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On stopping voluntary muscle relaxations and contractions: evidence for shared control mechanisms and muscle state specific active breaking

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3

4 **Abbreviated title:** Stopping muscle relaxations and contractions

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37

38 **Abstract**

39 Control of the body requires inhibiting complex actions, involving contracting
40 and relaxing muscles. However, little is known of how voluntary commands to
41 relax a muscle are cancelled. Action inhibition causes both suppression of
42 muscle activity and the transient excitation of antagonist muscles, the latter
43 being termed active breaking. We hypothesised that active breaking is present
44 when stopping muscle relaxations. Stop signal experiments were used to
45 compare the mechanisms of active breaking for muscle relaxations and
46 contractions in male and female human participants. In experiments 1 and 2, go
47 signals were presented which required participants to contract or relax their
48 biceps or triceps muscle. Infrequent stop signals occurred after fixed delays (0-
49 500ms), requiring that participants cancelled go commands. In experiment 3,
50 participants increased (contract) or decreased (relax) an existing isometric
51 finger abduction depending on the go signal, and cancelled these force changes
52 whenever stop signals occurred (dynamically adjusted delay). We found that
53 muscle relaxations were stopped rapidly, met predictions of existing race
54 models, and had stop signal reaction times that correlated with those observed
55 during the stopping of muscle contractions, suggesting shared control
56 mechanisms. However, stopped relaxations were preceded by transient
57 increases in EMG, while stopped contractions were preceded by decreases in
58 EMG, suggesting a later divergence of control. Muscle state specific active
59 breaking occurred simultaneously across muscles, consistent with a central
60 origin. Our results indicate that the later stages of action inhibition involve

61 separate excitatory and inhibitory pathways, which act automatically to cancel
62 complex body movements.

63 **Significance Statement**

64 The mechanisms of how muscle relaxations are cancelled are poorly
65 understood. We showed in three experiments involving multiple effectors that
66 stopping muscle relaxations involves transient bursts of EMG activity, which
67 resemble co-contraction and have onsets that correlate with stop signal reaction
68 time (SSRT). Comparison with the stopping of matched muscle contractions
69 showed that active breaking was muscle state specific, being positive for
70 relaxations and negative for contractions. The two processes were also
71 observed to co-occur in agonist-antagonist pairs, suggesting separate
72 pathways. The rapid, automatic activation of both pathways may explain how
73 complex actions can be stopped at any stage of their execution.

74

75 **Introduction**

76 Control of the body necessitates the rapid cancelling of actions in
77 response to changes in the environment. Such actions often involve multiple
78 muscles in different states of contraction and relaxation. However, most action
79 inhibition experiments have focused on simple button press responses and
80 saccades (Band and van Boxtel, 1999; Verbruggen and Logan, 2008). Less is
81 known about complex actions, particularly those caused by muscle relaxations.

82 Voluntary isometric and isotonic muscle relaxations are an active
83 process (Toma et al., 2000; Alegre et al., 2003), vital for updating posture and
84 initiating movement (Buccolieri et al., 2003). Relaxation commands activate the

85 supplementary and primary motor cortex (M1) to a similar degree to matched
86 muscle contractions (Toma et al., 1999; Spraker et al., 2009). Prior to a
87 relaxation command, readiness potentials are observed at fronto-central
88 electrodes (Dimitrov, 1985; Terada et al., 1995; Rothwell et al., 1998), while the
89 execution of the movement itself is preceded by a lateralized readiness
90 potential (LRP) of equal amplitude and latency to those preceding muscle
91 contractions (Pope et al., 2007). Understanding the control of voluntary muscle
92 relaxations is important because muscle relaxation deficits are prevalent in
93 stroke (Chae et al., 2002, 2006; Seo et al., 2009), dystonia (Yazawa et al.,
94 1999; Oga et al., 2002) and Parkinson's disease (Kunesch et al., 1995; Grasso
95 et al., 1996; Labyt et al., 2005).

96 Given the control similarities to muscle contraction, it should be possible
97 to cancel a 'go command' to relax a muscle if a 'stop command' is issued early
98 enough. The stop signal task involves participants responding to a go signal on
99 every trial and attempting to suppress their responses whenever an infrequent
100 stop signal occurs (Lappin and Eriksen, 1966; Logan et al., 1984). Stop signal
101 reaction time (SSRT) can then be calculated, which is a good predictor of
102 general inhibitory function (Logan et al., 1997; Williams et al., 1999). Stop and
103 go processes race to completion via largely separate pathways in the brain
104 (Logan and Cowan, 1984; Aron et al., 2007). If stopping muscle relaxations
105 conforms to this account, it may be possible to derive the general mechanisms
106 determining how complex actions are cancelled.

107 Action inhibition involves multiple central and peripheral pathways. In the
108 motor cortex, stopping is associated with increased short-interval intracortical

109 inhibition (SICI; Coxon et al., 2006; van den Wildenberg et al., 2010), indicating
110 greater inhibitory GABAergic activity. Increased SICI is also observed during
111 muscle relaxation (Buccolieri et al., 2004; Motawar et al., 2012). As such,
112 combining muscle relaxation go commands and action inhibition might produce
113 slowing of SSRT due to interference. However, excitatory pathways likely also
114 contribute to stopping, negating any such slowing. Excitation has been
115 observed at the periphery in the form of bursts of antagonist activity when
116 reaching movements are inhibited (Atsma et al., 2018). This 'active breaking'
117 may also explain how ballistic movements (McGarry and Franks, 1997) and
118 increases of isometric force (de Jong et al., 1990) can be arrested mid-way
119 through their execution. Nevertheless, it is unknown whether active breaking is
120 confined to the antagonist muscle, nor whether it occurs uniformly in relaxing
121 and contracting muscles, or manifests differently in both as a form of muscle
122 state specific response. Relatedly, it is uncertain whether active breaking arises
123 from changes at the agonist which in turn influence the antagonist (peripheral
124 command), or from a single central command, which may be similar to co-
125 contraction.

126 We therefore designed a series of stop signal experiments to determine
127 the general peripheral mechanisms underpinning the inhibition of relaxing and
128 contracting muscles. We test the hypotheses that: 1) voluntary relaxations are
129 cancelled as rapidly as muscle contractions, in accordance with the principles of
130 the race model, 2) Active breaking is present across movement types, but
131 manifests differently depending on the muscle state, 3) Active breaking involves
132 simultaneous action across muscles.

133

134 **Methods**135 *Equipment*

136 Participants were seated at an adjustable table, fitted with foam padding.

137 Modified stop signal tasks were run via MATLAB (2017a) and Psychtoolbox

138 (Brainard, 1997). Visual stimuli were displayed via a flat screen monitor (27-inch

139 LCD, 1902 x 1080 pixels, 60 Hz refresh rate). Auditory stimuli consisted of

140 tones delivered via a buzzer (3000 Hz, 50 dB, 160ms).

141 Electromyography (EMG) was recorded from bipolar, surface electrodes

142 (Ag-AgCl disposable electrode, GE Healthcare Japan, Tokyo, Japan) placed

143 over the middle of the right biceps brachii and right triceps brachii lateral head

144 muscles in experiment 1 and 2, and on the first dorsal interosseous (FDI)

145 muscle in experiment 3. EMG signals were sampled at 2000 Hz and amplified

146 using variable gain (MME-3132, Nihon Kohden, Tokyo, Japan). For the

147 purposes of training participants to relax their muscles, EMG data was

148 displayed via an oscilloscope (TDS2004C, Tektronix, Inc., Oregon, USA)

149 positioned at eye level in front of them. This was removed during the actual task.

150 In experiments 1 and 2, elbow angle was determined via the use of three

151 infrared reflective markers positioned at the forearm, elbow and shoulder of the

152 right arm. Marker position was detected via two motion tracking cameras

153 (Oqus300, Qualisys AB, Gothenburg, Sweden) with a sampling rate of 250 Hz.

154 In experiment 3, isometric finger abduction force was measured using a force

155 sensor (6-axis force sensor; ThinNANO 5/4-A, BL-Autotech, LTD., Kobe, Japan).

156 Maximum voluntary forces were recorded at the start of the experiment via a

157 different force sensor (Force gauge; FGC-5, Nidec-Shimpo, Kyoto, Japan).

158

159 *Participants*

160 For experiment 1 we recruited 13 participants (8 were female, mean age
161 = 34.5 yrs, SD = 9.2 yrs). Experiment 2 included 12 participants (8 were female,
162 mean age = 33.83 yrs, SD = 9.42 yrs), 5 of whom had previously participated in
163 experiment 1. We recruited 16 participants for experiment 3. However, 2
164 participants were excluded before the analysis phase due to equipment failure.
165 This left 14 participants who were included in the analysis (11 were female,
166 mean age = 36.71 yrs, SD = 8.63 yrs). Of these, 5 had participated in exps. 1
167 and 3 had participated in exps. 2. Experiments were undertaken with the
168 understanding and written consent of each participant in accordance with the
169 Code of Ethics of the World Medical Association (Declaration of Helsinki), and
170 with the NTT Communication Science Laboratories Research Ethics Committee
171 approval.

172

173 *General procedure*

174 Participants first completed a practice button press version of the stop signal
175 task, similar to those widely used in the stop signal literature (Verbruggen et al.,
176 2019). Maximum voluntary contractions (MVC) of relevant muscles were then
177 recorded via 5 s isometric contractions against a fixed surface. All experiments
178 used a modified version of the stop signal task. Participants maintained a
179 constant level of muscle contraction in the target effector whilst looking at a
180 fixation cross on a screen 70 cm in front of them. On every trial, a visual go
181 signal was presented in the centre of the screen in the form of a coloured shape
182 (subtending 4.7 ° visual angle). In the relaxation condition participants relaxed

183 the target muscle and in the contraction condition they contracted the target
184 muscle as quickly as possible in response to the go signal. On a randomly
185 selected subset of trials (30%), a stop signal (tone) was presented after a
186 specifically controlled stop signal delay (SSD). Experiments 1 and 2 compared
187 the stopping of relaxation of the biceps to the stopping of contractions of the
188 triceps and biceps muscle respectively. Thus in experiment 1, the movement
189 direction was matched (elbow extension) and in experiment 2 the primary
190 muscle was matched (biceps brachii) across movement conditions. Experiment
191 3 was conducted using isometric relaxations and contractions of the FDI muscle
192 to determine whether the findings of the previous experiments generalised to
193 other effectors.

194

195 *Procedure for experiment 1*

196 The target muscle was the right biceps brachii in the relax condition and the
197 triceps brachii in the contract condition. Electromyography (EMG) was recorded
198 from both muscles in all conditions. Uniquely in experiment 1, we tested the
199 relax and contract condition in separate blocks (counterbalanced). Participants
200 were informed of the block type on screen at the start of each block. They
201 rested their elbow on a padded table, supinated and bent at an angle of $\sim 30^\circ$
202 from the horizontal (Fig. 1. A.), which required activity in the biceps muscle
203 ($\sim 4\%$ MVC). Before each trial, arm angle was adjusted via visual feedback on
204 screen. Successfully matching current to required elbow angle triggered the
205 appearance of a fixation cross.

206 After a random inter-stimulus interval (ISI = 2-4s) the fixation cross was
207 replaced by green shape subtending 4.7° of visual angle (Go/No Go stimulus).

208 Participants moved their right arm downwards as rapidly as possible whenever
209 a green circle appeared on the screen (Go trials; 80% of trials), and remained
210 stationary whenever a green square appeared (No-go trials; 20% of trials). No
211 go trials ensured participants processed the go stimulus before acting
212 (Verbruggen and Logan, 2009), but were not analysed further. During
213 'relaxation blocks' participants moved their right arm downwards (elbow
214 extension) by relaxing the biceps muscle, but during 'contraction blocks'
215 extension was achieved by contracting the triceps muscle. On a random subset
216 (30%) of trials a stop signal (tone) was presented. Stop signal onset time in
217 relation to the go signal varied between 0 and 500ms (stop signal delay). There
218 were six tone intervals (0, 100, 200, 300, 400, 500ms; randomized).

219 Each block consisted of 80 trials (40 go trials, 24 stop trials, 8 no go trials
220 with tone, 8 no go trials without tone). There were 10 blocks in total (5 relax, 5
221 contract). The experiment lasted ~3 hours.

222

223 *Procedure for experiment 2*

224 Procedural details were the same as experiment 1, with three important
225 differences. Firstly, the target muscle in the contract condition was the biceps
226 and the go response in this condition involved flexing the elbow (Fig. 3. A.). The
227 relax condition was the same as experiment 1. Secondly, relax and contract
228 conditions were tested within the same blocks. Thirdly, we did not include No go
229 trials. Participants were instructed to relax their right biceps muscle in response
230 to a blue circle appearing on the screen (50% of trials; randomized) and
231 contract their right biceps muscle in response to a blue square (50% of trials;
232 randomized). Stimulus-action pairing was counterbalanced across participants.

233 Stop signals were the same as experiment 1. There were 10 experimental
234 blocks of 80 trials (28 relax condition go trials, 12 relax condition stop trials, 28
235 contract condition go trials, 12 contract condition stop trials).

236

237

238 *Procedure for experiment 3*

239 Experiment 3 was different to experiment 2 because the target muscle
240 was the right first dorsal interosseous (FDI), the go response involved isometric
241 increases and decreases of force, and the stop signal used dynamic rather than
242 fixed SSDs. Participants rested their arm on a table in a pronated position (Fig.
243 5. A.) with the outside of their right index finger touching a force sensor. They
244 pushed outwards (abduction) with ~10% MVC to move a small black circle
245 between two boundary lines on screen. These lines, and an additional target
246 line representing a force of 20% MVC, were shown during training, but removed
247 during the actual task once a stable force was achieved at the start of each trial.
248 Go signals were presented after variable ISI (1-4s). Participants had to rapidly
249 decrease FDI muscle activity to 0 if a *circle* appeared (relax condition) and
250 rapidly increase it by ~10% if a *square* appeared (contract condition).

251 SSD on stop trials (30% of trials) was adjusted from an initial 250ms by
252 increasing it by 50ms every time participants successfully stopped and
253 decreasing by 50ms every time they failed to stop. Staircasing was performed
254 separately for each condition (relax vs contract). This *one-up one-down* method
255 is commonly used in stop signal experiments and requires online trial
256 classification (Logan et al., 1997; Osman et al., 1986; Verbruggen and Logan,
257 2009). We classified stop trials by setting a force change threshold based on

258 the mean maximum force 'velocity' (first derivative of force increase/decrease)
259 during the go trials of the previous run. Exceeding 30% of this value during the
260 response window meant a stop trial was classified as a failed stop trial. Trials
261 under threshold were classified as successful stop trials. There were 10 runs of
262 60 trials (21 relax go trials, 9 relax stop trials, 21 contract go trials, 9 contract
263 stop trials per run). The experiment lasted ~ 3 hours.

264

265 *Analysis*

266 Elbow angle velocity was determined in exp. 1 and 2 by calculating the
267 angle change (relative to baseline: -2000ms to 0ms) between three reflective
268 marker located on the upper arm, elbow and forearm. The angular change data
269 was smoothed (zero-phase digital filtering; low pass 10 Hz) and the velocity was
270 calculated by taking the difference between successive time points.
271 Acceleration was calculated by taking the difference between successive time
272 points of the velocity data. Force velocity was calculated in exp. 3, by taking the
273 difference between successive time points of the smoothed force data (zero-
274 phase digital filtering; low pass 5 Hz).

275 Response time (RT) was the point after the Go signal where the angular
276 velocity first rose above 10% of the maximum velocity on that trial (Irlbacher et
277 al., 2006; De Havas et al., 2016). Outliers ($> 4 \times SD$) were re-assigned to the
278 value of the slowest non-outlier trial to ensure SSRT calculations were unbiased
279 (Verbruggen and Logan, 2009). In exp. 3 we used a threshold of 30% because
280 the force signal contained more noise.

281 EMG data in all experiments was band-pass filtered (zero-phase digital
282 filtering; 10–500 Hz) and rectified, before being smoothed (zero-phase digital

283 filtering; low pass 10 Hz). EMG velocity was calculated by taking the difference
284 between successive time points in the smoothed data. We determined EMG-RT
285 on Go trials via the threshold technique (first time point > 30% of the max, which
286 remained above threshold for at least 25ms).

287 Stop trial success was classified online in exp. 3 (*see procedure*), but in
288 exp. 1 and 2 it was classified offline by first calculating the mean peak go trial
289 velocity separately for each movement type for each participant. Stop trials
290 were coded as successful if the velocity never rose above 10% of the mean
291 max go trial velocity, but as failed if the threshold was exceeded. In all
292 experiments, failed stop trial RT was the point relative to the go signal where
293 the velocity first rose above the stop classifier threshold. Outliers were dealt
294 with in the same manner as for go trials. All calculations were performed
295 independently for the relax and contract conditions. Response times were
296 compared across conditions via within subjects t-tests.

297 In exp. 3 we calculated the mean probability of movement (pMov) in each
298 movement condition (total failed stop trials divided by total stop trials). In exp. 1
299 and 2, pMov was calculated at each SSD, in each movement condition in the
300 same manner. One-way within subjects ANOVAs were used to verify that pMov
301 increased across the six levels of SSD in each movement condition. Inhibition
302 functions were fit to the pMov data using the Weibull method (Hanes and Schall,
303 1995). We used Matlab (2017a) and the Palamedes curve fitting toolkit (Prins
304 and Kingdom, 2018), with a 4 parameter model (α = scale, β = shape, γ = lower
305 bias, λ = upper bias). Lower and upper bias in this context are synonymous with
306 error rate, and in the SST literature consist of trials where stop (trigger failure)

307 and go (forced inhibition) processes are not initiated (Band et al., 2003). Both
308 bias parameters were restricted to 0.05 (Livesey and Livesey, 2016). Beta
309 values were compared via t-test to determine whether inhibition functions
310 differed in shape across relax and contract conditions.

311 SSRT was calculated in exp. 3 by subtracting mean SSD from go trial
312 response time in each movement condition (Verbruggen et al., 2019). In exp. 1
313 and 2, SSRT was calculated using the integration method, which is commonly
314 used for fixed SSDs (Logan and Cowan, 1984) and was necessary because we
315 did not tightly constrain performance, leading to heterogeneity in SSD curves
316 (Verbruggen and Logan, 2009). Go RTs were rank ordered from smallest to
317 largest. For each SSD the total number of Go RT trials was multiplied by the
318 pMov at that SSD. This value was rounded to the nearest integer and the
319 corresponding Go RT in the rank ordered list was selected. The SSD itself was
320 then subtracted from this Go RT to give the SSRT at that SSD. Overall SSRT
321 was then calculated by taking the mean of all the SSRTs obtained, separately
322 for the relax and contract conditions.

323 SSRTs were compared across movement conditions in all experiments
324 via within-subjects t-tests and Pearson's r . To discount the possibility that
325 experiment 1 was underpowered and to avoid type II errors, we applied a
326 Bayesian analysis (Dienes, 2008, 2014) to experiments 2 and 3 using the mean
327 difference between conditions obtained in experiment 1 (half-normal distribution,
328 1-tailed).

329 In all experiments, active breaking was defined as a consistent stop-
330 signal related change in the elbow angle acceleration (exp. 1 and 2), force

331 velocity (exp. 3) or EMG velocity (exp. 1, 2 and 3) that could not be ascribed to
332 the go signal. In exp. 1 and 2, kinematic active breaking was determined from
333 angle acceleration to control for the large deviations in position and velocity on
334 failed stop trials. In all cases, we first removed any stop trials where early
335 movements occurred in the opposite direction to that which would be expected
336 if it was a Go trial (mean no. trials removed per subject in exp. 1 = 0.6%, exp. 2.
337 = 1.6%, exp. 3. = 2.18%). Next, through visual inspection of group average
338 kinematic, force and EMG data we determined the direction (positive or
339 negative) of active breaking, separately for successful and failed stop trials in
340 each movement condition. Active breaking onset times were calculated
341 separately for elbow angle acceleration, force velocity and EMG velocity. For a
342 given signal, onset time was defined at the individual participant level by taking
343 the first time point 0 – 500ms after the stop signal that reached 30% of its max
344 or minimum value (depending on active breaking direction) within the same
345 window. To determine if active breaking was significant at the group level, we
346 averaged the signal in a 50ms window time-locked to active breaking onset for
347 each participant and compared these values to 0 via one-sample t-tests (see
348 Fig. 2., 4. & 6.). After confirming the presence of active breaking at the group
349 level, the analysis was repeated across trials at the individual participant level
350 and tested via one-sample t-test (1-tailed).

351 In experiment 1, in the relax condition the primary muscle was the biceps
352 and the secondary muscle was the triceps. Some participants (n = 5) showed
353 evidence of increased triceps activity before the stop signal on some trials and
354 were excluded from the secondary muscle analyses to minimize false positive

355 detection of active breaking. In the contract condition the primary muscle was
356 the triceps and the secondary muscle was the biceps. Again, we had to remove
357 some participants (n = 5) who showed early pre-stop signal activation of the
358 secondary muscle. In experiment 2, as before the biceps was the primary
359 muscle and the triceps was the secondary in the relax condition (n = 5 excluded
360 from secondary muscle analysis, due to pre-stop signal activity on some trials).
361 Likewise, in the contract condition the biceps was the primary muscle and the
362 triceps was the secondary muscle. One participant was excluded from the
363 secondary muscle analysis because triceps EMG active breaking occurred very
364 late relative to the stop signal (>300ms), and after the kinematic active breaking
365 onset, meaning it was a reaction to the arm stopping rather than a cause.
366 Finally, in experiment 3, the FDI was the primary muscle in both movement
367 conditions.

368 We performed a further analysis on successful stop trials to determine: 1)
369 if active breaking occurred when kinematics/force traces were flat, and 2) how
370 active breaking magnitude changed with decreasing detected go response. In
371 all experiments, successful stop trials were split into quartiles at the individual
372 participant level after rank ordering them according to the amplitude of go
373 response detected in the smoothed kinematic or force traces prior to the stop
374 signal onset (-1000 – 0ms). Primary muscle active breaking amplitude was then
375 calculated for trials in each of the 4 bins. This was done at the group level in the
376 same manner as described above, using the already derived active breaking
377 onset times (based on all successful stop trials) for time-locking.

378 Mean individual active breaking onset times were calculated by taking
379 the average between successful and failed stop trials, separately for kinematics
380 force and primary muscle EMG in each movement condition. Mean active
381 breaking times were then correlated to SSRT in all experiments via Pearson's r ,
382 under a 1-tailed assumption that all correlations were expected to be positive.

383 Primary and secondary muscle active breaking onset times were
384 compared via paired sample t-tests in exp. 1 (relax $n = 8$, contract $n = 8$) and
385 exp. 2 (relax $n = 7$, contract $n = 11$). A time-shifted cross correlation analysis
386 was done to determine if active breaking in one muscle predicted active
387 breaking in the other after some reliable time-delay. Absolute triceps muscle
388 EMG velocity traces were shifted (-50 – 50ms) in 0.5ms increments relative to
389 the Biceps muscle active breaking onset time. After each shift, cross-correlation
390 was performed (Pearson's r) during a window that included the whole active
391 breaking response (EMG velocity traces in both muscles, -10 – 250ms relative
392 to biceps muscle active breaking onset). The reverse calculation was also
393 performed (i.e. biceps shifted relative to triceps) and the results averaged
394 together, after the time-axis had been reversed to ensure results were in the
395 same direction. This was done to control for any biases in how onset times were
396 detected across muscles. Thus in both cases, if correlations peaked reliably
397 before 0ms it indicated that the biceps activity predicted the triceps activity, and
398 if correlations peaked reliably after 0ms it indicated triceps activity predicted the
399 biceps activity. Positive peaks close to 0ms indicated simultaneous action. The
400 analysis was performed at the individual participant level. Successful and failed

401 stop trial results were then averaged, and group averages were calculated for
402 relax and contract conditions in each experiment (see Fig. 2. M. & Fig. 4. M.).

403

404 -----

405 Figure 1

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408 -----

409 Table 1

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413 **Results**414 **Go trial response times**

415 Go trial response times were faster in the contract condition than the
416 relax condition in experiment 1 (521.28ms vs 554.74ms; $t(12) = 2.511$, $p =$
417 0.027 , Cohen's $d = 0.57$; Table 1) and experiment 2 (549.82ms vs 588.36; $t(11)$
418 $= 3.879$, $p = 0.003$, Cohen's $d = 0.77$; Table 3), but did not differ in experiment 3
419 (595.11ms vs 598.01ms; $t(13) = 0.26$, $p = 0.799$, Cohen's $d = 0.05$; Table 5). In
420 experiment 1, for the relax condition the biceps EMG-RT was faster than the
421 triceps (311.01ms vs 390.61ms; $t(12) = -3.572$, $p = 0.004$, Cohen's $d = -1.17$).
422 In the contract condition the biceps muscle began to relax earlier than the
423 triceps muscle started to contract (331.05ms vs 417.18ms; $t(12) = -6.036$, $p <$
424 0.001 , Cohen's $d = -1.56$). In experiment 2, the biceps muscle EMG-RT was
425 again faster than the triceps muscle in the relax condition (331.76ms vs
426 416.21ms; $t(11) = -4.083$, $p = 0.002$, Cohen's $d = -1.39$). Biceps and triceps
427 muscle EMG-RT did not differ in the contract condition, when both were
428 contracting (437.79ms vs 440.08ms; $t(11) = -0.852$, $p = 0.412$, Cohen's $d = -$

429 0.07). However, comparing directly between the relax and contract conditions in
430 the biceps showed relaxation was indeed faster ($t(11) = -11.003$, $p < 0.001$,
431 Cohen's $d = -2.99$). Thus relaxation was faster than contraction in the forearm,
432 particularly in the biceps muscle. However, in the FDI muscle (exp. 3) there was
433 only a trend for relaxation being faster than contraction (424.29ms vs 445.21ms;
434 $t(13) = -1.982$, $p = 0.069$, Cohen's $d = -0.41$).

435

436

437

438

439 **Stopping muscle relaxations not impaired relative to stopping**
440 **contractions**

441 In experiment 1 contract SSRT (Mn = 238.51ms, SD = 49.42ms) and
442 relax SSRT (Mn = 220.46, SD = 49.42) did not significantly differ ($t(12) = 1.628$,
443 $p = 0.129$, Cohen's $d = 0.401$, observed power = 0.266). There was also no
444 significant difference in SSRT between contract and relax conditions in
445 experiment 2 (Mn = 256.5ms, SD = 31.69 vs. Mn = 268.1ms, SD= 48.31; $t(11)$
446 = -0.992, $p = 0.34$, Cohen's $d = -0.284$, observed power = 0.147) and
447 experiment 3 (Mn = 156.86ms, SD = 69.84 vs. Mn = 170.98ms, SD= 58.94; t
448 (13) = -1.31, $p = 0.213$, Cohen's $d = -0.219$, observed power = 0.118).

449 Bayesian analysis of experiments 2 and 3, using the mean difference
450 between conditions in experiment 1 (half-normal distribution, 1-tailed), was
451 consistent with weak support for the null hypothesis in experiment 2 (mean diff.
452 = -11.599, SE = 13.314; likelihood of theory = 0.0076, likelihood of null = 0.021,
453 Bayes factor = 0.369) and support for the null in experiment 3 (mean diff. = -

454 14.124, SE = 11.886; likelihood of theory = 0.0048, likelihood of null = 0.0166,
 455 Bayes factor = 0.291). This suggests that SSRT does not differ across muscle
 456 contractions and relaxations.

457 -----

458 Figure 2

459 -----

460

461 -----

462 Table 2

463 -----

464

465

466

467 **The race model can be applied to stopping muscle relaxations**

468 Experiment 1 and 2 had a fixed SSD design. In line with the race model,
 469 the probability of failing to inhibit a response increased with increasing SSD
 470 across movement conditions (Fig. 1. E. and Fig. 2. C.). The effect was strong
 471 for the relax ($F(5,60) = 100.937$, $p < 0.001$, $\eta^2 = 0.89$) and contraction
 472 conditions ($F(5,60) = 79.418$, $p < 0.001$, $\eta^2 = 0.87$) in experiment 1, and for the
 473 relax ($F(5,55) = 112.86$, $p < 0.001$, $\eta^2 = 0.91$) and contract conditions ($F(5,55)$
 474 $= 97.403$, $p < 0.001$, $\eta^2 = 0.9$) in experiment 2.

475 Weibull functions showed a high degree of fit to the probability of
 476 stopping data (Fig. 1. E. and Fig. 2. C.). In the first experiment, mean adjusted
 477 R^2 was 0.90 (SD = 0.13) in the relax condition and 0.81 (SD = 0.18) in the
 478 contract condition. In experiment 2, R^2 was 0.89 (SD = 0.15) in the relax
 479 condition and 0.84 (SD = 0.24) in the contract condition. The shape of these SD
 480 curves was similar across movement conditions. Mean Weibull β parameter

481 value were 4.23 (SD = 1.95) for the relax condition and 3.34 (SD = 1.48) for the
 482 contract conditions in experiment 1 ($t(12) = 1.73$, $p = 0.11$, Cohen's $d = 0.51$),
 483 and in experiment 2 were 4.35 (SD = 1.43) for the relax condition and 3.97 (SD
 484 = 1.6) for the contract condition ($t(11) = 0.88$, $p = 0.4$, Cohen's $d = 0.25$).

485 Experiment 3 used a variable SSD design. Despite independent
 486 staircasing, the probability of moving (i.e. failed stop trials/total stop trials) did
 487 not differ between relax ($M = 0.48$, $SD = 0.04$) and contract conditions ($M = 0.47$,
 488 $SD = 0.05$; $t(13) = 1.325$, $p = 0.21$, Cohens $d = 0.17$). Mean SSD was also
 489 similar for muscle relaxations ($M = 427.02\text{ms}$, $SD = 55.48\text{ms}$) and contractions
 490 ($M = 438.25\text{ms}$, $SD = 59.97\text{ms}$; $t(13) = -0.77$, $p = 0.45$, Cohens $d = -0.19$).

491 As predicted by the race model, mean response times were significantly
 492 faster on failed stop trials than on go trials across all three experiments. This
 493 held for the relax ($t(12) = 5.83$, $p < 0.001$, Cohens $d = 1.22$) and contract
 494 conditions ($t(12) = 6.69$, $p < 0.001$, Cohens $d = 1.15$) in experiment 1, for the
 495 relax ($t(11) = 6.62$, $p < 0.001$, Cohens $d = 0.9$) and contract conditions ($t(11) =$
 496 5.22 , $p < 0.001$, Cohens $d = 1.21$). in experiment 2, and for the relax ($t(13) =$
 497 10.58 , $p < 0.001$, Cohens $d = 1.58$) and contract conditions ($t(13) = 8.4$, $p <$
 498 0.001 , Cohens $d = 1.79$) in experiment 3.

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500 -----
 501 Figure 3

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505 Table 3

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508 Relax SSRT and contract SSRT positively correlated

509 We observed a positive correlation between relax SSRT and contract
510 SSRT in all three experiments. In experiment 1 ($r = 0.62$, $p = 0.024$; Fig. 1. F.)
511 and experiment 3 ($r = 0.82$, $p < 0.001$; Fig. 5. C.) the results were significant,
512 while in experiment 2 ($r = 0.55$, $p = 0.061$; Fig. 3. D.) they only showed a trend
513 towards significance.

514

515 Stopping involves active breaking

516 In all experiments significant active breaking was observed at the group
517 level for all conditions. Experiment 1 compared the stopping of elbow
518 extensions caused either by relaxation of the biceps (relax condition) or
519 contraction of the triceps (contract condition; Fig. 1. G-I.). Elbow acceleration
520 showed positive active breaking (flexion direction) when relaxations were
521 successfully ($t(12) = 6.903$, $p < 0.001$, Cohen's $d = 1.91$; Fig. 2. A.) and
522 unsuccessfully ($t(12) = 6.012$, $p < 0.001$, Cohen's $d = 1.67$; Fig. 2. B.) stopped.
523 Biceps EMG velocity showed positive active breaking earlier in these same
524 successful ($t(12) = 6.862$, $p < 0.001$, Cohen's $d = 1.9$; Fig. 2. C.) and failed stop
525 trials ($t(12) = 5.125$, $p < 0.001$, Cohen's $d = 1.42$; Fig. 2. D.). We also tested
526 active breaking at the individual subject level (1-tailed). All participants (13/13)
527 showed significant active breaking for kinematics and EMG on successful and
528 failed stop trials in the relax condition.

529 In the contract condition, active breaking (flexion direction) was observed
530 when examining elbow acceleration on successful ($t(12) = 6.168$, $p < 0.001$,
531 Cohen's $d = 1.71$; Fig. 2. A.) and failed ($t(12) = 7.188$, $p < 0.001$, Cohen's $d =$

532 1.99; Fig. 2. B.) stop trials. On failed stop trials, when the triceps was already
533 contracting, negative active breaking was observed ($t(12) = -5.812$, $p < 0.001$,
534 Cohen's $d = -1.61$; Fig. 2. F.). Conversely, on successful stop trials, when the
535 biceps muscle was relaxing, but the triceps had typically not begun to contract
536 (see table 1.), positive active breaking was observed ($t(12) = 3.167$, $p = 0.008$,
537 Cohen's $d = 0.88$; Fig. 2. E.). Positive triceps EMG velocity increases on
538 successful stop trials could therefore have been contaminated with 'go
539 response', since both involved signal increase. This was judged unlikely given
540 the onset timing and shape of the triceps response, which was more closely
541 related to the stop-signal across participants. After positive active breaking, a
542 negative dip was observed (Fig. 2. E., red line), indicative of a bi-phasic
543 response, but also of later voluntary response components. As such, we
544 analysed only the first stop-related components of the signal in each condition.
545 Active breaking as defined above could be detected at the individual participant
546 level in 10/13 participants from the kinematics and 8/13 from the EMG on
547 successful stop trials. For failed stop trials it reached threshold ($p < 0.1$) in 12/13
548 participants for kinematics and 13/13 for triceps EMG.

549 In experiment 2 we compared the stopping of biceps relaxations (elbow
550 extension) and contractions (elbow flexion; Fig. 3. E-G.). In the relax condition
551 the results replicated those seen in experiment 1. Positive active breaking was
552 observed from the kinematics for successful ($t(11) = 6.04$, $p < 0.001$, Cohen's d
553 $= 1.74$; Fig. 4. A.) and failed stops ($t(11) = 5.447$, $p < 0.001$, Cohen's $d = 1.57$;
554 Fig. 4. B.) and from the biceps EMG velocity for successful ($t(11) = 6.56$, $p <$
555 0.001 , Cohen's $d = 1.89$; Fig. 4. C.) and failed stops ($t(11) = 4.994$, $p < 0.001$,

556 Cohen's $d = 1.44$; Fig. 4. D.). In the contraction condition, negative active
557 breaking was observed from the kinematics for successful ($t(11) = -5.614$, $p <$
558 0.001 , Cohen's $d = -1.62$; Fig. 4. A.) and failed stops ($t(11) = -8.261$, $p <$
559 0.001 , Cohen's $d = -2.38$; Fig. 4. B.) and from the biceps EMG velocity for successful
560 ($t(11) = -6.037$, $p <$ 0.001 , Cohen's $d = -1.74$; Fig. 4. C.) and failed stops ($t(11) =$
561 -6.51 , $p <$ 0.001 , Cohen's $d = -1.88$; Fig. 4. D.). At the individual participant level
562 active breaking could again be detected in the majority of cases for the relax
563 condition (kinematics: succ. = 10/12, fail = 12/12, EMG: succ. = 11/12, fail =
564 12/12) and for the contract condition (kinematics: succ. = 9/12, fail = 12/12,
565 EMG: succ. = 10/12, fail = 12/12).

566 Experiment 3 had participants stopping isometric relaxations and
567 contractions of the FDI muscle (Fig. 5. D-E). Positive active breaking was found
568 in the relax condition when examining force velocity on successful ($t(13) = 6.972$,
569 $p <$ 0.001 , Cohen's $d = 1.86$; Fig. 6. A.) and failed stop trials ($t(13) = 6.742$, $p <$
570 0.001 , Cohen's $d = 1.8$; Fig. 6. B.), and when examining FDI EMG velocity for
571 successful ($t(13) = 8.139$, $p <$ 0.001 , Cohen's $d = 2.18$; Fig. 6. C.) and failed
572 stop trials ($t(13) = 4.936$, $p <$ 0.001 , Cohen's $d = 1.32$; Fig. 6. D.). In the
573 contract condition, we found negative active breaking for force velocity on
574 successful ($t(13) = -6.923$, $p <$ 0.001 , Cohen's $d = -1.85$; Fig. 6. A.) and failed
575 stop trials ($t(13) = -6.524$, $p <$ 0.001 , Cohen's $d = -1.74$; Fig. 6. B.), and for FDI
576 EMG velocity on successful ($t(13) = -7.656$, $p <$ 0.001 , Cohen's $d = -2.05$; Fig. 6.
577 C.) and failed stop trials ($t(13) = -4.306$, $p <$ 0.001 , Cohen's $d = -1.15$; Fig. 6. D.).
578 Active breaking was also detected in the majority of cases at the individual
579 participant level, both for the relax (Force: succ. = 13/14, fail = 13/14, EMG:

580 succ. = 13/14, fail = 14/14) and contract conditions (Force: succ. = 13/14, fail =
581 14/14, EMG: succ. = 14/14, fail = 14/14).

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584 Figure 4

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588 Table 4

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591 **Active breaking is muscle state specific**

592 In experiment 1 and 2 we observed active breaking of secondary
593 muscles. In both cases for the relax condition the triceps was considered the
594 secondary muscle, showing significant positive active breaking for successful
595 stop trials ($t(7) = 4.587$, $p = 0.003$, Cohen's $d = 1.62$; Fig. 2. E.) and failed stop
596 trials ($t(7) = 4.96$, $p = 0.002$, Cohen's $d = 1.75$; Fig. 2. F.) in experiment 1 and
597 significant positive active breaking for successful stop trials ($t(6) = 3.901$, $p =$
598 0.008 , Cohen's $d = 1.47$; Fig. 4. E.) and failed stop trials ($t(6) = 2.547$, $p = 0.045$,
599 Cohen's $d = 0.96$; Fig. 4. F.) in experiment 2. In experiment 1, the biceps
600 muscle was the secondary muscle in the contract condition and significant
601 positive active breaking was found on successful ($t(7) = 4.01$, $p = 0.005$,
602 Cohen's $d = 1.42$; Fig. 2. C.) and failed stop trials ($t(7) = 5.125$, $p < 0.001$,
603 Cohen's $d = 1.91$; Fig. 2. D.). Here the active breaking was positive because the
604 biceps muscle was relaxing in response to the go signal. Conversely, in the
605 contract condition in experiment 2, the triceps (secondary muscle) tended to
606 contract in response to the go signal, presumably to stabilise the elbow during

607 rapid flexion. As such, active breaking was significantly negative at the group
608 level on successful ($t(10) = -4.482$, $p = 0.001$, Cohen's $d = -1.35$; Fig. 4. E.) and
609 failed stop trials ($t(10) = -2.826$, $p = 0.018$, Cohen's $d = -0.85$; Fig. 4. F.).

610 In experiment 1, 8/8 participants showed significant (1-tailed) triceps
611 active breaking in the relax condition on successful stop trials and 7/8 showed
612 significant active breaking on failed stop trials, at the individual participant level
613 (1-tailed). For the contract condition, 7/8 showed significant biceps active
614 breaking on successful stop trials and 8/8 showed significant active breaking on
615 failed stop trials. In experiment 2, 5/7 participants showed significant triceps
616 active breaking in the relax condition on successful stop trials and 6/7 showed
617 significant active breaking on failed stop trials. For the contract condition 8/11
618 showed significant triceps active breaking on successful stop trials and 11/11
619 showed significant active breaking on failed stop trials.

620 Thus, with one exception (exp.1 contract triceps condition, successful
621 stop trials; *see discussion*), in all experiment and conditions we found that
622 muscles that relaxed in response to the go signal showed positive active
623 breaking after the stop signal, while muscles that contracted in response to the
624 go signal showed negative active breaking.

625

626 **Active breaking onset time correlated with SSRT**

627 If active breaking is causally involved in stopping its onset time should be
628 positively correlated with SSRT across participants. In experiment 1 (Fig. 2. K.
629 & L.), for the relax condition there was a moderate positive relationship between
630 kinematic active breaking onset time and SSRT ($r = 0.54$, $p = 0.054$) and

631 between biceps EMG active breaking onset time and SSRT ($r = 0.5$, $p = 0.085$),
 632 both of which reached significance (1-tailed). We did not observe significant
 633 correlations in the contract condition, either for kinematic active breaking and
 634 SSRT ($r = 0.11$, $p = 0.714$) or triceps EMG active breaking and SSRT ($r = 0.13$,
 635 $p = 0.6712$).

636 In experiment 2 (Fig. 4. K. & L.), while relax condition kinematic active
 637 breaking did not correlate with SSRT ($r = 0.2$, $p = 0.523$), biceps EMG active
 638 breaking onset time did correlate significantly with SSRT ($r = 0.66$, $p = 0.02$).
 639 Moreover, in the contract condition there was a significant correlation between
 640 kinematic active breaking onset time and SSRT ($r = 0.73$, $p = 0.007$), and a
 641 trend towards a significant correlation between biceps EMG active breaking and
 642 SSRT ($r = 0.46$, $p = 0.131$).

643 Finally, in experiment 3 (Fig. 6. I. & J.) all measures of active breaking
 644 onset time significantly correlated with SSRT. This held for force ($r = 0.67$, $p =$
 645 0.0088) and EMG ($r = 0.67$, $p = 0.0082$) in the relax condition and for force ($r =$
 646 0.83 , $p < 0.001$) and EMG ($r = 0.7$, $p = 0.005$) in the contract condition. Thus
 647 overall we found support for a relationship between active breaking onset time
 648 and SSRT for muscle relaxations, but weaker support for this relationship when
 649 stopping muscle contractions.

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652 Figure 5

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656 Table 5

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660 Active breaking in the absence of detectable Go response

661 Splitting the successful stop trials into quartiles based on the amount of
662 putative 'go response' present revealed that even when the arm kinematics or
663 finger force level were flat, active breaking was still detected after the stop
664 signal. This was particularly true in the relax condition, where EMG velocity in
665 the fourth quartile (least go response) was significantly positive in experiment 1
666 ($t(12) = 3.347$, $p = 0.0058$, Cohen's $d = 0.93$; Fig. 2. G. & I.), experiment 2 (t
667 (11) = 2.370, $p = 0.037$, Cohen's $d = 0.68$; Fig. 4. G. & I.) and experiment 3 (t
668 (13) = 3.564, $p = 0.0035$, Cohen's $d = 0.95$; Fig. 6. E. & G.). For muscle
669 contractions, EMG velocity in the fourth quartile was significantly negative in
670 experiment 3 ($t(13) = -3.23$, $p = 0.0066$, Cohen's $d = 0.86$; Fig. 6. F. & H.), was
671 not significant in experiment 2 ($t(11) = -0.5$, $p = 0.626$, Cohen's $d = 0.14$; Fig. 4.
672 H. & J.) and showed a trend towards being significant in experiment 1 ($t(12) =$
673 1.958, $p = 0.074$, Cohen's $d = 0.54$; Fig. 2. H. & J).

674

675 Active breaking amplitude decreased as Go response decreased

676 The quartile analysis showed that amount of active breaking decreased
677 as the amount of detected 'go response' decreased. This was true for the relax
678 ($F(3,48) = 10.361$, $p < 0.001$, $\eta^2 = 0.61$; Fig. 2. I.) and contract ($F(3,48) =$
679 8.142, $p < 0.001$, $\eta^2 = 0.4$; Fig. 2. J.) conditions in experiment 1, the relax (F
680 (3,44) = 4.734, $p = 0.007$, $\eta^2 = 0.3$; Fig. 4. I.) and contract ($F(3,44) = 9.9$, $p <$
681 0.001, $\eta^2 = 0.47$; Fig. 4. J.) conditions in experiment 2, and the contract (F
682 (3,52) = 10.393, $p < 0.001$, $\eta^2 = 0.44$; Fig. 6. H.) condition in experiment 3. The

683 relax condition in experiment 3 had active breaking which did not systematically
684 differ across quartiles ($F(3,52) = 1.328$, $p = 0.279$, $\eta p^2 = 0.09$; Fig. 6. G.).
685 Overall, the results indicate that active breaking amplitude was proportional to
686 the degree of go response at the muscle.

687

688 **Comparing active breaking in relax and contract conditions**

689 EMG active breaking onset times were around 80-100ms across
690 experiments (Table 2., 4. and 6.). A notable exception were successful stop
691 trials in the contract condition experiment 2, where negative active breaking
692 onset times were around 140ms for both muscles (Table 4.). Direct comparison
693 shows that active breaking onset times for the biceps muscle were significantly
694 slower on successful than on failed stop trials (144.29ms vs 91.42ms; $t(11) =$
695 3.62 , $p = 0.004$, Cohen's $d = 1.51$). However, the comparison is not strictly
696 appropriate, since we did not observe active breaking on all successful stop
697 trials when the data was split into quartiles (Fig. 4. J.). As such, we did not
698 compare active breaking across movement conditions in experiment 1 and 2.
699 The comparison was justified in experiment 3, where active breaking was
700 significant in all cases, but no significant difference was observed when mean
701 onset times were compared between relax and contract conditions (94.41ms vs
702 97.88ms; $t(13) = -0.732$, $p = 0.477$, Cohen's $d = -0.18$).

703

704 **Active breaking involves simultaneous action across muscles**

705 In experiment 1 active breaking was detected on average 18ms earlier in
706 the biceps than the triceps muscle ($t(7) = -3.88$, $p = 0.006$) in the relax
707 condition. However, when the entire active breaking waveform was compared in

708 both muscles via time-shifted cross-correlation, strong positive correlation was
 709 observed peaking at -6ms (mean peak $r = 0.83$, $SD = 0.09$; Fig. 2. M.). This
 710 suggests that the biceps may have contracted just before the triceps, but that
 711 both muscles were likely driven by a single input, rather than the biceps activity
 712 causing the triceps activity. This was confirmed in the other conditions. In the
 713 contract condition in experiment 1 the mean onset of the triceps and biceps
 714 active breaking did not significantly differ ($t(7) = 0.764$, $p = 0.47$). Cross-
 715 correlation was weaker than in the relax condition (mean peak $r = 0.56$, $SD =$
 716 0.17), but the peak correlation was again close to 0 (2.5ms; Fig. 2. M.),
 717 indicating simultaneous action. Moreover, in experiment 2 biceps and triceps
 718 active breaking onset times did not differ from one another in the relax ($t(6) = -$
 719 1.124 , $p = 0.304$) or contract conditions ($t(10) = -0.149$, $p = 0.884$) and cross-
 720 correlation suggested simultaneous action in both cases (relax peak = 1ms, $r =$
 721 0.69 , $SD = 0.22$; contract peak = -3ms, $r = 0.61$, $SD = 0.26$; Fig. 4. M.). Thus,
 722 regardless of whether active breaking was positive or negative, our results
 723 suggest it occurred simultaneously across the observed muscles in each
 724 condition.

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727 Figure 6

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731 Table 6

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735 **Discussion**

736 Muscle relaxations were inhibited as rapidly as muscle contractions
737 across all experiments. Our behavioural results were consistent with the race
738 model, which argues that stop and go processes separately race to completion
739 (Logan and Cowan, 1984). On this account successful stopping will become
740 less frequent as the delay between go and stop signal increases (Logan, 1994)
741 because increasing the delay gives the go process more time to finish. This was
742 confirmed in experiment 1 and 2. The resulting SSD functions were similar to
743 those obtained in previous studies (Logan et al., 1984; Schall et al., 2017). In all
744 conditions tested, mean response times for unsuccessful stop trials were
745 significantly faster than those for go trials. If there had been prolonged
746 interaction between stop and go processes, then we would have observed the
747 opposite result as the presence of the stop signal would slow the go response.
748 But our results were consistent with stop/go independence. Failed stop trials
749 were faster because for a given probability of stopping (e.g. 0.5), the
750 unsuccessful stop trials must be drawn predominantly from the left side (faster
751 50% of responses) of the RT distribution, while the go trial RT reflects the mean
752 of the entire distribution.

753 We found SSRTs on the order of 150-300ms for muscle relaxation and
754 muscle contraction conditions, consistent with previous literature (Atsma et al.,
755 2018; Verbruggen and Logan, 2008). Existing work had suggested shared
756 GABAergic control in M1 might underpin the initiation of muscle relaxation and
757 the stopping of actions (Buccolieri et al., 2004; Coxon et al., 2006; Kato et al.,
758 2019; Motawar et al., 2012). However, we found that relax SSRT did not differ
759 from contract SSRT. This lack of interference-related slowing in the relax

760 condition argues against shared GABAergic control. Other pathways are known
761 to be involved in action inhibition. Of particular importance is the fronto-striatal
762 hyperdirect pathway, which is involved in the reactive breaking of action (Aron
763 and Poldrack, 2006; Aron et al., 2003). A basal ganglia account of stopping
764 maps well onto the race model and can explain the inhibition of diverse
765 movement such as saccades, vocalizations, reaching, grip force and classic
766 finger movements (de Jong et al., 1990; Logan and Irwin, 2000; Hanes and
767 Carpenter, 1999; Mirabella et al., 2006; Xue et al., 2008; Bissett and Logan,
768 2012). Our finding, that relax SSRT and contract SSRT significantly correlated
769 across participants, argues for a shared control mechanism in the stopping of
770 relaxations and contractions, possibly involving this basal ganglia circuitry.

771 Action inhibition is known to involve muscle excitation as well as
772 suppression. During the cancellation of reaching, increases in motor unit firing
773 have been recorded from the pectoralis muscle (Atsma et al., 2018). Such
774 responses are termed active breaking because they involve additional input to
775 the muscle which opposes the go response (Kudo and Ohtsuki, 1998;
776 Goonetilleke et al., 2010, 2012). As predicted, we found evidence that this
777 mechanism operates during the stopping of muscle relaxations. Positive active
778 breaking was consistently found across all three experiments both at the group
779 and individual participant level for failed and successful stop trials. These
780 transient increases in surface EMG could be detected even on the subset of
781 successful stop trials where there was no detectable go signal (i.e.
782 kinematics/force was flat). Moreover, in all experiments we found a significant
783 positive correlation between the onset time of EMG active breaking and SSRT

784 in the relax condition. Active breaking may play a causative role in the stopping
785 of muscle relaxations, akin to the theorised peripheral stopping system, which
786 counters unwanted activity that escapes the motor cortex (De Jong et al., 1995;
787 de Jong et al., 1990).

788 We also found transient *decreases* in EMG when muscle contractions
789 were cancelled. EMG decreases are expected when the excitatory drive to the
790 muscle is interrupted, and the resulting partial responses have been reported
791 during the stop signal task (de Jong et al., 1990; McGarry and Franks, 1997;
792 McGarry et al., 2000). However, our analysis showed that EMG did not simply
793 cease to increase on these stop trials, rather, the velocity of EMG changed
794 direction and quickly became significantly negative, suggesting a suppressive
795 input to the muscle (Coxon et al., 2006). Again the effect was detectable at the
796 group and individual subject level across tasks. We therefore propose that this
797 form of response be termed *negative* active breaking. The relationship between
798 negative active breaking onset time and SSRT was less consistent than that
799 seen for positive active breaking, with correlations being significant in
800 experiment 3, but only showing a trend in experiment 2. Negative active
801 breaking onset times did not differ from positive active breaking times in
802 experiment 3, where a direct comparison was justified due to active breaking
803 being present under all conditions, consistent with the similar SSRTs observed
804 across conditions. However, negative active breaking was noticeably slower in
805 experiment 2 for successful stop trials, perhaps because proximal muscles,
806 unlike distal muscles, only show negative active breaking when the muscle is
807 actively shortening, due to control differences (Serrien and Baeyens, 2017).

808 However, we cannot rule out other explanations such as increased onset
809 variability at the trial-level.

810 Active breaking was muscle state specific. If the go command caused the
811 muscle to relax then active breaking was positive, while if the go command
812 caused contraction then active breaking was negative. However, in the contract
813 condition in experiment 1 we found negative active breaking on failed stop trials
814 and positive active breaking on successful stop trials. This was likely because
815 the biceps muscle began to relax significantly earlier in relation to the go signal
816 than the triceps started to contract; an EMG-RT difference that has previously
817 been observed (Buccolieri et al., 2003). Consequently, stop commands on
818 successful trials tended to be executed before the triceps began to contract, but
819 as the relax command was starting to be initiated. The critical factor determining
820 active breaking direction therefore may be whether a relaxation command is
821 present centrally rather than at the muscle itself. Under such conditions the
822 observed active breaking response may be a mix of positive and negative forms,
823 perhaps explaining the lack of a significant correlation with SSRT in experiment
824 1.

825 Active breaking was previously observed in antagonist muscles (Atsma
826 et al., 2018). It was therefore unclear if the excitatory signal originates centrally
827 or from the agonist muscle. We found active breaking in the triceps and biceps
828 muscle in all conditions, with independently derived onset times that were
829 similar regardless of the type of go command. When we took the entire active
830 breaking EMG waveform for each muscle and performed time-shifted cross
831 correlation, we found that in all conditions there was a positive correlation and

832 the peak was close to 0ms. So activity in one muscle did not predict the activity
833 of the other at some time-delay. Rather, both muscles were likely driven by
834 central signals and acted simultaneously.

835 Simultaneous positive active breaking functionally resembles co-
836 contraction, in in that it prevents unwanted movement by increasing joint
837 stiffness (Osu and Gomi, 1999). However, when active breaking was
838 concurrently negative in one muscle and positive in the other, the analogy with
839 co-contraction is less clear. Positive and negative active breaking may involve a
840 common cortico-striatal pathway, but diverge afterwards. Negative active
841 breaking likely results from activity in the striatal hyperdirect pathway suddenly
842 cutting excitatory drive to M1 (Aron et al., 2007), coupled with downstream
843 inhibitory activity in M1 (Badry et al., 2009; Coxon et al., 2006). Reactive
844 inhibition of this sort has a strong and generalised suppressive effect on action
845 (Aron, 2011), which accounts for why we observed reductions of baseline FDI
846 EMG in the absence of a go command and why negative active breaking
847 amplitude scaled with the current level of excitatory drive to the muscle.
848 Importantly, we show that this cutting of excitatory drive does not extend to the
849 positive active breaking signal, since both mechanisms can be observed on the
850 same trials. Positive active breaking may therefore involve a separate
851 descending pathway, perhaps driven by connections from the striatum to sub-
852 cortical postural control regions (Mena-Segovia and Bolam, 2017; Takakusaki,
853 2017) or gain increases in the gamma motoneuron system (Johansson et al.,
854 1986).

855

856 Conclusion

857 The cancelling of voluntary muscle relaxations was rapid, conformed to
858 the predictions of the race model, and correlated with the time taken to cancel
859 muscle contractions, suggesting shared high-level control. But in the periphery
860 we found evidence for a later divergence of control. Stopping relaxing muscles
861 involved transient increases in muscle activity, while stopping contractions
862 involved transient decreases in muscle activity. This active breaking correlated
863 with SSRT and occurred simultaneously in the biceps and triceps, indicating a
864 central origin. Negative active breaking, though known to have a general
865 suppressive effect on activity, did not prevent positive active breaking in other,
866 already relaxing muscles, suggesting the two forms of active breaking rely on
867 separate pathways. The automatic coordination of positive and negative active
868 breaking across muscle groups may explain how animals can rapidly inhibit
869 complex actions at different stages of their execution.

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1052 **Tables and Figures**

1053 **Table 1: Response times for experiment 1.** Table shows group mean (SD) behavioural
 1054 response times for go trials and response times for onset of muscle activity changes on go trials.
 1055 Both muscles were relaxing in the relax condition, but in the contract condition the triceps
 1056 muscle was contracting and the biceps muscle was relaxing. Behavioural response times on
 1057 stop trials where movement occurred are also shown along with stop signal reaction time
 1058 (SSRT).

Condition	Go Trial RT	Go Trial Biceps EMG-RT	Go Trial Triceps EMG-RT	Failed stop trial RT	SSRT
Relax	554.74ms (54.21ms)	311.01ms (32.3ms)	390.61ms (90.3ms)	497.02ms (39.44ms)	220.46ms (49.42ms)
Contract	521.28ms (62.02ms)	331.05ms (51.09ms)	417.18ms (59.17ms)	461.27ms (39.74ms)	238.51ms (40.09ms)

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1060 **Table 2: Active breaking onset times exp. 1.** Mean (SD) onset time of active breaking are
 1061 shown for arm acceleration and primary and secondary muscles. Active breaking onset was
 1062 calculated separately for each measure, in each movement condition, for successful and failed
 1063 stop trials. This was done at the individual participant level. Mean stop trial onset refers to the
 1064 average of successful and failed stop trial onset times. Secondary muscle data (triceps in relax
 1065 condition, biceps in contract condition) could only be analyzed in a subset of participants. Note
 1066 that triceps active breaking in the contract condition is labelled mixed because it was positive on
 1067 successful and negative on failed stop trials (see text for details).

Movement condition	Type of active breaking	Succ. stop trial onset	Failed stop trial onset	Mean stop trial onset
Relax	Positive arm acceleration	193.23ms (32.92ms)	200ms (26.98ms)	196.62ms (27.27ms)
	Positive biceps EMG velocity	83.12ms (22.04ms)	82.42ms (19.74ms)	82.77ms (18.64ms)
	Positive triceps EMG velocity (n = 8)	86.31ms (14.84ms)	94.69ms (28.31ms)	90.5ms (17.14ms)
Contract	Positive arm acceleration	199.38ms (27.37ms)	166.15ms (37.08ms)	182.77ms (22.44ms)
	Mixed triceps EMG velocity	92.62ms (30.97ms)	85.85ms (25.98ms)	89.23ms (22.5ms)
	Positive biceps EMG velocity (n = 8)	92.94ms (32.23ms)	95.81ms (24.31ms)	94.38ms (26.08ms)

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1069 **Table 3: Response times for experiment 2.** Table shows group mean (SD) behavioural
 1070 response times for go trials and response times for onset of muscle activity changes on go trials.
 1071 Both muscles were relaxing in the relax condition and contracting in the contract condition.
 1072 Behavioural response times on stop trials where movement occurred are also shown along with
 1073 stop signal reaction time (SSRT).

Condition	Go Trial RT	Go Trial Biceps EMG-RT	Go Trial Triceps EMG-RT	Failed stop trial RT	SSRT
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Relax	588.36ms (56.51ms)	331.76ms (36.9ms)	416.21ms (77.43ms)	543.98ms (41.17.ms)	268.1ms (48.31ms)
Contract	549.82ms (42.18ms)	437.79ms (33.97ms)	440.08ms (32.84ms)	504.46ms (32.47ms)	256.5ms (31.69.ms)

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1075 **Table 4: Active breaking onset times exp. 2.** Mean (SD) onset time of active breaking are
 1076 shown for arm acceleration and primary and secondary muscles. Active breaking onset was
 1077 calculated separately for each measure, in each movement condition, for successful and failed
 1078 stop trials. This was done at the individual participant level. Mean stop trial onset refers to the
 1079 average of successful and failed stop trial onset times. Secondary muscle data (triceps) could
 1080 only be analyzed in a subset of participants in the relax condition and one participant was
 1081 removed from the triceps analysis in the contract condition (see text for details).

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Movement condition	Type of active breaking	Succ. stop trial onset	Failed stop trial onset	Mean stop trial onset
Relax	Positive arm acceleration	204.33ms (29.42ms)	211.33ms (29.39ms)	207.83ms (24.16ms)
	Positive biceps EMG velocity	67.25ms (12.59ms)	86.96ms (26.46ms)	77.1ms (16.13ms)
	Positive triceps EMG velocity (n = 7)	91ms (34.87ms)	87.79ms (27.47ms)	89.39ms (25.7ms)
Contract	Negative arm acceleration	260.67ms (37.56ms)	193.33ms (28.51ms)	227ms (29.15ms)
	Negative biceps EMG velocity	144.29ms (38.84ms)	91.42ms (30.71ms)	117.85ms (24.18ms)
	Negative triceps EMG velocity (n =11)	140.73ms (42.77ms)	96.64ms (26.63ms)	118.68ms (18.14ms)

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1084 **Table 5: Response times for exp. 3.** Group mean (SD) behavioural response time for go trials,
 1085 go trial FDI EMG response time, failed stop trials behavioural response time, and stop signal
 1086 reaction time (SSRT) in the relax and contract condition.

Condition	Go Trial RT	Go Trial FDI EMG-RT	Failed stop trial RT	SSRT
Relax	598.01ms (54.8ms)	424.29ms (47.98ms)	511.11ms (55.52ms)	170.98ms (58.94ms)

Contract	595.11ms (63.59ms)	445.21ms (52.95ms)	489.73ms (53.79ms)	156.86ms (69.84ms)
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1090 **Table 6: Active breaking onset times exp. 3.** Mean (SD) onset time of active breaking are
1091 shown for force velocity and FDI EMG velocity. Active breaking onset was calculated separately
1092 for each measure, in each movement condition, for successful and failed stop trials. This was
1093 done at the individual participant level. Mean stop trial onset refers to the average of successful
1094 and failed stop trial onset times.

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Movement condition	Type of active breaking	Succ. stop trial onset	Failed stop trial onset	Mean stop trial onset
Relax	Positive force velocity	177.04ms (75.94ms)	212.79ms (62.58ms)	194.91ms (66.01ms)
	Positive FDI EMG velocity	88.21ms (23.85ms)	100.61ms (23.34ms)	94.41ms (20.05ms)
Contract	Negative force velocity	184.07ms (78.34ms)	199.54ms (29.61ms)	191.8ms (51.49ms)
	Negative FDI EMG velocity	99.04ms (14.92ms)	96.71ms (28.54ms)	97.88ms (19.42ms)

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1102 **Figure 1. Design and results for experiment 1.** Task set-up is shown (A). In Relax and
1103 Contract blocks, participants made elbow extensions in response to go signals and attempted to
1104 prevent these movements whenever a stop signal occurred. Elbow extensions were driven by
1105 biceps relaxation or triceps activation respectively, as shown in the representative go trials from
1106 the Relax (B) and Contract (C) conditions. Group mean relax and contract condition EMG and
1107 kinematic responses to Go trials (D). SSD functions for the relax and contract condition (E),
1108 showing how the probability of failing to stop on a stop trial increased as SSD increased. SSRT

1109 in relax and contract blocks significantly correlated ($r = 0.62$, $p = 0.024$) across participants (**F**).
 1110 Group mean relax and contract condition kinematics (**G**), biceps EMG (**H**), and triceps EMG (**I**),
 1111 for successful and failed stop trials, time-locked to the stop signal. Upper panels show
 1112 smoothed data, while lower panels show angle acceleration and EMG velocity traces. All error
 1113 bars are SEM.

1114

1115 **Figure 2. Active breaking results for experiment 1.** Group mean kinematic active breaking
 1116 results for successful (**A**) and failed stop trials (**B**) in the relax and contract conditions.
 1117 Successful stop trials also plotted on the failed stop trial graph for comparison. Traces always
 1118 aligned to active breaking onset times derived at the individual participant level. Inserts show
 1119 mean values obtained during the 50ms shaded window immediately after active breaking onset.
 1120 Group mean biceps EMG active breaking results for successful (**C**) and failed (**D**) stop trials for
 1121 the relax and contract conditions. Active breaking was significantly positive for all conditions.
 1122 Group mean triceps EMG active breaking results for successful (**E**) and failed (**F**) stop trials.
 1123 Active breaking was significantly positive in the relax condition and significantly negative for
 1124 failed stop trials in the contract condition. But for successful stop trials it was significantly
 1125 positive (see text for details). Successful stop trial kinematics in the relax (**G**) and contract (**H**)
 1126 conditions after trials were sorted into quartiles according to the amount of Go response.
 1127 Positive biceps EMG active breaking was present in the relax condition even when the
 1128 kinematics were flat (**I**). A trend was observed for the triceps in the contract condition (**J**). Active
 1129 breaking amplitude decreased significantly across bins in both cases, as detected go response
 1130 decreased. Positive correlations between kinematic (**K**) and EMG (**L**) active breaking onset
 1131 times and SSRT reached significance in the relax, but not the contract condition. Time-shifted
 1132 cross correlation (**M**) showed that peak correlation between biceps and triceps active breaking
 1133 was close to 0ms in both conditions, indicating simultaneous action. All error bars show SEM.
 1134 Results of statistical tests shown as asterisks (***) $p < 0.001$, ** $p < 0.01$, * $p < 0.05$.

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1136 **Figure 3. Design and results for experiment 2.** On relax go trials participants extended the
 1137 elbow by relaxing the biceps and on contract go trials they flexed the elbow by further

1138 contracting the biceps. They attempted to stop these movements if a stop signal was present
 1139 **(A)**. Group mean relax and contract condition EMG and kinematic responses to Go trials **(B)**.
 1140 SSD functions for the relax and contract condition **(C)**, showing how the probability of failing to
 1141 stop on a stop trial increased as SSD increased. SSRT in relax and contract conditions showed
 1142 a positive trend ($r = 0.55$, $p = 0.061$) across participants **(D)**. Group mean relax and contract
 1143 condition kinematics **(E)**, biceps EMG **(F)**, and triceps EMG **(G)**, for successful and failed stop
 1144 trials, time-locked to the stop signal. Upper panels show smoothed data, while lower panels
 1145 show angle acceleration and EMG velocity traces. All error bars are SEM.

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1147 **Figure 4. Active breaking results for experiment 2.** Group mean kinematic active breaking
 1148 results for successful **(A)** and failed stop trials **(B)** in the relax and contract conditions.
 1149 Successful stop trials also plotted on the failed stop trial graph for comparison. Inserts show
 1150 mean values obtained during the 50ms window immediately after active breaking onset. Group
 1151 mean biceps EMG active breaking results for successful **(C)** and failed **(D)** stop trials, and mean
 1152 triceps EMG active breaking results for successful **(E)** and failed **(F)** stop trials. Active breaking
 1153 was always significantly positive in the relax condition and significantly negative in the contract
 1154 condition. Successful stop trial kinematics in the relax **(G)** and contract **(H)** conditions after
 1155 sorting by amount of detected Go response. Positive biceps EMG active breaking was present
 1156 in the relax condition even when the kinematics were flat **(I)**. In the contract condition **(J)**
 1157 negative active breaking was not significant in B4. Active breaking amplitude decreased
 1158 significantly across bins in the relax and contract conditions. Positive correlations between
 1159 kinematic **(K)** and EMG **(L)** active breaking onset times and SSRT reached significance in the
 1160 relax condition EMG, but only showed a trend in other cases. Time-shifted cross correlation **(M)**
 1161 showed that peak correlation between biceps and triceps active breaking was close to 0ms in
 1162 both conditions, indicating simultaneous action. All error bars show SEM. Results of statistical
 1163 tests shown as asterisks (***) $p < 0.001$, ** $p < 0.01$, * $p < 0.05$.

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1165 **Figure 5. Design and results for experiment 3.** On relax go trials participants reduced force
 1166 by relaxing the FDI and on contract go trials they increased force by further contracting the FDI.

1167 They attempted to stop these force changes if a stop signal was present **(A)**. Group mean relax
1168 and contract condition EMG and force responses to Go trials **(B)**. SSRT in relax and contract
1169 blocks showed a significant positive correlation ($r = 0.82$, $p < 0.001$) across participants **(D)**.
1170 Group mean relax and contract condition force **(E)** and FDI EMG **(F)** for successful and failed
1171 stop trials, time-locked to the stop signal. Upper panels show smoothed data, while lower panels
1172 show velocity traces. Error bars show SEM.

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1174 **Figure 6. Active breaking results for experiment 3.** Group mean force active breaking results
1175 for successful **(A)** and failed stop trials **(B)** in the relax and contract conditions. Inserts show
1176 mean values obtained during the 50ms window immediately after active breaking onset. Group
1177 mean FDI EMG active breaking results for successful **(C)** and failed **(D)** stop trials. Active
1178 breaking was always significantly positive in the relax condition and significantly negative in the
1179 contract condition. Successful stop trial force in the relax **(E)** and contract **(F)** conditions after
1180 sorting by amount of detected Go response. Positive FDI EMG active breaking was present in
1181 the relax condition even when the force traces were flat **(G)** and negative active breaking was
1182 present in the contract condition when force traces were flat **(H)**. Active breaking amplitude
1183 decreased significantly across bins in the contract, but not relax condition. Significant positive
1184 correlations were found between force **(I)** and EMG **(J)** active breaking onset times and SSRT in
1185 the relax and contract conditions. All error bars show SEM. Results of statistical tests shown as
1186 asterisks (** $p < 0.001$, ** $p < 0.01$, * $p < 0.05$).











