Association of orexin/hypocretin receptor gene (HCRTR1) with reward sensitivity, and interaction with gender

Aleksander Pulvera, Evelyn Kiiveb, Margus Kanarikc and Jaanus Harroa,c

a School of Natural Sciences and Health, Tallinn University, Narva Road 29, Astra Building, 10120 Tallinn, Estonia
b Division of Special Education, Department of Education, University of Tartu, Näituse 2, 50409 Tartu, Estonia
c Division of Neuropsychopharmacology, Department of Psychology, University of Tartu, Ravila 14A Chemicum, 50411 Tartu, Estonia

Abstract

Orexins/hypocretins maintain wakefulness, increase appetite and participate in the coordination of stress response. We have recently provided evidence on the role of orexins in aggression, showing the association of the HCRTR1 genotype (rs2271933 G>A; leading to amino acid substitution Ile408Val) with aggressiveness or breach of law in four independent cohorts. Aggressive behaviour can be reward driven and hence we have examined the association of HCRTR1 rs2271933 genotype with different aspects of reward sensitivity in the birth cohort representative Estonian Children Personality Behaviour and Health Study. HCRTR1 genotype was associated with reward sensitivity in a gender dependent manner. Male HCRTR1 A/A homozygotes had higher Openness to Rewards and the overall reward sensitivity score while, in contrast, female A/A homozygotes scored lower than G-allele carriers in Openness to Rewards. In the
total sample, aggressiveness correlated positively with reward sensitivity, but this was on account of Insatiability by Reward. In contrast, the HCRTR1 A/A homozygotes had a positive association of aggressiveness and Openness to Rewards. Experience of stressful life events had a small but significant increasing effect on both aspects of reward sensitivity, and correlated in an anomalous way with reward sensitivity in the HCRTR1 A/A homozygotes. Conclusively, the higher aggressiveness of HCRTR1 A/A homozygotes appears based on a qualitative difference in sensitivity to rewards, in the form that suggests their lower ability to prevent responses to challenges being converted into overt aggression.

**Keywords:** orexins; HCRTR1; reward sensitivity; aggressiveness; gender

1. Introduction

Orexins or hypocretins are neuropeptides first described in 1998 (Sakurai et al., 1998; de Lecea et al., 1998) that are expressed in clusters of dorsomedial and lateral hypothalamus (Broberger et al., 1998) and involved in multiple physiological functions (Sutcliffe and de Lecea, 2000; Broberger and Hökfelt, 2001; Schwartz and Kilduff, 2015; Ferrario et al., 2016), including participation in the coordination of defence response (Johnson et al., 2012). Orexin release modulates the expression and extinction of fear memories (Flores et al., 2015) and the orexin neurons are activated in acute response to a variety of stressors; however, their role in chronic stress appears complex (Sargin, 2019), leading to the hypothesis that the role for orexins in neurotransmission is promoted during aversive conditions that elicit high arousal (Berridge et al., 2010). We have recently found evidence for a role of orexins in the other side of the flight/flight
response, aggressiveness, by association of the HCRTR1 gene encoding the orexin OX1 receptor (Harro et al., 2019). The HCRTR1 gene variant (rs2271933, G1222A) in exon 7 that leads to amino acid substitution (Ile408Val) (Meerabux et al., 2005) was linked to aggressive behaviour or breach of law in four independent population samples by use of self-reports, interviews, and databases. In two population-representative birth cohort samples, male HCRTR1 rs2271933 A/A homozygotes were more aggressive than G-allele carriers. Female A/A homozygotes were also more aggressive if they had experienced higher number of adverse life events. The HCRTR1 genotype was also associated with re-occurrence of driving while impaired by alcohol and with involvement in traffic accidents (Harro et al., 2019). Another recent study has revealed that with each A-allele of the HCRTR1 rs2271933 there is less fMRI-measured activation in the inferior frontal cortex and more activation in the locus coeruleus (Gottschalk et al., 2019).

The only previous implication of orexins in human aggressiveness-related states had come from a unique study on epileptic patients that monitored release of orexin A in amygdala, and found an increase while subjects reported higher levels of anger (Blouin et al., 2013). Curiously, the highest increase of orexin levels was however related to experiencing positive emotions and during episodes of social interactions. Concluding from this and their previous animal studies showing activation of orexin neurons during positively but not during negatively motivated tasks (MacGregor et al., 2011), the authors suggested that orexin activity is necessary for positive emotion and motivation that drives social interaction.
While motivation for positive social interaction does not pose direct health risks, simultaneous association of orexin release with anger brings about a possibility that the role of orexinergetic neurotransmission in motivational activation can in occasion lead to less desirable behaviours. If resources are limited, aggression is a strategy to obtain rewards, and neurobiology of aggression and reward-related behaviour is closely intertwined (Panksepp, 1998; Aleyasin et al., 2018). Animal studies have strongly implicated orexins in reward processing (Harris et al., 2005; Moorman et al., 2017) and OX1 receptor antagonists are actively pursued as a potential avenue for treating drug addiction (Perrey and Zhang, 2020). It has been proposed that the role of orexins in reward seeking is in coordinating motivational activation when higher effort is required or external stimuli interfere with reward stimuli (James et al., 2017), and that OX1 receptor mediated signaling is an evolutionally conserved promoter of foraging (Barson, 2020). All this evidence together suggests that orexins may be involved in the emergence of aggressive impulses in the context of reward seeking, possibly in particular in social contexts. Thus, we aimed at exploring whether the HCRTRI rs2271933 genotype found to be associated with aggressiveness would also be associated with reward sensitivity. Reward sensitivity reflects the individually characteristic level of behavioural activation (Gray, 1994) and is a major component of temperament and personality reflecting the tendency to detect, pursue and derive pleasure from positive stimuli (Corr, 2009; Gray and McNaughton, 2000). It appears to emerge from neural networks with the mesotelencephalic dopaminergic circuitry at its core (Fu and Depue, 2019), and the latter receives a prominent input from the orexinergic neurons (Peyron et al.,
Because the higher aggressiveness in \textit{HCRTR1} rs2271933 A/A homozygotes had a different relationship with lifetime adversities in males \textit{vs} females, being independent of stressful life events in males, we hypothesized that reward sensitivity could contribute to the higher aggressiveness in \textit{HCRTR1} rs2271933 A/A males.

2. Results

2.1 \textit{HCRTR1} genotype and reward sensitivity in population-representative birth cohorts

Based on previous results on the association of the \textit{HCRTR1} rs2271933 genotype with aggressiveness, we had hypotetized that the relationship of the \textit{HCRTR1} rs2271933 genotype with reward sensitivity would be found in males but possibly not in females. Indeed, two-way ANOVA revealed statistically significant interaction effects for the Openness to Rewards scale and three of the four subscales, and several main effects of gender or genotype were also present. The \textit{HCRTR1} genotype was statistically significantly associated with three subscales of the Reward Openness and Insatiability Scale (ROIS): Excitement and Novelty, Social Experiences, and Giving in to Cravings. The interaction of the gender factor with genotype in case of Excitement and Novelty and the Openness to Rewards overall was largely owing to differences between male and female A/A homozygotes (Table 1; Fig 1): Male \textit{HCRTR1} A/A homozygotes had significantly higher Excitement and Novelty scores than G-allele carriers [F(5,819) = 4.10, p =0.001, partial $\eta^2=0.02$], but female \textit{HCRTR1} A/A homozygotes had lower scores than female G-allele carriers. The score of Social Experiences was generally
higher in females than in males \(F(5,819) = 2.60, p = 0.024, \text{ partial } \eta^2=0.02\) but this was not the case for female \(HCRTR1\ A/A\) homozygotes. Altogether this means that while in males each A-allele was associated with an increase in both subscales of Openness to Rewards, in females the \(HCRTR1\ A/A\)-homozygotes had lower scores than female G-allele carriers, and the overall Openness to Rewards score was significantly higher in \(HCRTR1\ A/A\) males but lower in \(HCRTR1\ A/A\) females \(F(5,819) = 2.47, p =0.031, \text{ partial } \eta^2=0.02\). Insatiability by Reward was not statistically significantly associated with the \(HCRTR1\) genotype, but its Giving in to Cravings facet was significantly higher in \(HCRTR1\ A/A\) males than male G-allele carriers \(F(5,819) = 2.45, p =0.032, \text{ partial } \eta^2=0.02\); such a difference by genotype was not present in females. The Excessive Spending facet was neither statistically significantly associated with genotype nor with gender.
Table 1. The Reward Openness and Insatiability Scale (ROIS) scores in males and females by HCRTR1 genotype (mean±SD).

<table>
<thead>
<tr>
<th></th>
<th>OR_Openness to Reward</th>
<th>OR_Social Experiences</th>
<th>IR_Excessive Spending</th>
<th>IR_Giving in to cravings</th>
<th>Openness to Reward</th>
<th>Insatiability by Reward</th>
</tr>
</thead>
<tbody>
<tr>
<td>Males</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>A/A</td>
<td>3.86±0.55*** (n=53)</td>
<td>3.56±0.69 (n=53)</td>
<td>2.45±0.73 (n=52)</td>
<td>2.85±0.62</td>
<td>3.71±0.50*</td>
<td>2.65±0.58 (n=52)</td>
</tr>
<tr>
<td>A/G</td>
<td>3.81±0.62 (n=165)</td>
<td>3.50±0.63** (n=163)</td>
<td>2.42±0.80 (n=163)</td>
<td>2.56±0.73**</td>
<td>3.65±0.53</td>
<td>2.49±0.68</td>
</tr>
<tr>
<td>G/G</td>
<td>3.63±0.63* (n=122)</td>
<td>3.46±0.69* (n=120)</td>
<td>2.35±0.75 (n=120)</td>
<td>2.59±0.67*</td>
<td>3.54±0.55</td>
<td>2.47±0.61</td>
</tr>
<tr>
<td>Females</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>A/A</td>
<td>3.46±0.64 (n=67)</td>
<td>3.53±0.65 (n=67)</td>
<td>2.62±0.90 (n=67)</td>
<td>2.72±0.70</td>
<td>3.50±0.57¤</td>
<td>2.67±0.71 (n=67)</td>
</tr>
<tr>
<td>A/G</td>
<td>3.73±0.67** (n=255)</td>
<td>3.68±0.68 (n=254)</td>
<td>2.48±0.83 (n=254)</td>
<td>2.59±0.65</td>
<td>3.71±0.58</td>
<td>2.53±0.62</td>
</tr>
<tr>
<td>G/G</td>
<td>3.65±0.66* (n=163)</td>
<td>3.64±0.75 (n=162)</td>
<td>2.47±0.89 (n=162)</td>
<td>2.73±0.74</td>
<td>3.64±0.60</td>
<td>2.60±0.70</td>
</tr>
</tbody>
</table>

OR – Openness to Reward; IR – Insatiability by Reward;
*p<0.05; **p<0.01 different from the corresponding HCRTR1 A/A group; #p<0.05; #p<0.01 different from the corresponding HCRTR1 A/G group;
*p<0.05; **p<0.01; ***p<0.001 different from the corresponding female group.

Figure 1. The OR Excitement and Novelty (A) and IR Giving in to Cravings (B) scores by gender and the HCRTR1 genotype Data are shown as mean ± SEM. OR- Openness to Rewards; IR- Insatiability by Reward. ***p<0.0001 different from the corresponding female group, #p<0.05; #p<0.01 different from the corresponding HCRTR1 A/A group.
Table 2. Pearson correlations between the Reward Openness and Insatiability Scale scores, and Life History of Aggression and Buss-Perry Aggression Questionnaire total scores by HCRTR1 genotype.

<table>
<thead>
<tr>
<th></th>
<th>OR_Excitement and Novelty</th>
<th>OR_Social experiences</th>
<th>IR_Excessive spending</th>
<th>IR_Giving in to cravings</th>
<th>Openness to Reward</th>
<th>Insatiability by Reward</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>A/A genotype (n=116)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>LHA Total</td>
<td>0.24*</td>
<td>0.12</td>
<td>0.22*</td>
<td>0.25**</td>
<td>0.21*</td>
<td>0.26**</td>
</tr>
<tr>
<td>BPAQ Total</td>
<td>0.20*</td>
<td>0.11</td>
<td>0.18*</td>
<td>0.29**</td>
<td>0.18</td>
<td>0.27**</td>
</tr>
<tr>
<td><strong>A/G genotype (n=407)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>LHA Total</td>
<td>0.07</td>
<td>0.02</td>
<td>0.23***</td>
<td>0.25**</td>
<td>0.05</td>
<td>0.27***</td>
</tr>
<tr>
<td>BPAQ Total</td>
<td>0.05</td>
<td>-0.04</td>
<td>0.44**</td>
<td>0.44**</td>
<td>0.01</td>
<td>0.44**</td>
</tr>
<tr>
<td><strong>G/G genotype (n=278)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>LHA Total</td>
<td>0.04</td>
<td>0.03</td>
<td>0.19**</td>
<td>0.20**</td>
<td>0.04</td>
<td>0.23**</td>
</tr>
<tr>
<td>BPAQ Total</td>
<td>0.01</td>
<td>-0.02</td>
<td>0.41***</td>
<td>0.53***</td>
<td>-0.02</td>
<td>0.54***</td>
</tr>
<tr>
<td><strong>All groups (n=801)</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>LHA Total</td>
<td>0.08*</td>
<td>0.04</td>
<td>0.21***</td>
<td>0.24***</td>
<td>0.07</td>
<td>0.26***</td>
</tr>
<tr>
<td>BPAQ Total</td>
<td>0.05</td>
<td>-0.01</td>
<td>0.46***</td>
<td>0.46***</td>
<td>0.02</td>
<td>0.45***</td>
</tr>
</tbody>
</table>

ROIS - Reward Openness and Insatiability Scale; OR - Openness to Reward; IR - Insatiability by Reward; LHA - Life History of Aggression questionnaire; BPAQ - Buss-Perry Aggression Questionnaire; *p<0.05; p**<0.01; p***<0.0001.

Figure 2. Overall and HCRTR1 genotype-wise divergent associations between aggressiveness and components of reward sensitivity. Panels A and D, the total sample of subjects; panels B and E, HCRTR1 A/A homozygotes; panels C and F, HCRTR1 G/G homozygotes. BPAQ - Buss and Perry Aggression Questionnaire.
2.2 Relationship of aggressiveness and reward sensitivity, and its dependence on the HCRTR1 genotype

In the whole sample, Insatiability by Reward was positively correlated with scores of both Buss-Perry Aggression Questionnaire and the interview-based Life History of Aggression (Table 2). Openness to Reward had virtually no correlation with aggressiveness in the whole sample. We were particularly interested in the correlation between aggressiveness and reward sensitivity in the HCRTR1 A/A homozygotes, because this is the group previously found to present high aggressiveness. Intriguingly, the A/A-homozygotes had a weaker positive correlation between self-rated aggressiveness and Insatiability by Reward than the G-allele carriers (in A/A homozygotes the correlation between BPAQ total score and Insatiability by Reward was statistically significantly weaker than in A/G heterozygotes, z = 1.84, p = 0.033, and in G/G homozygotes, z = 2.93, p = 0.002). In contrast, they had a positive correlation between aggressiveness and Openness to Rewards. Respective correlations in males and females were however largely similar (data not shown). An unexpected observation was made in comparison of correlations between reward sensitivity and the two measures of aggressiveness (Table 2; Fig. 2): In the HCRTR1 A/A homozygotes both self-rated and interview-based assessments correlated moderately with Insatiability by Reward and the correlations of the BPAQ total score and LHA total score with Insatiability by Reward were similar (z = 0.023, p = 0.49). In contrast, correlation of the two measures of aggressiveness with this aspect of reward sensitivity was different in G-allele carriers so that each G-allele increased the correlation with self-assessed aggressiveness but not with interview-based assessment [correlations of the two measures of aggressiveness with Insatiability by Reward
were statistically different \((z = 3.75, p < 0.0001 \text{ and } z = 5.71, p < 0.0001 \text{ for A/G and G/G, respectively})\).

2.3 Reward sensitivity and stressful life experiences, relevance to aggressiveness

Because we had previously found that in females the \(\text{HCRTR1}\) genotype effect on aggressiveness depends on exposure to stressful life events (Harro et al., 2019), the interaction effect of the stressful life events (SLE), gender and \(\text{HCRTR1}\) genotype on reward sensitivity was examined. However, no three-way interaction was found, but ANOVA revealed a statistically significant SLE main effects of stressful life events as well as the above described (2.1) interaction effect between genotype and gender on Openness to Reward. Both Openness to Rewards and Insatiability by Reward were slightly higher in the high SLE group as compared to the low SLE group \([F(1,754) = 3.92, p = 0.048, \text{ partial } \eta^2 = 0.05 \text{ and } F(1,748) = 3.92, p = 0.048, \text{ partial } \eta^2 = 0.05; \text{ data not shown}]\). Neither did the covariance analysis reveal a role for stressful life events in the genotype by gender interaction. We also examined the correlation of the dimensional SLE measure with reward sensitivity. The SLE score was positively correlated with Insatiability by Reward, but the correlations were weak (Table 3). SLE did not correlate at all with Openness to Rewards in the whole sample, but in the \(\text{HCRTR1} \text{ A/A-homozygotes} \) a weak positive correlation between SLE and Openness to Rewards, in particular the Social Experiences subscale was revealed.
3. Discussion

We have found that the HCRTR1 rs2271933 (G1222A) genotype that causes an amino acid substitution (Ile408Val) of the OX₁ receptor protein in a region that is suggested to interact with G proteins or other proteins (Thompson et al., 2014) and has been associated with aggressiveness or accident-prone law-breaching behaviour in multiple cohorts (Harro et al., 2019) is also associated with reward sensitivity. Aggressiveness belongs to the behavioural repertoire of coordinated adaptive response to environmental challenges (van Kampen, 2015) and in this response different nuclei of the hypothalamus play a critical role (Krük, 2014). Reward sensitivity can shape the emergence of aggressive behaviour through distinct pathways and, interestingly, the association of aspects of reward sensitivity and aggressive behaviour was HCRTR1 rs2271933 genotype dependent.

In a previous analysis (Pulver et al., 2020) we dissected reward sensitivity into two major components with hardly any co-variance, one representing striving...
toward reward multiplicity and another describing the fixation to a given reward. We found that only the latter, Insatiability by Reward, was associated with symptoms of ADHD and the $TPH2\ -703G/T$ polymorphism; also, only this component of reward sensitivity was associated with the ANGER factor of the Affective Neuroscience Personality Scale (Davis and Panksepp, 2011). Subsequently, we have found that Insatiability by Reward, but not Openness to Rewards, is associated with the increase of body mass index and other measures of obesity from childhood to young adulthood (Katus et al., 2020). Thus, the two aspects of reward sensitivity have substantially different consequences to behaviour. Herewith we have extended these findings to demonstrate that the trait of being strongly fixated on a particular reward is positively correlated to both self-assessed and interview-based aggressiveness, while striving toward multiple rewards, generally, is not.

A role for orexins in reward seeking has been proposed in a number of experimental studies (Tyree et al., 2018), but the results have not been uniform. In order to explain the apparent controversies, orexins have been suggested to coordinate motivational activation in effortful conditions (James et al., 2017). This would be in line with the notion that neuropeptides gain significance at higher neural activity (Hökfelt et al., 2000; Hökfelt et al., 2018). Neuroanatomical and functional circuitry studies have indeed suggested that the orexin system modulates reward processing through neural substrates that are crucial in the control of arousal (Li and de Lecea, 2020). For example, the locus coeruleus received the densest extrahypothalamic projections from the orexin neurons (Peyron et al., 1998), and predominantly express the OX1 receptors. Of note, the
*HCRTR1* rs2271933 genotype was recently found to be associated with activation of the locus coeruleus, this being higher with the A-allele in an additive manner (Gottschalk et al., 2019).

No significant gender differences have emerged in the structure of reward sensitivity or the relationship of its components with a number of personality and behavioural measures. But a large body of evidence suggests the existence of relevant gender differences in personality and behaviour. For example, men in general display higher levels of narcissism, Machiavellianism, and psychopathy (Muris et al., 2017), impulsivity (Cross et al., 2011), and are physically more aggressive than women (Björkqvist, 2018). Important gender differences in dopaminergic mechanisms mediating reward exist (Becker and Chartoff, 2019) and the present study suggests that the orexin system may contribute to the differences in the development of male and female aggressiveness through neurobiological underpinnings of reward sensitivity. Orexin neurons in the lateral hypothalamus are in bidirectional cross-talk with dopamine neurons in the ventral tegmental area (Liu et al., 2020) and dopamine can elicit opposite behavioural effects via input to orexin neurons through distinct receptor subtypes (Linehan et al., 2019). Indeed significant sex differences in the orexin system have been described, with higher orexin system activity in females and sex-specific interactions with the HPA axis (Grafe and Bhatnagar, 2020). Studies using pharmacological OX₁ receptor blockade have demonstrated that while in both male and female animals sucrose self-administration in condition of food restriction involves orexin signaling, only in males the OX₁ receptor mediated
signal is involved in cue-reinstated sucrose seeking (Cason and Aston-Jones, 2014).

Aggressive behaviour had been found to relate to the HCRTR1 rs2271933 genotype irrespective of life events in males but dependent on higher exposure to adverse events in females (Harro et al., 2019). We hypothesized that male HCRTR1 rs2271933 A/A homozygotes, with markedly higher aggressiveness, are driven by higher Insatiability by Reward. Indeed we did find significantly higher scores in one of the two sub-scales of Insatiability by Reward, and a tendency toward higher overall score of this measure in male HCRTR1 rs2271933 A/A homozygotes. We did, however, also observe in males higher scores by each A-allele in the other aspect of reward sensitivity, Openness to Rewards. Moreover, the most evident genotype effect was the difference between male and female HCRTR1 rs2271933 A/A homozygotes with regard to Openness to Rewards and, in particular, its sub-scale Excitement and Novelty, with male A/A homozygotes having the highest but female A/A homozygotes the lowest scores. In the Social Experiences subscale the male A/A homozygotes did not display any difference from other genotypes: While the behaviour of A/A males has been found as more frequently anti-social, this would refer to single acts of violationg social norms (Harro et al., 2019) while the subscale measures motivation of social exchange using items such as 'enjoys big parties' or 'is modest in socializing' (reversed item; Pulver et al., 2020).

Because the association of the Openness to Rewards with aggressiveness was not expected, we examined the correlation between aggressiveness and the
components of reward sensitivity by the *HCRTR1* rs2271933 genotype. Overall, both self-reported and interview-based aggressiveness measures were in positive correlation with Insatiability by Reward, but not with Openness to Rewards. Nevertheless, these relationships were different in the *HCRTR1* rs2271933 A/A homozygotes (who form 15% of the total sample): Only in this genotype the aggressiveness measures were related to Openness to Rewards, whereas correlation of Insatiability by Reward with the self-rated aggressiveness measure was lower than in other genotypes. Different measures of aggressiveness are in significant but only moderate correlation (Kiive et al., 2017), and the major difference in the measures obtained by the BPAQ and LHA interview, besides one being self-administered and another interview-based, is in the fact that the former includes feelings and thoughts while the latter is devoted to aggressive behaviour that has actually occurred, and quantitates its frequency. This difference is also reflected in the highest correlation of the LHA score with the Physical Aggression component of the BPAQ (Kiive et al., 2017).

Direct evidence at the biochemical level regarding functionality of *HCRTR1* rs2271933 (Ile408Val) variation is not available, but it has been suggested that the valine in the respective position could promote the phosphorylation of the immediately adjacent serine and alter signal transduction upon activation of the orexin *OX1* receptor (Meerabux et al., 2005). The *HCRTR1* rs2271933 genotype has previously been associated with migraine (Rainero et al., 2011b; Kowalska et al., 2018) and mood disorders (Rainero et al., 2011a), the ANGER factor of the Affective Neuroscience Personality Scale, and with aggressiveness (Harro et al., 2019). Interestingly, the A-allele of the *HCRTR1* rs2271933 genotype was also
associated with higher proneness to traffic accidents according to police registry and traffic insurance database, and the A/A homozygotes had the risk of recidive of drunk driving (Harro et al., 2019). The latter finding may offer insight into the role of this genetic variation and of the OX1 receptor in regulation of behaviour, as in this longitudinal study the HCRTR1 rs2271933 A/A homozygosity was associated with drunk driving only in the group with previous drunk driving record but not in control subjects or in drivers with speed limit violations. Irresponsible use of alcohol thus appears as a specific risk factor for the HCRTR1 rs2271933 A/A homozygotes, and is known to strongly correlate with manifest aggressive behaviour. In the stop-signal reaction time task, administration of a OX1-receptor antagonist reduced motivation to perform the task but did not affect inhibitory control (Wiskerke et al., 2019). This finding suggests that OX1-receptors are not contributing to aggression control through simple real-time inhibition, but intoxication by alcohol could change this. As a working hypothesis, it should be further examined whether the distinct relationship of aggressiveness with reward sensitivity in HCRTR1 rs2271933 A/A homozygotes is derived from misinterpretations and failed inhibitions while engaging in novel and exciting situations instead of the more common ground of aggressiveness in the inflexible fixation on specific rewards.

Higher exposure to adversity during childhood and adolescence increased reward sensitivity but the effect was minor and could not explain the genotype by gender interactions. Interestingly, the matrix of weak but statistically significant correlations again suggested a dissociation between HCRTR1 rs2271933 A/A homozygotes and G-allele carriers, as only in the former the
exposure to stressful events was (positively) related to Openness to Rewards, in particular to Social Experiences in young adulthood. Limited sample size did not permit a detailed mediation analysis, but speculatively a hypothesis could be put forward that in the HCRTR1 rs2271933 A/A homozygotes life stress increases strive toward social acceptance, that could increase the risk of occasionally resulting in transformation into inappropriate aggressive behaviour.

Increase in orexin A levels in amygdala was found during anger (Blouin et al., 2013) but even higher release of orexin was observed during experiencing positive emotions and in episodes of social interaction. The two aspects of reward sensitivity are differently related to not only to anger but all basic emotions: Of the facets of the Affective Neuroscience Personality Scale, Insatiability by Reward scores co-varied with ANGER, SADNESS and FEAR, but Openness to Rewards with SEEKING, PLAY and CARE (Pulver et al., 2020). In animal experiments the orexin neurons were found to fire during positively but not during negatively motivated tasks (MacGregor et al., 2011). Thus, previous studies and the present analysis converge to suggest, albeit indirectly, that the involvement of orexin in aggressive and antisocial behaviour is mediated by inappropriate interpretation of social cues. Proper function of orexin neurons is crucial to appropriately timed transitions between arousal states, and for taking context-specific actions (Burdakov, 2020). The major genotype by gender interaction with regard to openness to rewards, especially excitement and novelty, may help to explain the difference between expression of aggressiveness in male and female HCRTR1 rs2271933 A/A homozygotes. Higher striving towards exciting and novel situations can expose the male A/A homozygotes to
situations that provoke aggression. In females, this risk appears as being compensated for if not exposed to adversities. Nevertheless, this study is limited in collecting relevant data in young adulthood and thus any hypotheses regarding developmental strategies remain speculative; also the sample size does not support a detailed analysis given the prevalence of the genotype of particular interest. Another limitation concerns the novel reward sensitivity instrument that was developed post hoc using a large item pool in a Fennic language, and requires further characterization together with related instruments. The strength of the study however includes the exceptional representation of eligible population-based cohorts and uniform data collection procedure in a laboratory setting.

Conclusively, male HCRTR1 rs2271933 A/A homozygotes, previously found to display higher aggressiveness, reported higher striving towards multiple rewards, whereas A/A females, previously found to show higher aggressiveness only in association with severe life events, were less open to excitement and novelty. These findings support the role of orexins in reward sensitivity and the notion of different pathways to aggressiveness in males and females.

4. Method

4.1. Sample

This study was carried out on the Estonian sample of the European Youth Heart Study (1998/1999), which was subsequently incorporated into the longitudinal Estonian Children Personality Behaviour and Health Study (ECPBHS). The rationale and procedure of sample formation, and further data collection waves
have been described elsewhere in detail (Harro et al., 2001; Harro et al., 2019). ECPBHS is highly representative of two birth cohorts of a local population, as 79% of subjects of the randomized regional sample participated in the original data collection. All the subjects are of European descent. Data collection for the presented analyses was conducted at ages 25 (both cohorts) and 33 (only the older cohort). The original size of the total sample is n=1238, but all data necessary for the analyses presented in this paper was n=818 to 825. This study was carried out in accordance with the code of ethics as per Declaration of Helsinki, it was approved by the Ethics Review Committee on Human Research of the University of Tartu, and written informed consent was obtained from all the participants.

4.2. Assessment of reward sensitivity

In order to take into account the inner structure of reward sensitivity, the Reward Openness and Insatiability Scale (ROIS) was used to assess reward sensitivity (Pulver et al., 2020). In brief, ROIS comprises 28 items equally divided between two higher-order factors, Openness to Rewards and Insatiability by Reward, that describe striving toward multiplicity of rewards and excessive fixation to particular reward, respectively. Openness to Rewards includes two facets, Excitement and Novelty, and Social Experiences. Insatiability by Reward comprises Excessive Spending and Giving in to Cravings. While items belonging to the two factors were the product of an exploratory principal component analysis, the factors do not correlate significantly with each other, whereas the facets that form the factors correlate with each other moderately (r=0.45 and
0.48). ROIS scores were available for 818 or 825 participants for the two subscales.

4.3. Measures of aggressiveness

4.3.1. Self-report questionnaire

The 29-item Buss-Perry Aggression Questionnaire (BPAQ; Buss and Perry, 1992) assesses four aspects of aggressive behaviour: Physical aggression, Verbal aggression, Anger, and Hostility. Participants rated each statement on a 5-point Likert Scale (uncharacteristic=1, characteristic=5). Data of Buss-Perry Aggression Questionnaire were available for 926 subjects.

4.3.2. Aggressiveness assessment by interview

The Life History of Aggression interview (LHA; Coccaro et al., 1997) was carried out by an experienced clinical psychologist. Items were scored only for the history of actual behavior. LHA has three subscales: Aggression (temper tantrums, verbal aggression, indirect aggression, non-specific fighting, and physical assault against people); Consequences/Antisocial Behaviour (school disciplinary problems, problems with supervisors, antisocial behaviour not resulting in police involvement, and antisocial behaviour involving the police); and Self-Directed Aggression (self-injurious behavior, and suicide attempts). Each item was rated on a 5-point scale, ranging from 0=No events to 5=More events than can be counted. Data of LHA were available for 922 subjects.
4.4. Assessment of stressful life experiences

Stress experienced in childhood was measured at age 15 with a list of 21 adverse life experiences reported by participants or their parents, which included questions about parental death and divorce/separation, absence of both parents, unemployment in the family, financial problems and poverty in the household, poor living conditions, poor health and chronic diseases, serious illness of a family member, death of a close relative, trauma, fear of school, bullying at school and humiliation at home (Laas et al., 2014). The events were recorded as dichotomous variables (present or not present) and were then counted to form the number of experienced stressful life events (SLE). For group-wise comparison, two categories of the SLE (Low SLE, 0-2 events and High SLE, >2 events) were constructed on the basis of median value of SLE score ($Md = 2$). Data were available for 557 subjects in the ECPBHS younger cohort and for 594 subjects in the older cohort.

4.5 Genotyping HCRTR1 rs2271933

This was carried out as previously described (Harro et al., 2019). Genomic DNA was extracted from venous blood samples using Qiagen QIAamp® DNA Blood Midi Kit. The real-time polymerase chain reaction (RT-PCR) for genotyping the HCRTR1 rs2271933 polymorphism was performed using a TaqMan Pre-Designed SNP Genotyping Assay (Applied Biosystems; Foster City, CA, USA) C__15961465_10 containing primers and fluorescent probes. Genotyping reactions were performed in a total volume of 10 µl with ~25 ng of template
DNA. RT-PCR reaction components and final concentrations were as follows: 1:5
5 x HOT FIREPol® Probe qPCR Mix Plus (ROX) (Solis BioDyne) and 1:20 80 x
TaqMan Primers Probe. Context sequence [VIC/FAM] was as follows:
CTTGTCCCTTGCAGAGCGATGCTCC[A/G]TCTCCAAAATCTCTGAGCATGTGGT.
Reactions were performed on the Applied Biosystems ViiA™ 7 Real-Time PCR
System. The amplification procedure consisted of an initial denaturation step at
95 °C for 12 min and 40 cycles of 95 °C for 15 s and 60 °C for 1 min. Positive and
negative controls were added to each reaction plate. No inconsistencies
occurred. Genotyping was performed blind to all phenotypic data. Allele
frequencies agreed with National Center for Biotechnology Information database
(minor allele frequency around 0.4;
https://www.ncbi.nlm.nih.gov/snp/rs2271933#frequency_tab) and published
reports. All the subjects were of European origin, and genotype frequencies of all
samples were in Hardy-Weinberg equilibrium.

4.6. Statistical analysis
Statistical analysis was carried out using the SPSS v. 25 software. Two-way
between-groups analyses of variance (ANOVA) were performed to investigate
gender and genotype interactions in reward sensitivity. The dependent variables
included the Reward Openness and Insatiability Scale (ROIS) summary score, the
Openness to Reward (OR) scale and its sub-scales Excitement and Novelty
subscale and Social experiences, and the Insatiability by Reward (IR) scale and its
sub-scales Excessive spending and Giving in to cravings.
The independent variables (fixed factors) were the HCRTR1 rs2271933 genotype
(A/A, A/G, G/G) and gender. A three-way between-groups univariate analysis of
variance (ANOVA) was also performed to investigate the stressful life experience, gender and genotype interactions in reward sensitivity. Two-way analysis of covariance (ANCOVA) was used to examine the possible covariate effect of stressful life events on the association between genotype, gender and reward sensitivity. Preliminary assumption testing was conducted to check for normality, outliers, homogeneity of variance, with no serious violations noted. In case of significant ANOVA interaction effects the differences between means of the 6 groups (two gender levels by three genotype levels) was investigated by Fisher LSD post hoc test. Associations between the test scores were assessed by Pearson correlation. The z-statistic was used for comparison of correlations. Analysis was performed and data are presented as the mean item score (i.e., sum of the items is divided by number of items in scale). In the statistical analysis, the conventional 5% level was used to assess the significance.

Acknowledgements

We are grateful to the ECPBHS study participants, their parents and the whole ECPBHS Team. This work was supported by Estonian Research Council Project IUT20-40, the EC Horizon 2020 project CoCA (H2020-PHC-2015-667302), and the Tallinn University ASTRA project TU TEE financed by the European Union European Regional Development Fund (2014-2020.4.01.16-0033).

Declarations of Interest: None.

References


