1	Title

- 2 The influence of angiotensin converting enzyme and bradykinin receptor B2 gene variants on
- 3 voluntary fluid intake and fluid balance in healthy men during moderate intensity exercise in
- 4 the heat.
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Angiotensin converting enzyme (ACE) and bradykinin receptor B2 ( $B_2R$ ) genetic variation 27 28 may affect thirst due to effects on angiotensin II production and bradykinin activity 29 respectively. To examine this, 45 healthy Caucasian men completed 60 minutes of cycle exercise at  $62 \pm 5\%_{2peak}$  in a room heated to  $30.5 \pm 0.3$ °C with *ad libitum* fluid intake. Blood 30 31 samples were collected pre-, mid-, and immediately post-cycle. Fluid intake, body mass loss (BML), sweat loss (determined via changes in body mass and fluid intake) and thirst 32 sensation were recorded. All participants were genotyped for the ACE insert fragment (I) and 33 the  $B_2R$  insert sequence (P). Participants were homozygous for the wild type allele (WW or 34 MM), heterozygous (WI or MP) or homozygous for the insert (II or PP). No differences 35 between genotype groups were found in mean ( $\pm$  SD) voluntary fluid intake (*WW*: 613  $\pm$  388, 36 37 *WI*: 753 ± 385, *II*: 862 ± 421 mL, *P* = 0.31; *MM*: 599 ± 322, *MP*: 745 ± 374, *PP*: 870 ± 459 mL, P = 0.20), percentage BML or any other fluid balance variables for both the ACE and 38  $B_2R$  genes, respectively. Mean thirst perception in the  $B_2R PP$  group, however, was higher (P 39 < 0.05) than both MM and MP at 30, 45 and 60 minutes. In conclusion, the results of this 40 study suggest that voluntary fluid intake and fluid balance in healthy men performing 60 41 42 minutes of moderate intensity exercise in the heat are not predominantly influenced by ACE or  $B_2R$  genetic variation. 43

44

## 45 Key Words

46 Thirst; Angiotensin converting enzyme; Bradykinin receptor B2; Dipsogenic; Exercise; Fluid
47 Balance

Page 3 of 27

#### 48 Introduction

It has long been observed that the majority of humans performing physical work or exercising do not voluntarily drink enough water to replace sweat losses despite adequate fluid supply (Pitts *et al.* 1944). This phenomenon has been termed 'voluntary dehydration' (Rothstein *et al.* 1947) and has been found to frequently occur in athletes (Maughan *et al.* 2004; Passe *et al.* 2007) and the general population even in favourable conditions for fluid intake. Based on these observations, it is apparent that relying on thirst mechanisms does not always guarantee sufficient total water intake (Armstrong and Grandjean 2007).

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Intracranial and systemic administration of physiological doses of angiotensin II, a 57 component of the renin-angiotensin aldosterone system (RAAS), has been shown to promote 58 59 drinking in animals (Fitzsimons 1972) and humans (Fitzsimons 1998). Angiotensin II is produced from the cleavage of angiotensin I by angiotensin converting enzyme (ACE). A 287 60 base pair (bp) Alu repeat insert within intron 16 of the ACE gene has been identified as a 61 common allelic variant (Rigat et al. 1990) unique to humans (Montgomery et al. 2002). The 62 wild type allele (W) is significantly associated with both higher tissue (Danser et al. 1995), 63 and serum (Rigat et al. 1990) ACE activity. 64

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Another known function of ACE is the degradation of vasodilator kinins within the kallikrein kinin system, particularly bradykinin (Dzau *et al.* 1988). Bradykinin exerts its effects on bradykinin B2 receptors (B<sub>2</sub>R) and has been shown to be a powerful dipsogen. Endogenous bradykinin (Cadnapaphornchai *et al.* 2004) and exogenous infusion of bradykinin (Fregly and Rowland 1991), during acute ACE inhibition has been observed to cause polydipsia and polyuria. Furthermore, the antagonism of bradykinin receptors reverses such effects (Cadnapaphornchai *et al.* 2004). An allelic variant containing a nine bp repeat insertion (*P*) has been identified in exon 1 of the gene encoding for  $B_2R$  (Braun *et al.* 1996). The wild type allele (*M*) is significantly associated with higher gene transcriptional and receptor activity (Braun *et al.* 1996; Lung *et al.* 1997). It is possible, therefore, that variants in *ACE* and  $B_2R$  genes may affect thirst and fluid intake in humans. Understanding the potential role of these, and other, dipsogenic gene variants on thirst and fluid intake may explain some of the large differences in these variables that are often observed in athletes and recreational exercisers.

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To date, only one study has previously investigated the influence of these allelic variants on thirst and voluntary dehydration. Saunders *et al.* (2006) investigated the association of these allelic variants with weight changes in 428 Caucasian male ironman triathletes during competition. The authors concluded that the  $B_2R PP$  genotype was associated with greater weight loss during the ironman triathlon with a significant linear trend for the distribution of this genotype across weight loss groups. On the other hand, no association was found with *ACE* genotype.

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The aim of the present study was to conduct a controlled laboratory study investigating the influence of *ACE* and  $B_2R$  allelic variations on thirst, voluntary fluid intake and fluid balance during moderate intensity exercise in the heat. It was hypothesized that individuals homozygous for the wild type *ACE* or  $B_2R$  allele would drink more fluid during exercise in the heat.

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95 Materials and methods

96 Participants

Forty-five Caucasian men aged 18-45 years (mean  $\pm$  SD, age 28  $\pm$  8 y, height 178.3  $\pm$  6.7 cm, 97 body mass  $78.93 \pm 14.29$  kg, body mass index  $24.74 \pm 3.59$  kg.m<sup>-2</sup>, and peak oxygen uptake 98  $50.95 \pm 10.10 \text{ ml.kg}^{-1}$ .min<sup>-1</sup>) volunteered to participate in the present investigation. All 99 participants were healthy non-smokers, not concurrently taking medication, and not 100 101 supplementing with creatine. Verbal and written explanations of the experimental procedures were given before participants provided written consent and completed a medical screening 102 103 questionnaire. The study had prior approval from the Manchester Metropolitan University's Faculty of Science and Engineering ethical advisory committee, in accordance with the 104 105 Declaration of Helsinki.

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## 107 Preliminary trials

Each participant completed two preliminary trials, separated by a minimum of 48 hours, prior 108 to their main experimental trial. In the first of these, peak oxygen uptake (2peak) was assessed 109 through the performance of a continuous, incremental cycle ergometer (Lode Excalibur Sport, 110 Groningen, Netherlands) test to volitional fatigue in a thermoneutral environment (21.4  $\pm$ 111  $1.0^{\circ}$ C;  $33 \pm 7\%$  relative humidity). Pulmonary gas-exchange (Cosmed K4b<sup>2</sup>, Italy) and heart 112 rate (HR) (Polar FS2c, Kempele, Finland) were continuously analysed and recorded using a 113 breath-by-breath system. Rating of perceived exertion (RPE) (Borg, 1998) was obtained 114 every 2 minutes. Utilising the rolling mean of 10 breaths and work rate, the work-rate to 115 oxygen uptake (2) relationship was determined. The highest mean value of the rolling 10 116 breath oxygen consumption values was accepted as 2peak. The desired work rate eliciting 55% 117 of <sub>2peak</sub> was then calculated, with account taken for the mean response time for <sub>2</sub> during ramp 118 119 exercise (i.e. two thirds of the ramp rate was deducted from the calculated work rate) (Whipp 120 *et al.* 1981).

Page 6 of 27

The second preliminary trial involved familiarising the participants with the experimental trial protocol. All procedures undertaken in the main trial, described in detail below, were performed in a room heated to 30°C, with the exception of body mass (BM) weighing and urine and blood sampling.

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## 127 Experimental protocol

In the 24 h period preceding the experimental trial, participants refrained from strenuous exercise and abstained from alcohol and caffeine ingestion. In addition, participants were also asked to fast from 2100 hours the evening before the experimental trial, and to consume 500 mL of water approximately 90 minutes prior to arrival at the laboratory in an effort to ensure an adequate level of hydration.

133 All participants reported to the laboratory for their experimental trial between 0800 and 1000 hours. Upon arrival at the laboratory, participants were asked to completely empty 134 their bladder into a container from which a 5 mL urine sample was retained for future 135 analysis. Participants were then seated in an upright position for 15 minutes in an air 136 conditioned environment maintained at approximately 20°C before a 9 mL blood sample was 137 obtained by venous puncture of an antecubital vein. Mid- and post-exercise blood samples 138 were taken with the participants seated on the cycle ergometer in an upright position in an 139 140 attempt to reduce the influence posture has on circulating blood and plasma volumes (Hagan 141 et al. 1978). During the 15 minute seating period, resting HR and thirst sensation (on adapted 142 Borg scales) were recorded.

Following the pre-exercise blood sample, participants were weighed nude to the nearest 0.01 kg (Adam Equipment Co Ltd., GFK 150, Milton Keynes, UK) before initiation of the exercise protocol. Participants cycled at a constant work-rate equivalent to an initial load of 55% of <sub>2peak</sub> for 60 minutes on an electronically braked cycle ergometer (Lode

Page 7 of 27

Excalibur Sport, Groningen, Netherlands), with a three to five minute rest interval at 30
minutes for blood sample collection. Breath-by-breath expired air samples, analysed for
oxygen uptake and HR, were collected for 3 minutes pre-exercise and at every 15 minutes of
cycling. Ratings of perceived exertion and thirst sensations were recorded pre-exercise then
every 15 minutes of cycling. Post-exercise nude BM was recorded after participants towel
dried.

153 During the 60 minute cycle, participants were provided with constant access to a commercially available hypotonic sports drink containing 2% carbohydrate. The participants 154 155 were given a choice of two flavours and asked to consume ad libitum. Drinks were provided in specialised drinks bottles (CamelBak Better Bottle<sup>TM</sup>) kept in opaque insulating sleeves to 156 prevent participants from viewing the volume of fluid contained and aimed to maintain drinks 157 at regular room temperature. Drink bottles were also changed every 15 minutes. The volume 158 of fluid consumed was determined by weighing of drinks bottles. A 5 mL sample of the drink 159 was obtained for analysis. Total sweat loss was calculated through change in BM corrected 160 for fluid intake. For the purposes of this investigation, it was assumed that all mass loss 161 during exercise was due to sweating since respiratory water loss and mass loss due to 162 substrate exchange would have accounted for only a small component of total mass loss 163 (Maughan et al. 2007). 164

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#### 166 Biochemical analysis

Drink and urine samples were stored at 4°C until analysis for osmolality by freezing point depression (Gonotec Osmomat 030 Cryoscopic Osmometer; Berlin, Germany). EDTA blood samples were analysed for haemoglobin concentration by the cyanmethaemoglobin method, packed-cell volume by microcentrifugation). The haemoglobin and packed-cell volume values were used to estimate percentage changes in blood, erythrocyte and plasma volumes, 172as described by Dill and Costill (1974). Remaining whole blood samples were stored at -80°C173until DNA extraction. Serum tube blood samples were centrifuged at 1500 g for 15 min at1744°C before serum was removed and kept at 4°C until the analysis of osmolality using the175method described previously. All analyses were performed in duplicate, with the exception of176the packed-cell volume measurements, which were made in triplicate.

177

## 178 Genotyping

179 Genomic DNA was extracted from 5 mL of EDTA whole blood using Qiagen FlexiGene

180 DNA Kit (West Sussex, UK) according to the manufacturer's instructions. The participants

181 were genotyped for the W/I allelic variants within intron 16 of the ACE gene, and the M/P

allelic variants within exon 1 of the  $B_2R$  gene. For each gene to be assayed, primers unless

183 stated, were designed using publicly available genomic sequences obtained through GenBank

184 (http://www.ncbi.nlm.nih.gov/genbank/) and the public domain primer design software

185 Primer3 (http://frodo.wi.mit.edu/primer3/; Rozen and Skaletsky 2000). Target gene sequences

to be amplified within whole gene sequences were identified from published literature.

187 Primers were designed with Primer3 using the human mispriming library.

188 ACE genotype was ascertained using two separate PCR reactions. Firstly, the

189 following forward (5'-GGGGACTCTGTAAGCCACTG-3') and reverse (5'-

190 TCGGGTAAAACTGGAGGATG-3') primers were used to detect the presence of a *W* allele.

191 The presence of a *W* allele resulted in an approximate 300 bp PCR product whilst the absence

of a *W* allele resulted in no PCR products being formed. Due to the preferential amplification

193 of the *W* allele observed with the above primers and furthermore owing to the preferential

amplification of the W allele in heterozygous samples (Shanmugam et al. 1993), a second

195 insertion specific PCR was performed on all samples using the following forward (5'-

196 TGGGACCACAGCGCCCGCCACTAC-3') and reverse (5'-

TCGCCAGCCCTCCCATGCCCATAAC-3') primers as previously utilised by Settin et al. 197 (2009). The presence of an *I* allele resulted in an approximate 350 bp PCR product whilst for 198 samples homozygous for the W allele, no products were visualised. The PCR reactions were 199 carried out in a total volume of 20 µl containing 20 ng of DNA, 1 x NH<sub>4</sub>, 1.5 mM MgCl<sub>2</sub>, 25 200 µM each of dATP, dTTP, dCTP and dGTP, 250 nM of each primer, and 0.02 units of Taq 201 DNA polymerase. For the first ACE PCR reaction, the PCR conditions consisted of an initial 202 203 denaturing step at 95°C for 5 min; followed by 10 cycles of denaturing at 95°C for 30 seconds, annealing for 30 seconds at 58°C and extension for 45 seconds at 72°C; then 30 204 205 cycles of denaturing at 89°C for 20 seconds, annealing for 30 seconds at 58°C and extension for 45 seconds at 72°C; and a final extension step at 72°C for 5 min. The ACE insertion 206 specific PCR conditions consisted of these identical steps except for an annealing temperature 207 of 59°C. The amplified fragments were resolved by electrophoresis on 2% agarose gel stained 208 with ethidium bromide and visualised under ultra violet (UV) light. 209  $B_2R$  genotype was ascertained using the following forward (5'-CACT 210 CCAGCTCTGGCTTCTG-3') and reverse (5'-TTCAGTCGCTCCCTGGTACT-3') primers 211 to produce approximately 100 and/or 90 bp fragments. The PCR reactions were carried out in 212 volumes and concentrations as above. The PCR conditions were also as above except for an 213 annealing temperature of 55°C. The amplified fragments were subsequently resolved on 4.5% 214 microfagorose gel stained with ethidium bromide and visualised under UV light. All 215 fragments were sized using a molecular weight marker (Hyperladder<sup>TM</sup> V, Bioline, UK). 216 217

## 218 Statistical analysis

Differences in participant characteristics, environmental conditions during exercise,
 and pre-exercise hydration status between genotype groups were examined using one-way
 ANOVA. One-way ANOVA were also used to examine differences in percentage BML, fluid

intake, sweat loss, sweat rate, and drink osmolality between genotype groups. Significant F-222 tests were followed by multiple independent Student's *t*-tests. The false discovery rate 223 224 procedure was applied to control for type 1 error. Pre and post BM data within groups was analysed using paired Student's *t*-test. Differences in <sub>2</sub> and subjective measures during the 60 225 minute submaximal exercise between genotype groups were determined using two-way 226 227 mixed model ANOVA (time x group). Two-way mixed model ANOVA (time x group) were 228 also used to examine differences in serum osmolality, blood volume, and plasma volume. Sphericity for repeated measures was assessed and where appropriate, Greenhouse-Geisser 229 230 corrections were applied for epsilon <0.75, and the Huynh-Feldt correction adopted for less severe asphericity. Significant *F*-tests were followed with one-way ANOVA at time points 231 and multiple independent Student's t-tests. The false discover rate procedure was applied to 232 control for type 1 error. All data were analysed using SPSS for Windows version 14.0 233 234 (Chicago, IL). Statistical significance was accepted at the 5% level and results presented as mean  $\pm$  standard deviation (SD). 235

236

237 Results

238

239 Subject characteristics and environmental conditions

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The genotype distributions for *ACE* and  $B_2R$  of the participants in this study were in Hardy-Weinberg equilibrium. Participant characteristics and exercise conditions grouped according to genotype are presented in Table 1 (*ACE*) and Table 2 ( $B_2R$ ). Genotype groups for both variants were not different in age, height, body mass, BMI, <sub>2peak</sub>, average exercise intensity or pre-exercise urine osmolality (P > 0.05). Six participants were classed as dehydrated at the start of exercise with pre-exercise urine osmolalities of >900 mOsm.kg<sup>-1</sup>. There were no

247	differences in the environmental exercise conditions between the ACE genotypes. However,
248	for $B_2R$ , ambient temperature for the <i>MP</i> genotype group was significantly higher than the
249	<i>PP</i> group ( $P < 0.05$ ). Relative humidity was also significantly higher for <i>PP</i> compared to
250	<i>MM</i> and <i>MP</i> ( $P < 0.05$ ). Data is presented in Tables 1 and 2.
251	
252	Mean exercise intensity and HR significantly increased over time ( $P < 0.05$ ) but no
253	interaction effect or main effect of genotype was observed in any of these variables ( $P > 0.05$ )
254	for both variants.
255	
256	Fluid intake and fluid balance
257	ACE
258	Both the WW and WI genotype groups exhibited a significant decrease in body mass at the
259	end of the 60 minute cycle. The <i>II</i> group also tended to a significant decrease in body mass
260	(P = 0.07). There were no differences in mean percentage BML, fluid intake, and total sweat
261	loss (Table 1).
262	
263	$B_2R$
264	A significant decrease in body mass occurred in all three genotype groups at the end of 60
265	minute cycle ( $P < 0.05$ ). There were no differences in mean percentage BML, fluid intake,
266	and total sweat loss between genotypes
267	
268	Blood variables
269	ACE
270	Plasma volume significantly decreased by $9.1 \pm 5.5$ , $6.0 \pm 7.8$ and $7.8 \pm 6.6\%$ from pre to
271	post for WW, WI, and II, respectively. No interaction effect ( $P = 0.61$ ) or main effect of

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genotype (P = 0.91) was found. Serum osmolality (Figure 1a) significantly increased over
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time (P < 0.05) but no interaction effect (P > 0.05) or main effect of genotype (P > 0.05) was present.

275

276  $B_2 R$ 

- 277 Plasma volume significantly decreased by  $6.0 \pm 8.4$ ,  $6.8 \pm 6.0$  and  $9.0 \pm 6.7\%$  from pre to
- post for *MM*, *MP*, and *PP*, respectively. No interaction effect (P = 0.69) or main effect of
- 279 genotype (P = 0.51) was found. Serum osmolality (Figure 1b) significantly increased over
- time (P < 0.05) but no interaction effect (P > 0.05) or main effect of genotype (P > 0.05) was
- 281 present.

282

- 283 Subjective measures
- 284 *ACE*
- Ratings of perceived exertion and thirst (Figure 2a) significantly increased over time ( $P \le P$
- 286 0.05). However, no interaction effects (P = 0.92, P = 0.81) or main effects of genotype (P =

287 0.57, P = 0.83), were found for these subjective measures, respectively.

288

- 289  $B_2R$
- 290 Ratings of perceived exertion and thirst (Figure 2b) significantly increased over time (P <

291 0.05). No interaction effect (P = 0.22) or main effect of genotype (P = 0.20) were found for

292 RPE. Both an interaction effect and main effect of genotype (P < 0.05) was found for thirst,

- however. Thirst was significantly higher (P < 0.05) in the PP genotype group compared to
- both MM and MP in the latter half of the 60 minute cycle at 27-30 min, 42-45 min, and 57-60
- 295 min.
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Page 13 of 27

#### 297 Discussion

The primary finding of the present investigation was that both ACE and  $B_2R$  allelic variation 298 did not significantly influence voluntary fluid intake or fluid balance during moderate 299 intensity exercise in the heat. All six genotype groups exhibited indications of the voluntary 300 dehydration phenomena however, with five of the groups manifesting decreases in body mass 301 from pre-exercise that reached significance. Furthermore, the pattern of results surprisingly 302 303 appeared the reverse of that expected. Participants with the ACE II genotype and participants with the  $B_2R$  PP genotype did not exhibit the greatest level of voluntary dehydration but 304 305 rather exhibited the intermediate and lowest level of dehydration, respectively. 306

With regards to *ACE*, the findings of the present study lend support to those of Saunders *et al.*(2006) who reported no significant association of *ACE* genotype and weight change in
competing Caucasian male ironman triathletes. The present study was the first controlled
laboratory study to be conducted that confirms this previous finding to the wider population
of healthy Caucasian males partaking in a period of moderate intensity exercise in the heat.

It is acknowledged that a limitation of this study is sample size and that more participants should be investigated or added to the dataset for firm conclusions to be drawn. However, it is worth noting that previous studies such as de Souza *et al.* (2013), Folland *et al.* (2000), Santana *et al.* (2011) and Williams *et al.* (2011) have found positive *ACE* results/associations with sample sizes smaller than in the present study (range n = 27 to n = 41).

318

319 Given the influences of *ACE* genotype on inter-individual variation in circulating ACE

320 concentrations (Day et al. 2007; Rigat et al. 1990), it was theorised that ACE allelic variation

321 may have a resulting effect on angiotensin II concentrations and thus thirst and fluid intake

Page 14 of 27

during exercise. The results of the present study suggest that this may not be the case. Instead,
other rate limiting steps in the production of angiotensin II or its activity such as angiotensin I
concentration or angiotensin II receptor activity may be responsible for inter-individual
differences in thirst, voluntary fluid intake and voluntary dehydration.

326

It may be that the RAAS plays little role in the regulation of fluid intake and balance during exercise in the heat. Greenleaf *et al.* (1983) have suggested the RAAS is influenced by reductions in total body water and plasma volume is the predominant stimulus to thirst during exercise in the heat. Maresh *et al.* (2004), on the other hand, suggest that the RAAS has a minor role whilst the AVP system has a dominant role in regulating thirst during exercise heat challenge. The lack of influence of *ACE* genotype on thirst and voluntary fluid intake found in the present study appears to support the conclusion of Maresh *et al.* (2004).

334

The greater role and importance of the AVP system in regulating thirst and fluid balance 335 336 during exercise in the heat is supported by serum osmolality data presented in this investigation. In healthy humans, AVP is released in response to small increases in blood 337 plasma osmolality. Mid-exercise serum osmolality in the present study increased a significant 338 amount over time to elicit enhanced AVP secretion and stimulated thirst response. It is also 339 340 worth noting that only three participants exhibited a serum osmolality above 295 mOsm.kg<sup>-1</sup> 341 at any point of measurement, suggesting that serum osmolality was generally well regulated within the normal range of 280-295 mOsm.kg<sup>-1</sup> when free access to fluid was available. 342 343

The  $B_2R$  results of the present study are in disagreement to the findings of Saunders *et al.* (2006) who found a significant linear trend for the distribution of the *PP* genotype across three percentage weight loss groups. The lack of genotype influence obtained in the present

Page 15 of 27

study compared to Saunders et al. (2006) is likely due to the difference in study population 347 and exercise stimulus. With a few exceptions, the group of healthy participants who 348 volunteered for the present study were active recreational exercisers who were relatively fit. 349 The 60 minutes of moderate intensity exercise at 30°C and approximately 30% relative 350 humidity would have therefore been no doubt challenging and hard, but comparably less 351 physiologically challenging than an ultra-endurance event lasting 10-15 h in a hot and humid 352 353 climate. It may therefore only be in such extreme situations that an influence of  $B_2R$  variation is observed. 354

355

Despite the absence of a significant influence of genotype for voluntary fluid intake and fluid 356 balance in the present study, subjective perception of thirst was significantly higher in the PP 357 genotype group compared to the others in the latter half of the 60 minute cycle. Since serum 358 osmolality was regulated within normal range, there was no significant difference in serum 359 osmolality and no significant differences in blood or plasma volume decrease between 360 genotypes, it is unlikely that the excess thirst experienced by this group was either osmotic or 361 hypovolemic thirst due to a greater physiological need for water. This finding supports the 362 perception that thirst in humans is not solely influenced by physiological need but other 363 factors such as habit and psychology may also play a role. The results of the other subjective 364 measures are suggestive towards this. 365

366

Sweat loss during this study (approximately 1.1 L/h) was relatively low compared to other studies. Byrne *et al.* (2006) observed a sweat rate of  $1.47 \pm 0.34$  L/h in runners participating in a half marathon in environmental conditions of 25.6 - 27.3 °C and a relative humidity of 75 - 90%. Similarly, Lee *et al.* (2010) observed a sweat rate of  $1.45 \pm 0.32$  °C during a run at 26.4 °C and 81% humidity and O'Neal *et al.* (2014) observed a sweat rate of  $1.35 \pm 0.4$  L/h

during a run at  $20 \pm 3$  °C and  $54 \pm 14\%$  humidity. This difference could be attributed to the 372 relatively low humidity in the present study. Rate of fluid intake amounted to  $0.37 \pm 0.26$  L/h 373 and  $0.075 \pm 0.062$  L/h in the studies by Byrne *et al.* (2006) and Lee *et al.* (2010) which are 374 much lower than in the present study (approximately 0.74 L/h). A likely reason for this 375 difference in ingestion rate is the mode of exercise. Cycling allows for greater ingestion due 376 to easier access to fluid and also less likely to suffer from gastrointestinal comfort compared 377 378 to running. Furthermore, fluid intake is highly variable between individuals and subject to the effect of a number of variables including palatability of fluid provided (Minehan et al. 2002). 379 380 A greater volume of fluid is ingested during exercise if the drink is considered to be acceptable in comparison to less acceptable or to water (Passe et al. 2000). Similarly, greater 381 fluid intake during exercise is observed with ingestion of a carbohydrate electrolyte solution 382 compared to water and other solutions that are rated less palatable (Passe et al. 2004; Rivera-383 384 Brown et al. 1999). Some of the difference in fluid ingestion volume may therefore be attributed to greater subjective acceptability and palatability of the flavoured 2% 385 carbohydrate drink provided in this present study compared to the water and 6% carbohydrate 386 electrolyte solutions provided in the studies by Byrne et al. (2004) and Lee et al. (2010). 387 388

In conclusion, the results of the present study suggests that ACE and  $B_2R$  allelic variation do 389 not play a major role in governing voluntary fluid intake and fluid balance in normal healthy 390 391 males partaking in a 60 minute period of moderate intensity exercise in the heat. It appears that other factors such as subjective feeling and subjective perception of the exercise 392 environment may be more important in determining thirst and fluid intake in exercise of this 393 394 nature and in this population. Additional or a larger number of participants should be 395 investigated whilst future work should also examine the responses of hormones and enzymes involved in body fluid regulation in order to elucidate the underlying mechanism involved 396 397 and attempt to explain the observed individual differences in thirst, fluid intake and fluid balance. References 398

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# 549 Tables

550

551 Table 1: Participant characteristics, environmental conditions during exercise, and fluid

balance grouped according to angiotensin converting enzyme (ACE) genotype. WW;

homozygous wild type, *WI*; heterozygous, *II*; homozygous insertion. Values are mean  $\pm$  SD.

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Variable	WW	WI	II	P-value
n	12	21	12	-
Age (yr)	$30 \pm 9$	$27\pm8$	$28 \pm 7$	0.50
Height (cm)	$177.2 \pm 5.9$	$178.6\pm6.1$	$178.8\pm8.7$	0.82
Body Mass (kg)	$76.73 \pm 14.84$	$82.79 \pm 15.15$	$74.39 \pm 11.10$	0.22
BMI (kg.m <sup>-2</sup> )	$24.31\pm3.54$	$25.87 \pm 3.94$	$23.19\pm2.38$	0.11
<sub>2peak</sub> (mL.min <sup>-1</sup> .kg <sup>-1</sup> )	$49.98 \pm 12.83$	$48.44 \pm 8.50$	$56.30\pm8.24$	0.09
Temperature (°C)	$30.4\pm0.3$	$30.5\pm0.2$	$30.5\pm0.3$	0.50
Humidity (%)	$30\pm7$	$30 \pm 6$	$27\pm8$	0.48
% <sub>2peak</sub>	$63.0\pm3.2$	$61.7 \pm 5.1$	$59.7\pm4.1$	0.19
Urine osmolality (mOsmol.kg <sup>-1</sup> )	$459\pm298$	$502\pm291$	$472\pm311$	0.92
Drink osmolality (mOsmol.kg <sup>-1</sup> )	$116 \pm 4$	$116 \pm 4$	$115 \pm 3$	0.70
Body mass loss (Kg)	$0.40\pm0.48$	$0.30\pm0.37$	$0.40\pm0.68$	0.79
Body mass loss (%)	$0.51\pm0.63$	$0.35\pm0.48$	$0.46\pm0.90$	0.78
Sweat loss (L)	$1.013\pm0.257$	$1.048\pm0.254$	$1.257\pm0.674$	0.28
Fluid intake (L)	$0.613 \pm 0.388$	$0.753 \pm 0.385$	$0.862 \pm 0.421$	0.31

- 556 Table 2: Participant characteristics, environmental conditions during exercise, and fluid
- balance grouped according to bradykinin receptor B2 ( $B_2R$ ) genotype. MM; homozygous wild

Variable	MM	MP	PP	P-value
n	13	17	15	-
Age (yr)	$31 \pm 9$	$26 \pm 8$	$27\pm8$	0.36
Height (cm)	$177.9\pm5.7$	$180.0\pm7.5$	$177.4\pm6.6$	0.42
Body Mass (kg)	$77.87 \pm 15.58$	$78.89 \pm 15.78$	$79.89 \pm 12.13$	0.94
BMI (kg.m <sup>-2</sup> )	$24.78\pm4.43$	$24.21\pm3.66$	$25.30\pm2.79$	0.70
$_{2peak}$ (mL.min <sup>-1</sup> .kg <sup>-1</sup> )	$49.45\pm10.12$	$50.64 \pm 11.68$	$52.59\pm8.49$	0.71
Temperature (°C)	$30.5\pm0.3$	$30.6\pm0.2^{\ast}$	$30.3\pm0.2$	< 0.05
Humidity (%)	$27 \pm 6$	$27 \pm 6$	$33\pm6^{\dagger}$	< 0.05
⁰∕₀ <sub>2peak</sub>	$61.3\pm4.6$	$62.1 \pm 5.2$	$61.0\pm3.8$	0.78
Urine osmolality (mOsmol.kg <sup>-1</sup> )	$477\pm288$	$436\pm270$	$537\pm329$	0.63
Drink osmolality (mOsmol.kg <sup>-1</sup> )	$117 \pm 5$	$116 \pm 4$	$116 \pm 3$	0.57
Body mass loss (Kg)	$0.39\pm0.36$	$0.34\pm0.50$	$0.33\pm0.58$	0.95
Body mass loss (%)	$0.50\pm0.49$	$0.41\pm0.64$	$0.37\pm0.77$	0.87
Sweat loss (L)	$0.986\pm0.254$	$1.089\pm0.401$	$1.195\pm0.521$	0.41
Fluid intake (L)	$0.599 \pm 0.322$	$0.745\pm0.374$	$0.870\pm0.459$	0.20

<sup>560</sup> \*Significantly higher than *PP*; <sup>†</sup>Significantly higher than *MM* and *MP* 

<sup>558</sup> type, *MP*; heterozygous, *PP*; homozygous insertion. Values are mean  $\pm$  SD.

#### **Figure captions**

Fig. 1: (a) Serum osmolality pre, mid and immediately post 60 min of cycling for each angiotensin converting enzyme (ACE) genotype group. WW, homozygous wild type; WI, heterozygous; II, homozygous insertion. (b) Serum osmolality pre, mid and immediately post 60 min of cycling for each *bradykinin receptor B2* ( $B_2R$ ) genotype group. *MM*, homozygous wild type; MP, heterozygous; PP, homozygous insertion. Values are mean  $\pm$  SD. \*Significant increase from baseline pre-exercise for ACE WW and WI genotype groups and all B2R genotypes (P < 0.05). \*Significant increase from baseline pre-exercise for MM genotype group (P < 0.05). Fig. 2: (a) Subjective feeling of thirst at baseline, pre-exercise and 15 min intervals throughout 60 min of cycling for each *angiotensin converting enzyme (ACE)* genotype group. WW, homozygous wild type; WI, heterozygous; II, homozygous insertion. (b) Subjective feeling of thirst at baseline, pre-exercise and 15 min intervals throughout 60 min of cycling for each *bradykinin receptor B2* ( $B_2R$ ) genotype group. *MM*, homozygous wild type; *MP*, heterozygous; *PP*, homozygous insertion. Values are mean  $\pm$  SD. \*Significant difference between *PP* and *MM* and *MP* (P < 0.05).