( $n = 12$ ),  $p < 0.05$  FWE-corrected (Monte Carlo-based simulations).

consumption of several drugs including methamphetamines, heroin, cocaine, and alcohol (Kovacs *et al*, 1998; Qi *et al*, 2009; Carson *et al*, 2010; Bowen *et al*, 2011; Peters *et al*, 2013; Lee *et al*, 2016; Leong *et al*, 2016; MacFadyen *et al*, 2016;

Peters *et al*, 2016; King *et al*, 2017). Concerning alcohol, most of the reports suggest a potential role of OXT as a therapeutic target (Lee *et al*, 2016). However, to our knowledge so far no study has been done in alcohol-dependent rats, that is, in an

animal model with alcohol-induced long-term neuroadaptations comparable to those in humans. Here, we used post-dependent rats—a well-characterized and widely accepted animal model with predictive validity and utility for medication development in alcoholism (Meinhardt and Sommer, 2015).

Intracerebroventricular-administered OXT decreased cue-reinstatement in dependent, but not in non-dependent rats. Many compounds tested so far are not specific for alcohol-dependent subjects (Meinhardt and Sommer, 2015), although several systems appear to be additionally recruited in alcohol dependence, including stress systems such as the corticotropin-releasing hormone system in the amygdala (Hansson *et al*, 2006, 2007; Sommer *et al*, 2008), as well as dynorphin (Walker *et al*, 2011) or vasopressin (Edwards *et al*, 2012). This might suggest that in alcohol dependence, the OXT system is particularly involved in and sensitized to processes mediating alcohol-seeking behavior. Thus, OXT represents a promising target for treatment development, because it would selectively affect patients with an OXT deficiency.

A striking feature of the present study is the high level of concordance in the alterations of OXTRs in alcohol-dependent humans and rats. These relations support the validity of this pathophysiological mechanism for alcohol dependence as well as the translational value of this animal model. We found increased OXTR mRNA and binding levels in striatal and prefrontocortical post-mortem samples from alcohol-dependent subjects of both species. The animal model provides important additional information about the time course of dependence-induced neuroadaptation of the OXT system. First, OXTRs undergo important changes upon cessation of alcohol exposure, being downregulated during alcohol intoxication and early withdrawal before a robust upregulation after a few weeks. Similar dynamic regulation has been found in the dopamine system of dependent rats during abstinence (Hirth *et al*, 2016). A possible link for this similar dynamical regulation of the OXT and dopamine systems has been recently provided by the finding that intracerebroventricular OXT prevents alcohol-evoked dopamine release in the Acb (Peters *et al*, 2016). It also suggests different mechanisms to be involved in the actions of OXT on acute alcohol withdrawal (Pedersen *et al*, 2013) compared with the decreased alcohol cue-reactivity reported here.

A second important finding from our animal study is that the changes in OXTR expression are widespread and region specific. Increases were not restricted to the striatum and the prefrontal cortex but also included amygdala and hippocampal regions. Interestingly, OXTRs in hypothalamic regions seemed unaffected. In the PVN and SON of dependent rats, we instead found a persistent decrease of OXT. This may be due to a loss of hypothalamic OXT neurons (Silva *et al*, 2002; Stevenson *et al*, 2017) and thus may explain why OXT treatment is effective at acute (Pedersen *et al*, 2013) and protracted abstinence. The findings agree with severe morphological changes in OXT neurons of hypothalamic nuclei and decrease of OXT immunoreactivity in the SON and the posterior pituitary lobe from human patients afflicted with chronic alcoholic disease (Sivukhina *et al*, 2006). The mechanisms underlying the dynamic changes of the receptor during abstinence remain unknown.

There are obvious mismatches of OXTR at mRNA and protein expression (eg, dorsal hippocampal neurons contain high levels of *Oxtr* mRNA with no or too low detectable amount of OXTR binding signal). One explanation may be that after synthesis the OXTR protein is transported along the axons and/or into distal dendrites at a different brain site (Peris *et al*, 2017). On the contrary, Mitre *et al* (2016) and Yoshida *et al* (2009) demonstrate a widespread expression of OXTR in different forebrain regions (including the dorsal hippocampus) using an immunohistochemistry and a transgenic reporter line approach, respectively. Thus, the observed mismatches of mRNA and receptor binding sites may be due to methodological issues.

In line with our observations, increased OXTR and decreased OXT levels were also observed after long-term cocaine, morphine, and methamphetamine administration (Sarnyai and Kovacs, 1994; You *et al*, 2000; Zanos *et al*, 2014a, b; Georgiou *et al*, 2015; Lee *et al*, 2016), and might reflect an adaptive reaction within the OXT target regions to decreased peptide expression upon chronic drug exposure. Therefore, supplementing OXT should positively affect drug-related behaviors, which has been demonstrated here for alcohol cue-reactivity in abstinent rats.

Alcohol cue-reactivity may also be affected by supplementing OXT in heavy social drinkers. Our pilot study in humans showed that a single intranasal dose (24 IU) OXT decreased neural cue-reactivity in brain regions implicated by our analysis of receptor expression in rats and deceased humans. Reduced alcohol cue-reactivity was also found in regions relevant for craving (eg, IC). The decrease of neural activation might suggest a decline of the anticipation of alcohol cues by OXT.

Intranasal OXT has been proposed as a treatment for many different conditions, including schizophrenia (Woolley *et al*, 2014; Rich and Caldwell, 2015), autism spectrum disorders (Okamoto *et al*, 2016), and substance dependence (McRae-Clark *et al*, 2013; Kirkpatrick *et al*, 2014; Lee *et al*, 2014; Flanagan *et al*, 2015; Woolley *et al*, 2016). A recent study investigated the effect of intranasal OXT on social perception, craving, and approach behavior in alcohol use disorder (Mitchell *et al*, 2016). While these authors did not find a significant effect on cue-induced craving between groups, OXT appeared to differentially influence subjects depending on their level of attachment anxiety. In another small study, intranasal OXT was effective in reducing alcohol withdrawal symptoms in alcohol-dependent subjects (Pedersen *et al*, 2013).

Recent studies by Quintana *et al* (2015, 2017) assessed the effects of a single dose of OXT (IN, 24 IU) on plasma levels up to 2 h after treatment and suggested a direct nose-to-brain transport with the IN delivery mechanism. The IN OXT administration seems to achieve therapeutic benefit, while minimizing adverse effects due to peripheral OXT activation (Bales *et al*, 2007; Bales *et al*, 2013). Overall, these findings combined with our results certainly warrant further preclinical and clinical investigation of this treatment possibility, especially since intranasal OXT has been shown to be safely tolerated by the subjects.

The present study has the following limitations. First, the radioligand, [<sup>125</sup>I]OVTA, has been used before for OXTR autoradiography in rodents and humans (Liberzon *et al*, 1994; Liberzon and Young, 1997; Olazabal and Young, 2016a,b;



Uhrig *et al*, 2016). However, some studies on non-human primates suggest significant binding to other nonapeptide receptors, most notably vasopressin 1a receptors (Toloczko *et al*, 1997; Freeman *et al*, 2014a, b). Given that we found no differences between alcoholic patients and controls in *AVPR1A* and *AVPR1B* mRNA levels, we attribute the observed differences in ligand binding to the effects on OXTR. Transcriptional levels do not universally reflect protein levels, but for OXTR we found a high concordance between these levels of assessment. Important to mention is that the selective OXTR antagonist ALS-II-69 has been recently used for competitive binding assays (Freeman *et al*, 2014b, 2017), although this compound is currently not commercially available. Second, the difference in responding under reinstatement is quite small between dependent and non-dependent groups. Examination of several OXT doses might better show potential differences in responses. Thus, the usage of a single dose of OXT represents a limitation of the study. Third, the sample size of the human pilot study is low and the fMRI results did not show a cue-induced activation in striatal regions. In a previous study (Vollstädt-Klein *et al*, 2010) using the same paradigm as here we found increased ventral striatal activation in light drinkers and more dorsal striatal activation in heavy drinkers. With regard to measures of severity of dependence (see sample description in Supplementary Information), the participants of the present study are ranging between light social drinkers and already dependent heavy social drinkers, and thus might be the reason why we neither found a ventral nor a dorsal striatal activation. Furthermore, the use of a lower dose of IN OXT (8 IU) may be more efficacious than the higher dose used in this study, at least for single administration trials (Quintana *et al*, 2017). Nevertheless, the results from the present fMRI study are in line with other independent experiments in humans and rats, but should be confirmed in a larger cohort. Fourth, only male subjects participated in the studies. This is justified by the fact that alcohol use disorders are much more prevalent in males. Nevertheless, also females are affected, and for generalization of the present results, studies in both sexes will be required. Finally, there was no direct assessment of the agonal state available by the provider of the postmortem tissues, the New South Wales Tissue Resource Centre (University of Sydney, Australia). However, tissues from this brain bank were of good quality and have been already successfully used in numerous studies (Sheedy *et al*, 2008; Meinhardt *et al*, 2013; Hirth *et al*, 2016; Hermann *et al*, 2017).

In conclusion, we here show an increase in OXTR expression (mRNA and binding sites) both in alcoholic patients and alcohol-dependent rats, as well as a decrease in *Oxt* mRNA and OXT peptide in alcohol-dependent rats. Furthermore, we could demonstrate that OXT decreased alcohol cue-reactivity in rats at the behavioral level by the cue-induced reinstatement test and in heavy social drinkers upon cue exposure at the neural level. The activity differences found in the human fMRI task, although preliminary, comprised similar brain networks as found in the mechanistic postmortem studies. Our highly translational, multilevel approach underlines the importance of the OXT system in alcohol dependence and supports current efforts made to understand the underlying mechanism and potential treatment applications for this disorder.

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The authors declare no conflict of interest.

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## AUTHOR CONTRIBUTIONS

ACH, RS, WHS, VG, and RC were responsible for study design of the rat and post-mortem brain study and procured study funding. SU performed post-mortem experiments. SU, WHS, and ACH analyzed post-mortem data. SU and ED performed behavioral animal experiments. SU, EK, and VG performed expression experiment in rats. RCF provided OXTR-2 antibodies for western blots. SU, WHS, ACH, and RS analyzed rat data. SV-K, AK, and FK were responsible for the concept and design of the fMRI study, and FK procured funding. SB and AK performed clinical experiments. SV-K analyzed the fMRI data. SU, WHS, SV-K, RS, and ACH drafted the manuscript. All authors revised the manuscript critically for important intellectual content and approved the final version.

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Supplementary Information accompanies the paper on the Neuropsychopharmacology website (<http://www.nature.com/npp>)