

1 Muscle synergies are flexibly recruited during gait pattern
2 exploration using motor control-based biofeedback

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27 **ABSTRACT**

28 Understanding how the central nervous system coordinates diverse motor outputs has been a topic of
29 extensive investigation. While it is generally accepted that a small set of synergies underlies many
30 common activities, such as walking, whether synergies are equally robust across a broader array of gait
31 patterns or can be flexibly modified remains unclear. Here, we evaluated the extent to which synergies
32 changed as nondisabled adults ($n = 14$) explored gait patterns using custom biofeedback. Secondarily, we
33 used Bayesian Additive Regression Trees to identify factors which were predictive of synergy
34 modulation. Participants performed 41.1 ± 8.0 gait patterns using biofeedback, during which synergy
35 recruitment changed depending on the type and magnitude of gait pattern modification. Specifically, a
36 consistent set of synergies was recruited to accommodate small deviations from baseline, but additional
37 synergies emerged for larger gait changes. Synergy complexity was similarly modulated; complexity
38 decreased for 82.6% of the attempted gait patterns, however, distal gait mechanics were highly predictive
39 of these changes. In particular, greater ankle dorsiflexion moments and knee flexion through stance, as
40 well as greater knee extension moments at initial contact corresponded to a reduction in synergy
41 complexity. Taken together, these results suggest that the central nervous system preferentially adopts a
42 low-dimensional, largely invariant control strategy, but can modify that strategy to produce diverse gait
43 patterns. Beyond improving understanding of how synergies are recruited during gait, study outcomes
44 may also help identify parameters that can be targeted with interventions to alter synergies and improve
45 motor control following neurological injury.

46

47 1. INTRODUCTION

48 Humans are capable of producing a broad array of movements, allowing for robust locomotion in diverse
49 and unpredictable environments. To achieve this range of motor outputs, it has been hypothesized that the
50 central nervous system (CNS) recruits a small number of synergies (i.e., modes, modules), defined as
51 groups of coactivating muscles; this architecture is believed to simplify control beyond activating muscles
52 independently (Bizzi and Cheung, 2013; Ting et al., 2015; Tresch and Jarc, 2009). Numerous studies have
53 evaluated this hypothesis experimentally, employing matrix decomposition techniques, such as non-
54 negative matrix factorization, to extract synergies and their corresponding activation patterns from
55 electromyography (EMG) data (Lee, 1999; Tresch, 2005). These studies revealed that tasks such as
56 walking (Allen and Neptune, 2012; Ivanenko et al., 2004), running (Cappellini et al., 2006; Hagio et al.,
57 2015) and cycling (Barroso et al., 2014) share a small set of muscle synergies, despite being
58 biomechanically distinct. Further, across tasks, changes in speed (Rozumalski et al., 2017), incline
59 (Ivanenko et al., 2004; Rozumalski et al., 2017), cadence (Rouston et al., 2014), and body-weight loading
60 (Ivanenko et al., 2004; McGowan et al., 2010) are shown to shift the phase or duration of synergy
61 activations rather than the structure of the synergies themselves. These observations suggest that modest
62 changes in sensory input or biomechanical demand are accommodated by altering the activation of
63 invariant synergies and lend credence to their centralized role in coordination (Cheung et al., 2005;
64 Torres-Oviedo and Ting, 2010). However, whether synergies are equally robust across a greater subset of
65 achievable gait patterns or can be actively modified during gait is largely unknown (Jason J. Kutch and
66 Valero-Cuevas, 2012; Tresch and Jarc, 2009).

67 Because synergies generally align with the sub-tasks of walking (e.g., push-off, weight
68 acceptance), gait patterns which impose additional mechanical requirements or present large changes in
69 somatosensory feedback may alter synergy recruitment (Cheung et al., 2005; Ivanenko et al., 2005; Nazifi
70 et al., 2017; Torres-Oviedo and Ting, 2010). This is supported by prior work in animal models which
71 demonstrated that frogs recruit task-specific synergies during swimming, jumping, and walking which

72 correspond to the unique biomechanical demands of each movement (d'Avella and Bizzi, 2005).
73 Similarly, humans recruit specific synergies during perturbation recovery tasks to maintain mediolateral
74 stability and reduce center of mass movement (Krishnamoorthy et al., 2004; Martino et al., 2015; Nazifi
75 et al., 2017; Torres-Oviedo and Ting, 2010). Importantly, such synergies emerge in addition to those
76 shared with other tasks, which suggests that the CNS flexibly draws from a limited library rather than
77 deploying unique control strategies to accommodate task demand (Torres-Oviedo and Ting, 2010).

78 Taken together, prior results indicate that a relationship exists between the biomechanical
79 constraints of a given task and the recruited control strategy. That is, the CNS may preferentially tune the
80 activation timing of a consistent set of synergies but is simultaneously capable of recruiting different
81 synergies to produce diverse outputs. Understanding when and how synergies are modulated across
82 changing biomechanical contexts and the factors driving this modulation is critical to better inform how
83 the CNS coordinates complex movement. While this relationship has been previously characterized
84 across broad balance (Torres-Oviedo and Ting, 2010) and finger force generation tasks (Jason J Kutch
85 and Valero-Cuevas, 2012; Valero-Cuevas et al., 2009), gait has not been studied to the same extent
86 (Rouston et al., 2014; Zelik et al., 2014).

87 Beyond enhancing understanding of the neural control of gait, characterizing whether synergies
88 can be modulated in walking may also inform methods for targeting aberrant synergy recruitment.
89 Individuals with cerebral palsy (Schwartz et al., 2016; Steele et al., 2015; Tang et al., 2015), Parkinson's
90 disease (Rodriguez et al., 2013), and spinal cord injury (Fox et al., 2013) as well as stroke survivors
91 (Cheung et al., 2012; Clark et al., 2010) recruit fewer synergies than nondisabled peers which impacts
92 independent mobility (Bowden et al., 2010; Clark et al., 2010; Mehrabi et al., 2019) and may reduce the
93 efficacy of traditional interventions (Schwartz et al., 2016). Because available interventions for these
94 populations often fail to alter synergies (Shuman et al., 2019), developing new paradigms to directly
95 improve synergy recruitment has become a critical priority in gait rehabilitation. This has spurred the
96 development of biofeedback and robotic gait training paradigms which have thus far yielded promising,

97 yet still highly variable results (Booth et al., 2019; Conner et al., 2021; Rouston et al., 2013). As such,
98 mapping the relationship between biomechanical constraints and synergy modulation may further inform
99 the design of these systems by highlighting gait parameters that can be directly targeted to produce greater
100 and more consistent changes in motor control.

101 The aim of this study was to characterize the robustness of synergies to changing biomechanical
102 constraints during walking. Specifically, we evaluated the extent to which nondisabled individuals could
103 modulate both synergy structure and complexity during walking while using motor control biofeedback to
104 drive broad gait pattern exploration. These data were then used to build a Bayesian Additive Regression
105 Trees (BART) model to identify biomechanical variables that were predictive of synergy modulation. We
106 hypothesized that changing biomechanical constraints would alter the recruitment but not the structure of
107 muscle synergies, but that different synergies may be recruited to accommodate large deviations from
108 baseline. The results from this investigation will provide further insight into the extent to which motor
109 control can be altered and, importantly, improve understanding of how the CNS shapes its control
110 strategy to produce a repertoire of motor outputs. The latter will support the development of intervention
111 strategies to improve motor control among individuals with neurological injury.

112 **2. METHODS**

113 **2.1 Experimental Protocol**

114 Fourteen nondisabled individuals (7M/7F; Age: 24.1 ± 4.7 years; Height: 1.7 ± 0.1 m; Mass: 65.7 ± 20.1
115 kg) were recruited to evaluate synergies during gait pattern exploration. Prior to participation, all provided
116 written informed consent and the experimental protocol was approved by the University of Washington
117 Institutional Review Board.

118 Participants walked on a treadmill at a self-selected speed (1.07 ± 0.13 m/s; Bertec, Columbus,
119 OH) while responding to a custom biofeedback system, designed to encourage gait pattern exploration.
120 Briefly, this system presented the participant with a real-time score of their dominant-limb synergy

121 complexity, defined as the total variance accounted for by one synergy, on a graphical display (Steele et
122 al., 2015). To facilitate participant interpretation, the displayed score was normalized to baseline walking
123 and scaled such that a value of 100 corresponded to baseline and higher values indicated more complex
124 control (see S1 for additional system details). Participants performed one baseline walking trial with the
125 feedback system turned off followed by feedback trials during which they were instructed to either (1)
126 raise or (2) lower their complexity score; two trials were performed in each target direction. All trials
127 were three minutes long and separated by mandatory one-minute rest periods. During the feedback trials,
128 participants were encouraged to explore a broad range of gait patterns to modify their score. The only
129 imposed restrictions were that they must (1) maintain forward-facing walking and (2) take at least five
130 consecutive strides in the pattern selected.

131 Surface EMG data (Delsys Inc, Natick, MA) were recorded bilaterally for seven lower limb
132 muscles: gluteus maximus (GM), lateral hamstrings (LH), medial hamstrings (MH), vastus medialis
133 (VM), soleus (SO), tibialis anterior (TA), and medial gastrocnemius (MG). Raw EMG signals were low
134 passed filtered (4th order Butterworth; 20 Hz), rectified, and high pass filtered (4th order Butterworth; 10
135 Hz) to establish linear envelopes (Shuman et al., 2017). After filtering, non-physiological signal spikes
136 were removed using a robust-PCA algorithm (Lin et al., 2013) and the data were normalized to the 95th
137 percentile of maximum muscle activity across all trials.

138 Full-body motion data were collected using a 10-camera motion capture system (120 Hz) and a
139 modified Helen Hayes marker set (Kadaba et al., 1990). Joint kinematics and kinetics were derived from
140 marker data in OpenSim v3.3 using a 33 degree-of-freedom model, scaled to each subject (Delp et al.,
141 2007; Rajagopal et al., 2016). The average root-mean-squared (RMS) and maximum error for all
142 developed models were 1.3 cm and 2.5 cm, respectively, which fall below the recommended thresholds
143 for model fidelity (Hicks et al., 2015).

144 2.2 Gait Analysis

145 Because participants explored many different gait patterns using the biofeedback system, we first had to
146 extract each pattern attempted across trials and participants (Figure 1). To do this, the gait deviation index
147 (GDI) was calculated from the kinematic data for every stride in each trial (Schwartz and Rozumalski,
148 2008). The GDI is a summary measure of deviations in pelvis, hip, knee, and ankle kinematics from
149 ‘normative’ trends and was, therefore, expected to change during gait pattern exploration. For each trial,
150 groups of five or more consecutive strides with similar GDI values were automatically labeled as unique
151 gait patterns; each unique pattern identified was then subsequently confirmed via manual inspection to
152 ensure appropriate labeling. Following labeling, average kinematic and kinetic trends at the pelvis, hip,
153 knee, and ankle were quantified for each unique pattern. To identify kinematically-similar strategies
154 adopted by multiple participants, the average kinematics for all unique patterns were separated into
155 clusters (K_1 to K_N) using k-means clustering (Rozumalski and Schwartz, 2009).

156 2.3 Synergy Analysis

157 Muscle synergies were quantified from EMG data for each unique gait pattern using non-negative matrix
158 factorization (NMF). NMF is a linear matrix decomposition technique which is commonly used to
159 identify non-negative synergies (W) and their corresponding activations (C) from EMG data, such that
160 $EMG_{mxt} = W_{mxi} * C_{ixt} + error$ where m is the number of muscles, i is the number of synergies, and t is the
161 time points (Lee, 1999; Ting and Chvatal, 2010). The structure of the W and C matrices provide insight
162 into how muscles coactivate across the gait cycle. Similarly, the total variance accounted for (tVAF) by a
163 given number of synergies (i) can provide a summary measure of synergy complexity and has been
164 frequently used as a marker for impairment level; individuals with neurological injury (Cheung et al.,
165 2012; Clark et al., 2010; Fox et al., 2013; Rodriguez et al., 2013; Schwartz et al., 2016; Steele et al.,
166 2015) have higher tVAF values (*i.e.* less complex control) for a given synergy solution (i) than
167 nondisabled peers. Therefore, if synergies were sensitive to imposed biomechanical constraints, we may
168 expect to see changes in both synergy structure and complexity measures.

169 We calculated $i = 1$ to 7 synergies using the dominant-limb EMG data for five concatenated
170 strides for each unique gait pattern attempted. Because synergy analysis is sensitive to the amount of
171 EMG data used, we elected to analyze a consistent number of strides across all patterns and participants
172 (Oliveira et al., 2014). If a participant took more than five strides in a unique pattern, we performed a
173 bootstrapping procedure by quantifying synergies using five random strides, selected with replacement
174 from the available set, and replicating this process until a normal distribution was achieved; average
175 synergies and tVAF values were then reported. The same bootstrapping procedure was performed on the
176 baseline walking data with sets of five concatenated strides (replicates = 200) to ensure accurate
177 comparisons between baseline and feedback conditions.

178 We evaluated synergy structure during gait pattern exploration in two ways. We first compared
179 the inter-cluster (K_1 to K_N) similarity of synergy weights (W) and activation patterns (C) for the $i = 3$
180 synergy solution. This solution was evaluated, as three synergies explained over 90% of the variance in
181 EMG data for the majority of unique gait patterns. We sorted synergy weights for all unique gait patterns
182 attempted during exploration as well as baseline walking into k clusters (MacQueen, 1967). Because
183 individuals may recruit different synergies during exploration compared to baseline gait, we varied k
184 between $k = 3$ (i.e., synergies were consistent between baseline and exploration) and $k = 10^*3$ (i.e.,
185 different synergies emerged during exploration) and selected k as the number of clusters with the
186 maximum silhouette coefficient (Rousseeuw, 1987); the upper bound on k was highly conservative and
187 based on our expectation that synergies would be predominantly shared across gait patterns (Torres-
188 Oviedo and Ting, 2010). Synergy weights and activations for each unique gait pattern were then sorted
189 into their respective clusters (K_1 to K_N) and the average values were calculated. Secondarily, we evaluated
190 the intrasubject similarity of baseline synergies with those recruited during exploration. This was done by
191 fixing the W matrix as the synergy weights extracted from baseline walking for the three-synergy solution
192 ($i = 3$) and using the multiplicative update rule from NMF to find a C matrix which minimized the error
193 between W^*C and the EMG data for each unique gait pattern that an individual attempted. From this, we

194 were able to calculate the total variance that could be explained in each unique gait pattern by baseline
195 weights ($tVAF_{3_BASE}$) which was then compared to the $tVAF_3$ values (i.e., those calculated directly from
196 the EMG data for each unique gait pattern), yielding a measure of synergy similarity. If similar synergies
197 were recruited during gait pattern exploration and baseline walking, we would expect $tVAF_{3_BASE}$ and
198 $tVAF_3$ to be similar.

199 **2.4 Statistical Analysis**

200 **2.4.1 Cluster-wise comparisons**

201 For each cluster (K_1 to K_N), we compared mean $tVAF$ values to (1) baseline walking and (2) $tVAF_{BASE}$
202 using paired t-tests to evaluate if synergy complexity or structure changed during gait pattern exploration,
203 respectively. Secondarily, one-way ANOVA tests were used to compare if synergy complexity and
204 structure were similar between clusters (K_1 to K_N); for any test that reached significance, t-tests were used
205 to perform pairwise comparisons. Average kinematic trends at key phases within the gait cycle (e.g.,
206 push-off, initial contact) for each cluster (K_1 to K_N) were also compared to baseline walking using paired
207 t-tests. To characterize stride-to-stride variability, the standard deviation of each kinematic parameter
208 during exploration was also compared to baseline using paired t-tests. For all comparisons to baseline
209 walking and post-hoc analyses, p-values were adjusted using a Holm-Šídák correction to account for
210 multiple tests. We defined significance as $p < \alpha$ for $\alpha = 0.05$ and report mean values ± 1 SD unless
211 otherwise indicated. All cluster-wise statistical analysis was performed using the MATLAB Statistical
212 Toolbox (MathWorks, Natick, USA).

213 **2.4.2. BART analysis**

214 To further examine the relationship between gait pattern exploration and synergy complexity, we
215 developed a Bayesian Additive Regression Trees (BART) statistical model (Chipman et al., 2010). BART
216 is a ‘sum-of-trees’ machine learning method used for non-parametric function estimation, similar to other
217 techniques such as boosting (Freund and Schapire, 1997; Schapire, 1990) and random forests (Breiman,
218 2001). However, unlike other methods, BART uses a regularization prior to control tree depth and

219 shrinkage, effectively constraining individual trees as ‘weak learners’ to prevent data overfitting
220 (Chipman et al., 2010; Kapelner and Bleich, 2016). BART was selected for this application due to its
221 favorable predictive performance compared to other machine learning algorithms and because it can
222 capture the non-linear relationships inherent in motion data (Chipman et al., 2010; Dorie et al., 2019; Tan
223 and Roy, 2019).

224 We developed a BART model to predict changes in synergy complexity during exploration
225 compared to baseline walking, quantified as the difference in total variance accounted for by a one-
226 synergy solution (i.e., $\Delta tVAF_1$). Our predictor set (Table 1) included kinematic and kinetic variables that
227 characterized each unique gait pattern as well as other metrics which could influence the type of gait
228 patterns a participant attempted. When defining kinematic and kinetic predictor variables, we prioritized a
229 set that captured salient trends at the pelvis, hip, knee, and ankle, while simultaneously maintaining
230 predictor set conciseness. These criteria resulted in the variables outlined in Figure 2 (n = 31). For each of
231 the identified kinematic and kinetic variables, both the mean and standard deviation values are included in
232 the predictor set, normalized to baseline walking. We elected to include standard deviation measures in
233 the model, as $tVAF_1$ is sensitive to the amount of variance in the data and could, therefore, be affected by
234 individuals simply moving with greater stride-to-stride variability, as might be expected during novel gait
235 pattern exploration (Sawers et al., 2015). We tuned hyperparameters for the developed BART model
236 using 10-fold cross-validation (parameters: k = 3, q = 0.9, nu = 3, num_trees = 200, seed = 30) and report
237 both pseudo- R^2 and the out-of-sample root-mean-squared error (RMSE) as metrics of model quality.

238 Outputs from the BART model were interpreted with accumulated local effect (ALE) plots
239 (Apley and Zhu, 2020). ALE plots are used to visualize the effect that individual predictors have on the
240 specified response variable (i.e., $\Delta tVAF_1$), conditioned on all other model covariates. Unlike partial
241 dependence plots, which are also commonly used, ALE plots are generated by averaging and
242 accumulating the local rather than marginal effects of each predictor, making them unbiased in cases

243 where predictors are highly correlated; this is particularly advantageous for this application, due to the
244 high level of correlation between kinematic and kinetic variables during gait.

245 Because ALE plots are generated by sampling from the available data, some discrepancy between
246 the ‘true’ and ‘estimated’ effect is expected (Apley and Zhu, 2020). To capture this uncertainty, we
247 performed a bootstrap analysis ($n = 100$ replicates), drawing samples with replacement from the original
248 data set to generate a series of ALE plots from which the average and standard deviation could be
249 quantified. Using these average plots, we approximated net effects for each predictor as the difference
250 between the 95th and 5th percentile of the response. If synergies were sensitive to biomechanical
251 constraints during gait pattern exploration, we would expect both kinematic and kinetic variables to have
252 large net effects on $\Delta tVAF_1$. BART model development and ALE plot generation were performed in
253 RStudio (RStudio Team, 2020) using the *bartMachineCV* and *ALEPlot* packages (Apley and Zhu, 2020;
254 Kapelner and Bleich, 2016).

255 **3. RESULTS**

256 **3.1 Gait Exploration**

257 Participants explored 10.3 ± 2.8 unique gait patterns per feedback trial on average, resulting in 575 total
258 patterns across all participants. These data were separated into five clusters, representing the common
259 kinematic strategies attempted (Figure 3). K_2 and K_4 represented 24 and 78 unique gait patterns,
260 respectively, and were characterized by increased hip flexion ($K_2: 47.7 \pm 11.3^\circ$; $K_4: 23.1 \pm 8.8^\circ$), knee
261 flexion ($K_2: 70.6 \pm 9.1^\circ$; $K_4: 40.9 \pm 11.1^\circ$), hip abduction ($K_2: 7.4 \pm 7.0^\circ$; $K_4: 7.7 \pm 7.6^\circ$), anterior pelvic
262 tilt ($K_2: 14.1 \pm 6.6^\circ$; $K_4: 8.4 \pm 7.9^\circ$), and ankle dorsiflexion ($K_2: 20.7 \pm 2.7^\circ$; $K_4: 15.6 \pm 5.2^\circ$) through
263 stance compared to baseline. K_3 represented 98 unique gait patterns defined by greater anterior pelvic tilt
264 ($2.8 \pm 3.7^\circ$), hip abduction ($3.9 \pm 4.2^\circ$), and plantarflexion ($6.3 \pm 8.9^\circ$) through stance, as well as
265 decreased knee flexion through swing ($29.8 \pm 9.1^\circ$). K_5 included patterns with increased hip ($44.7 \pm$
266 12.2°) and knee flexion ($80.0 \pm 10.1^\circ$) during swing and increased hip abduction ($4.5 \pm 3.5^\circ$) in stance.

267 Finally, K_1 included 322 unique gait patterns that aligned closely with baseline trends ($p > 0.054$ for all
268 angles), capturing points within the feedback trials in which participants were minimally exploring.

269 As expected, stride-to-stride variability increased for all kinematic parameters in K_2 to K_5 ($p <$
270 0.05 for all parameters), with the largest variability seen in gait patterns in K_2 . This increase in variability
271 highlights an inherent learning effect associated with unique gait pattern reproduction. Even when
272 participants were minimally exploring (*i.e.*, K_1), there was generally an increase in variability compared
273 to baseline, likely due to the added attentional demand of responding to the biofeedback system.

274 3.2 Synergy Analysis

275 All participants were able to significantly modify synergy complexity during exploration (Figure 4). A
276 one-synergy decomposition ($i = 1$) accounted for $66.1 \pm 5.9\%$ of the variance in the EMG data during
277 baseline. When clustered, tVAF₁ was $71.6 \pm 7.2\%$ (K_1), $78.3 \pm 6.6\%$ (K_2), $76.5 \pm 6.4\%$ (K_3), $76.0 \pm 6.2\%$
278 (K_4), and $69.8 \pm 5.7\%$ (K_5), indicating that all of the explored patterns significantly decreased complexity
279 ($p < 0.05$). Interestingly, there were also significant inter-cluster differences in tVAF₁, suggesting that the
280 type of gait pattern modification impacted complexity ($p << 0.001$). It should be noted that although
281 participants were instructed to either raise or lower their synergy complexity scores, minimal differences
282 existed between these trials; participants generally decreased complexity regardless of the target direction.
283 As such, we did not conduct further analyses comparing target directions.

284 A three-synergy solution ($i = 3$) accounted for $92.6 \pm 2.3\%$ of the variance for all exploration and
285 baseline walking patterns. Clustering yielded four distinct synergy structures (Figure 5) that were
286 dominated by the TA (W_1), hamstrings (W_2), the quadriceps and gluteus maximus (W_3), and the
287 plantarflexors (W_4). All four synergy structures were observed across K_1 to K_5 as well as baseline walking
288 but were recruited with varying frequency. For example, baseline walking was primarily defined by W_2 ,
289 W_3 , and W_4 , which were present in 85.7%, 78.6%, and 100% of gait patterns in the group, respectively.
290 These synergies align with those previously reported in nondisabled adults during steady-state walking

291 (Allen and Neptune, 2012; Clark et al., 2010). In contrast, K_3 was dominated by W_1 (76.5%), W_2
292 (82.7%), and W_4 (91.8%). Interestingly, the plantarflexor synergy (W_4) emerged for the majority of
293 patterns in all clusters (K_1 to K_5) whereas W_1 , W_2 , and W_3 were differentially recruited. These results
294 suggest that a small pool of synergies exists that can be selectively drawn from depending on the
295 biomechanical constraints of a given pattern. Across groups, synergy activation patterns were also distinct
296 from baseline and aligned with key kinematic trends. For example, K_2 was characterized by increased
297 knee flexion and ankle dorsiflexion through the gait cycle, which was reflected in the increased activation
298 of W_1 in swing and W_3 through stance.

299 The observed change in synergies recruited during exploration corresponded to an overall
300 decrease in $tVAF_{3_BASE}$ when baseline synergy weights were used to reconstruct EMG data from
301 exploration trials (Figure 6; $p << 0.001$ for all groups). For the three-synergy solution, baseline synergy
302 weights accounted for $6.0 \pm 6.0\%$ (K_1), $17.7 \pm 11.0\%$ (K_2), $10.6 \pm 8.0\%$ (K_3), $15.3 \pm 8.7\%$ (K_4), and 11.3
303 $\pm 7.1\%$ (K_5) less of the variance in EMG data than weights extracted directly from each unique pattern.
304 Further, reconstruction quality was different between clusters ($p << 0.001$), with the largest change in
305 synergy structure seen in K_2 . This suggests that baseline synergy weights captured muscle activity for
306 certain gait patterns better than others, further confirming the flexible recruitment of synergies to
307 changing biomechanical constraints.

308 3.3 BART Analysis

309 The BART model was able to explain changes in synergy complexity observed during exploration ($R^2 =$
310 0.88 ; RMS error = 4.4). Baseline $tVAF_1$ emerged as the top predictor of $\Delta tVAF_1$ (net effect = 4.6%), as
311 individuals with higher baseline complexity increased $tVAF_1$ to a greater extent during exploration than
312 those with lower baseline complexity (Figure 8). However, this observation partially reflects the effects of
313 regression to the mean. After baseline $tVAF_1$, kinematic and kinetic predictors, especially those at the
314 knee and ankle, had the largest effects on $\Delta tVAF_1$ (Figure 9A). In particular, greater knee flexion (net
315 effect = 3.2%), anterior pelvic tilt (2.3%), hip extension moment (2.7%), and ankle dorsiflexion moment

316 (2.8%) through stance corresponded to a greater decrease in synergy complexity. Increased knee
317 extension moment (net effect = 3.1%) at initial contact also corresponded to less complex control.
318 Interestingly, only one swing-phase variable had a large effect on $\Delta tVAF_1$; decreased knee flexion during
319 swing resulted in greater decreases in synergy complexity (net effect = 3.3%). Further, two measures of
320 kinematic and kinetic variability emerged among the top predictors in the BART model (Figure 9B),
321 highlighting the sensitivity of synergy complexity to the increased stride-to-stride variability observed
322 during gait pattern exploration.

323 Beyond gait mechanics, both participant number (net effect = 2.1%) and speed (1.6%) emerged
324 among the top predictors in the model. Although the former effect was largely driven by one participant
325 (P8), it still indicates that differences may exist in how individuals interacted with the biofeedback
326 system, including both the range of patterns they explored and their comprehension of the presented
327 metric. Further, the moderate effect of speed on synergy complexity, whereby slower speeds were
328 associated with greater decreases in complexity (Figure 8), could suggest differences in the feasibility of
329 performing certain gait patterns at different speeds.

330 4. DISCUSSION

331 This study demonstrated that a small library of synergies was sufficient to characterize a broad repertoire
332 of gait patterns attempted during biofeedback walking, and that recruitment from this library was
333 dependent on both the type and magnitude of gait pattern deviation. Specifically, small deviations from
334 baseline walking were generally accommodated by altering the activations of a consistent set of synergies
335 whereas different synergies were recruited to produce larger gait changes. Participants were also able to
336 widely modulate synergy complexity during gait pattern exploration. However, the majority of gait
337 patterns corresponded to an increase in $tVAF_1$ (i.e., decreased complexity); across all participants, only
338 17.4% of attempted patterns decreased $tVAF_1$. Collectively, these results suggest that although synergy
339 structures appear to be invariant, synergies can be flexibly recruited in response to changing sensory input
340 or biomechanical constraints. This organizational strategy is advantageous for enabling rapid learning of

341 new movement patterns and ensuring successful navigation in complex environments (Torres-Oviedo and
342 Ting 2010, Chiel et al 2009, McKay 2007).

343 Our observation that a small pool of synergies emerged during gait pattern exploration aligns
344 closely with prior literature in both animal and human models. These studies have demonstrated that
345 synergies are consistent across a repertoire of motor outputs (Allen and Neptune, 2012; Barroso et al.,
346 2014; Cappellini et al., 2006; Hagio et al., 2015; Ivanenko et al., 2004) and can be flexibly combined to
347 accommodate changes in sensory input (Cheung et al., 2005; Ivanenko et al., 2004; Kargo et al., 2010;
348 McGowan et al., 2010; Rozumalski et al., 2017) or biomechanical constraints (Krishnamoorthy et al.,
349 2004; Nazifi et al., 2017; Torres-Oviedo and Ting, 2010). However, beyond identifying differences in
350 synergy recruitment across movements, the nature of our protocol enabled us to understand the factors
351 associated with these differences with greater precision. For example, we demonstrated that small
352 deviations at the hip, knee, and ankle, as observed in K_1 , were accommodated by baseline synergies, as
353 baseline synergy weights largely captured the variance in EMG activity during feedback walking (i.e.,
354 tVAF_{3_BASE} for K_1). Baseline synergies were also recruited for the majority of patterns in K_5 , as the large
355 increase in knee flexion through swing could be accommodated by altering the activation timing of the
356 hamstring synergy (W_2). In contrast, patterns which were defined by large deviations in sagittal plane
357 mechanics through stance (e.g., K_2 and K_4), had synergy structures more dissimilar from baseline (Figure
358 6). A similar relationship emerged when considering synergy complexity. The results from our BART
359 analysis demonstrated that deviations at the knee and ankle during stance largely predicted changes in
360 tVAF₁ during gait pattern exploration. Specifically, greater knee extension moment, ankle dorsiflexion
361 moment, and knee flexion through stance corresponded to reduced complexity. This finding aligns with
362 observations in clinical crouch gait in cerebral palsy, where a crouched posture places greater demand on
363 the quadriceps to accelerate the center of mass upward and counteract gravitational force, resulting in
364 increased coactivation of the hamstrings and quadriceps through stance and, therefore, reduced control
365 complexity (Spomer et al., 2022; Steele et al., 2013). Hip extension moment through stance also emerged

366 among the top predictors in the BART model, which further indicates that increased hamstring-quadricep
367 co-contraction had a large effect on $\Delta tVAF_1$.

368 Beyond identifying those variables which were most predictive of changes in synergy
369 recruitment, the results from the BART analysis also allowed us to capture the non-linear relationship
370 between gait pattern deviations and synergy complexity. Specifically, a stepwise relationship consistently
371 emerged for kinematic and kinetic predictor variables wherein $tVAF_1$ was similar to baseline values up
372 until a certain threshold, after which changes in $tVAF_1$ were larger, but generally consistent. The stability
373 of synergy complexity measures for gait patterns similar to baseline walking further confirms the
374 propensity for the CNS to maintain a consistent control strategy to accommodate small gait deviations.
375 Further, the plateau in $\Delta tVAF_1$ observed at the extremes of each gait variable suggest that bounds exist on
376 the extent to which synergy complexity can be modulated, at least when limited to a specific muscle set.

377 While outcomes from the BART analysis also revealed a monotonic relationship between
378 baseline complexity and $\Delta tVAF_1$, partially reflecting regression to the mean, the overwhelming majority
379 of patterns selected during exploration increased $tVAF_1$. Although these results could reflect participant
380 comprehension of the biofeedback system and the task instructions, they may also be indicative of the
381 underlying control strategy employed by the CNS during learning. In novel task execution, the CNS may
382 initially assume a less complex strategy, sacrificing efficiency for stability. This hypothesis is consistent
383 with studies demonstrating that long-term training facilitates more efficient use of neural resources
384 (Krings et al., 2000; Picard et al., 2013) and increased supraspinal excitability (Christiansen et al., 2020;
385 Pascual-Leone et al., 1995; Rosenkranz et al., 2007). For example, Sawers et al (2015) demonstrated that
386 trained dancers recruited a larger number of synergies than novices during both beam and overground
387 walking and that the synergies recruited were sparser, both of which were used to suggest that training
388 promoted greater selective motor control. In our study, because individuals typically explored each
389 unique gait pattern for a short bout (~10 strides) during exploration, the CNS may have had insufficient

390 time to tune its control strategy, contributing to the observation that participants could increase, but not
391 consistently decrease tVAF₁ values.

392 Whether synergy complexity is similarly flexible and can be consistently increased following
393 neurologic injury is largely unknown but is especially salient for informing gait rehabilitation. Individuals
394 with central nervous system damage recruit fewer synergies than nondisabled peers (Cheung et al., 2012;
395 Clark et al., 2010; Fox et al., 2013; Rodriguez et al., 2013; Schwartz et al., 2016; Steele et al., 2015).
396 Further, within these populations, both synergy complexity and structure have been associated with
397 impairment level, as those with more severe impairments have less complex control (Cheung et al., 2012;
398 Steele et al., 2015). This is hypothesized to reflect increased reliance on spinal circuitry over supraspinal
399 input to shape motor outputs following neurologic injury, which may reduce the overall flexibility of
400 synergy recruitment (Leonard et al., 1991). This relationship has been demonstrated in CP, where prior
401 literature has reported that synergies are unchanged following surgery and biofeedback training, despite
402 both interventions yielding measurable improvements in gait (Booth et al., 2019; Shuman et al., 2019).
403 Further, stroke survivors with less severe impairment appear to maintain the capacity to modulate
404 synergies during locomotor training better than those with more severe impairment (Rouston et al., 2013).
405 Understanding whether individuals with neurologic injury can consistently alter synergy complexity and
406 improve movement, or how interventions can support sustained changes in control remain active and
407 important areas for future investigations. While recent literature has indicated that providing richer
408 afferent information via spinal stimulation or sensorimotor biofeedback may promote greater supraspinal
409 involvement and, therefore, more flexible synergy recruitment during movement, studies are still ongoing
410 (Cheng et al., 2019; Conner et al., 2021; Gad et al., 2021).

411 Our observation that participant number was predictive of $\Delta tVAF_1$ further accentuates the need to
412 evaluate personalized responses to biofeedback. This result suggests that even when controlling for all
413 other model covariates, including baseline complexity, interparticipant differences in response persisted.
414 Heterogeneous response to biofeedback training has been cited previously and may stem from both

415 individual capacity to modify the parameter targeted by biofeedback as well as system design choices
416 (Booth et al., 2019; Huang et al., 2006; MacIntosh et al., 2019; Sigrist et al., 2013; Spencer et al., 2021;
417 van Gelder et al., 2017). The latter likely contributed to the results observed here. Because synergy
418 complexity is derived from multiple data streams, some participants reported feeling unsure about how
419 specific gait changes affected the displayed metric or struggled to conceptualize what ‘more’ or ‘less’
420 complex gait patterns entailed, both of which likely influenced their exploration strategy. These results
421 highlight an inherent challenge of using motor control-based biofeedback in gait training applications and
422 presents an opportunity to explore more interpretable biofeedback metrics that can still be used to
423 improve control patterns. For example, the output from our model suggests that providing information on
424 joint moments to reduce hamstring-quadricep co-contraction in early stance may elicit changes in synergy
425 complexity, although further work is needed to extend these findings to populations with neurologic
426 injury. Our results also demonstrate the unique advantage of using non-linear function estimation
427 techniques such as BART in order to better interpret the inherently complex and multifactorial user-
428 system interactions present during biofeedback training to inform future system design.

429 4.1 Methodological Considerations

430 Although the decision to use a biofeedback system and minimal researcher coaching allowed us to
431 capture a broader array of patterns than have been previously examined in studies of synergies in gait,
432 there are limitations to this approach that should be considered when interpreting the results. Because we
433 wanted participants to freely explore using the biofeedback system, we only required them to take five
434 strides in a selected gait pattern. This meant that the novelty of the attempted patterns was likely reflected
435 in our results, as previously described. In order to reduce this effect, we calculated synergies from the
436 same number of strides during exploration and baseline walking ($n = 5$); however, it is possible that
437 synergies may have adapted further if we had collected a larger number of strides for each unique pattern
438 (Oliveira et al., 2014). The unstructured nature of the protocol also introduced the likelihood of observing
439 extreme outliers, as a given gait pattern may only be attempted by a single participant. The opportunity

440 for outliers and observed heterogeneity of participant response informed our decision to use BART as a
441 modeling paradigm. Because BART natively constrains tree structure, it prevents data overfitting, thereby
442 reducing the likelihood that outliers in our data set could significantly affect model outputs (Chipman et
443 al., 2010). Finally, despite the diversity of patterns attempted, our analysis was still limited to a subset of
444 gait patterns making it challenging to draw definitive conclusions about the relationship between
445 biomechanical constraints and synergies. In future studies, biofeedback systems may be useful to guide
446 users through a sample of possible walking configurations in order to develop a more comprehensive
447 landscape of user response. Simulation paradigms, such as those employed by Kutch and Valero-Cuevas
448 (2012), which involve systematically iterating over a range of achievable outputs, could also be a valuable
449 compliment to the present study to provide further insight into how synergies change as a function of gait
450 exploration. Importantly, such analyses need to be performed in both nondisabled populations and those
451 with neurologic injury in order to understand how injury impacts one's ability to flexibly alter control
452 strategies during walking.

453 5. CONCLUSION

454 Using motor control-based biofeedback to encourage exploration and capitalizing on non-linear machine
455 learning methodology allowed us to identify salient features which influence how the CNS flexibly
456 shapes control during walking. This analysis revealed that a small library of spatially invariant synergies
457 can be flexibly recruited to produce a diverse array of motor outputs and that recruitment changes as a
458 function of the imposed biomechanical constraints. Specifically, our results suggest that large deviations
459 in distal joint mechanics during stance resulted in the greatest overall change in synergy recruitment from
460 baseline walking. Further, they indicate that other participant-level factors may affect one's ability to
461 modify synergy recruitment during walking, which must be considered when designing interventions to
462 this end. Whether the recruitment flexibility observed in this study is a luxury of the unimpaired
463 neurological system or is maintained following neurological injury is a critical next step of this work. By
464 modeling how synergies are modulated during locomotion, we believe that this study presents both

465 theoretical and methodological contributions towards bolstering understanding of the neural control of
466 movement and may aid in improving interventions for individuals with neurological injury.

467 **Conflict of Interest**

468 The authors declare no conflict of interest regarding the publication of this manuscript.

469

470 **Author Contributions**

471 A.M.S., conceived and designed this research, collected and analyzed data, prepared figures, and drafted
472 the manuscript. R.Z.Y collected and analyzed data and revised the manuscript. M.H.S. and K.M.S.
473 conceived and designed the research, interpreted results, and revised the manuscript. All authors approved
474 the final version of the manuscript of publication.

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481

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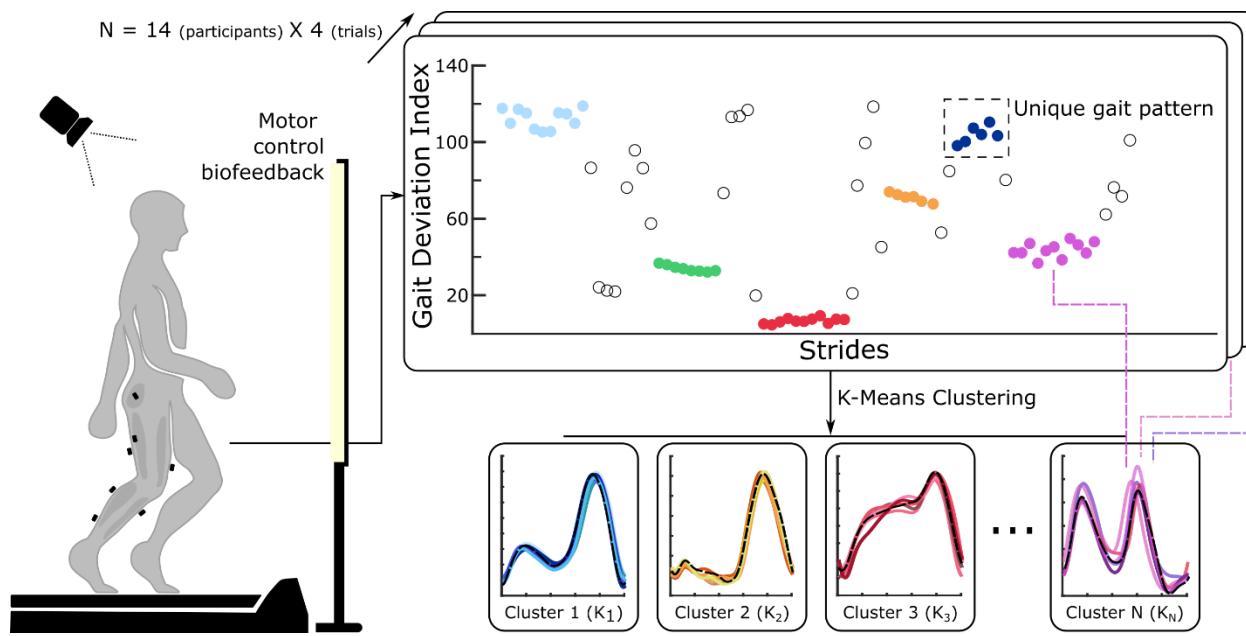
688 *Table 1: BART Model Variables*

Response	Description
$\Delta tVAF_1$	Difference in motor control complexity between each unique gait pattern and baseline walking.
Predictors	Description
Baseline tVAF₁	Measure of motor control complexity during the baseline walking condition. Values range from 0 - 1, where a higher value indicates less complex control.
Participant Number	Values range from 1-14. This variable was used to evaluate if participant-level differences in biofeedback exploration emerged.
Speed	Nondimensional walking speed normalized to participant leg length.
Kinematics/Kinetics	Mean values of all variables outlined in Figure 2. Variables input as z-scores, normalized to baseline walking. These variables were selected as they sufficiently capture kinematic/kinetic trends during the gait cycle.
Kinematic/Kinetic Variability	Difference in the standard deviation of all variables outlined in Figure 2 between each unique gait pattern and baseline walking. These variables reflect the participant's ability to consistently produce the attempted gait patterns.

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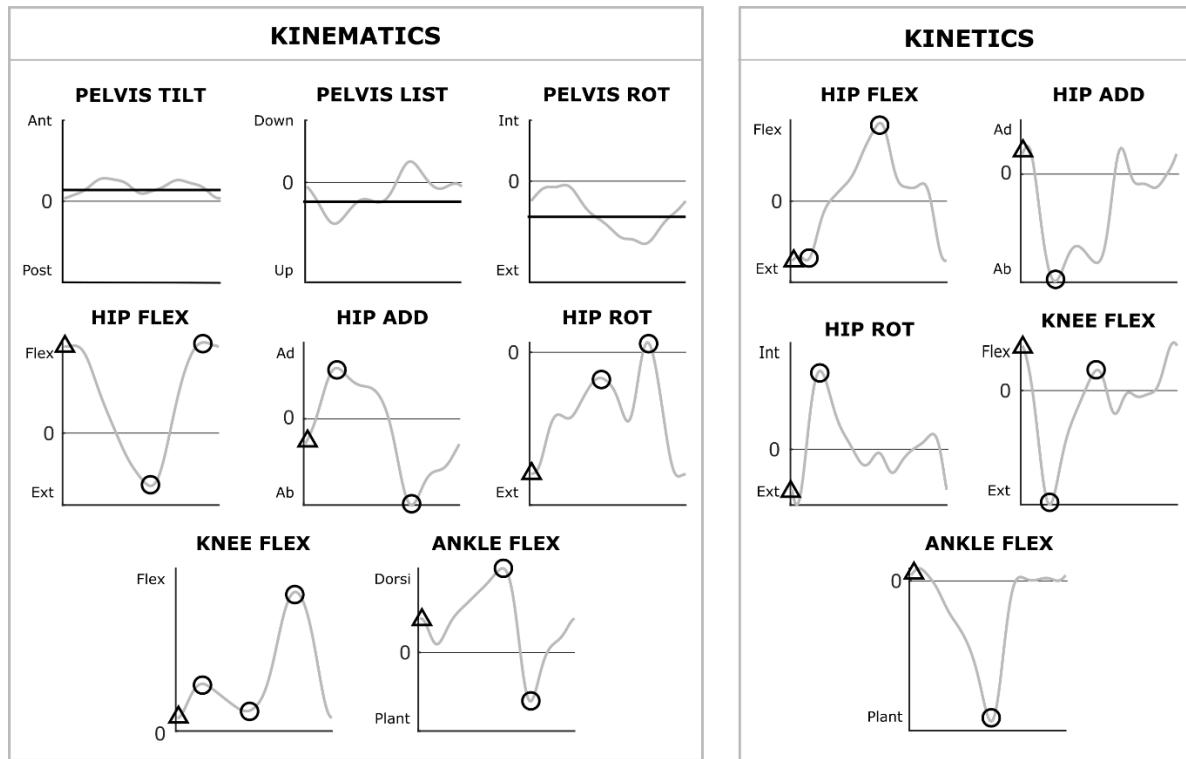
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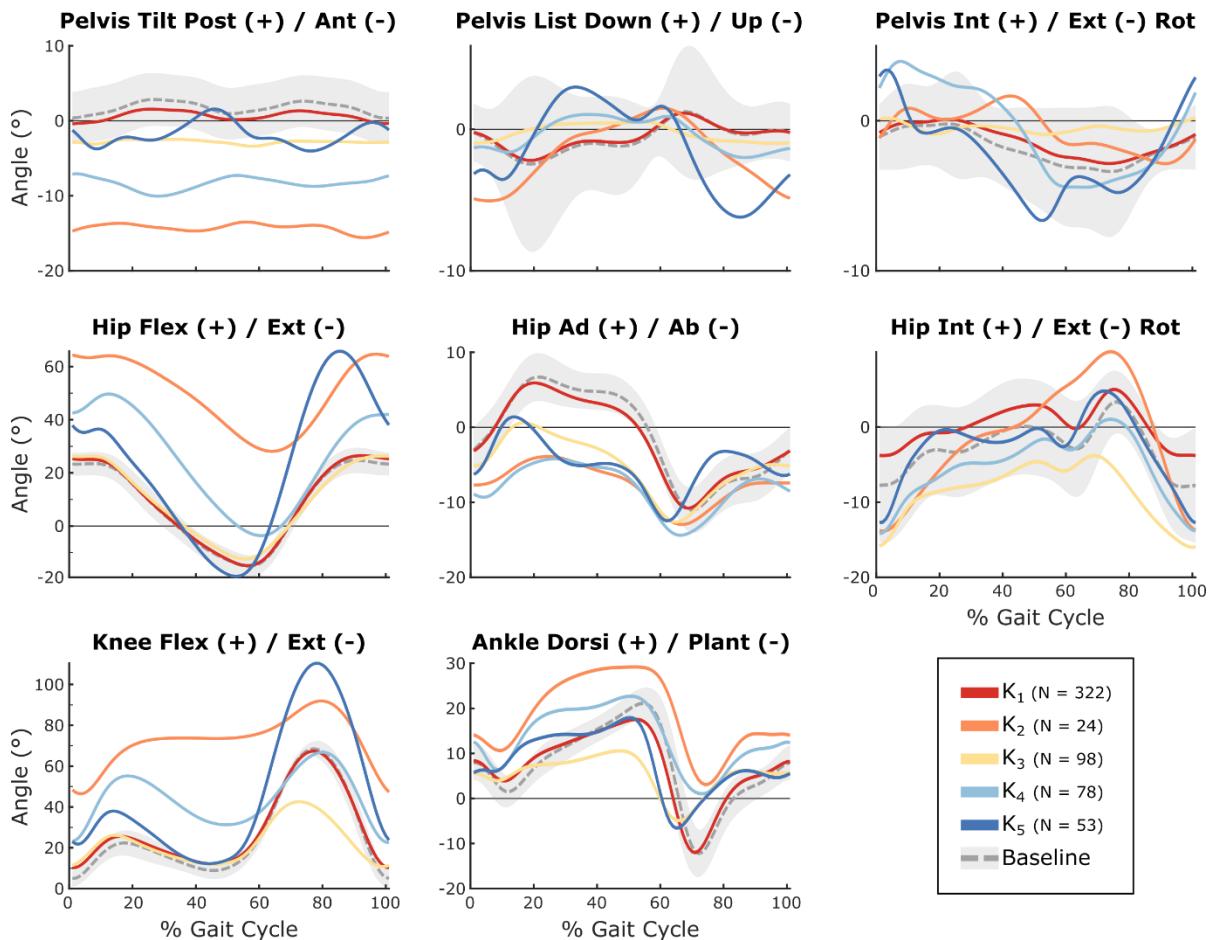
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693 *Figure 1: Methodology used to identify clusters (K₁ to K_N), representing kinematically-similar gait*
694 *patterns attempted by participants during feedback walking. Full-body kinematic and kinetic data and*
695 *lower-limb EMG data were collected while participants explored a broad range of movement patterns*
696 *using biofeedback. The Gait Deviation Index (GDI) was calculated from kinematic data for each*
697 *recorded stride for every participant and trial (56 data sets). Unique gait patterns were labeled as five or*
698 *more consecutive strides with similar GDI values and manually confirmed. Kinematic data for these*
699 *unique patterns were input into a k-means clustering algorithm to identify clusters (K₁ to K_N) across*
700 *participants and trials.*



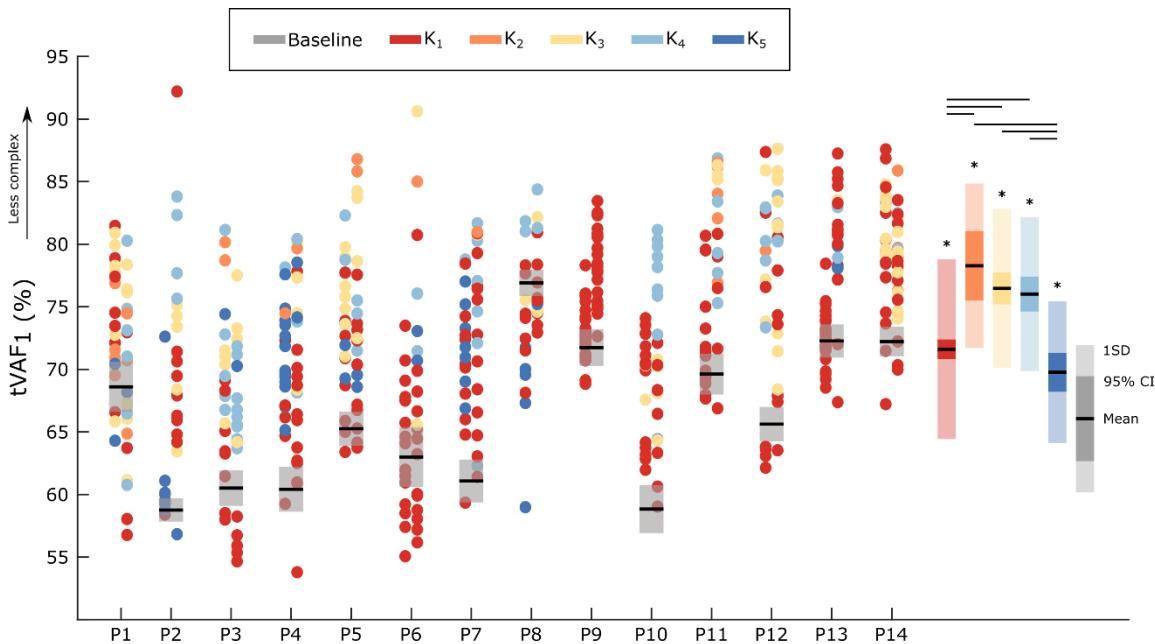
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Figure 2: Kinematic and kinetic predictor variables included in the BART model. Each icon indicates a variable ($n = 31$ total) that was identified for every unique gait pattern. Variables were selected to capture trends at the pelvis, hip, knee, and ankle that could change during exploration. Circles indicate local maximum or minimum values, and triangles indicate initial contact points, calculated as the mean value over the first 5% of the gait cycle. Average pelvis list, tilt, and rotation angles across the gait cycle were included to capture asymmetries. The standard deviations of each kinematic and kinetic variable, used to capture stride-to-stride variability during gait pattern exploration, were also included in the predictor set.



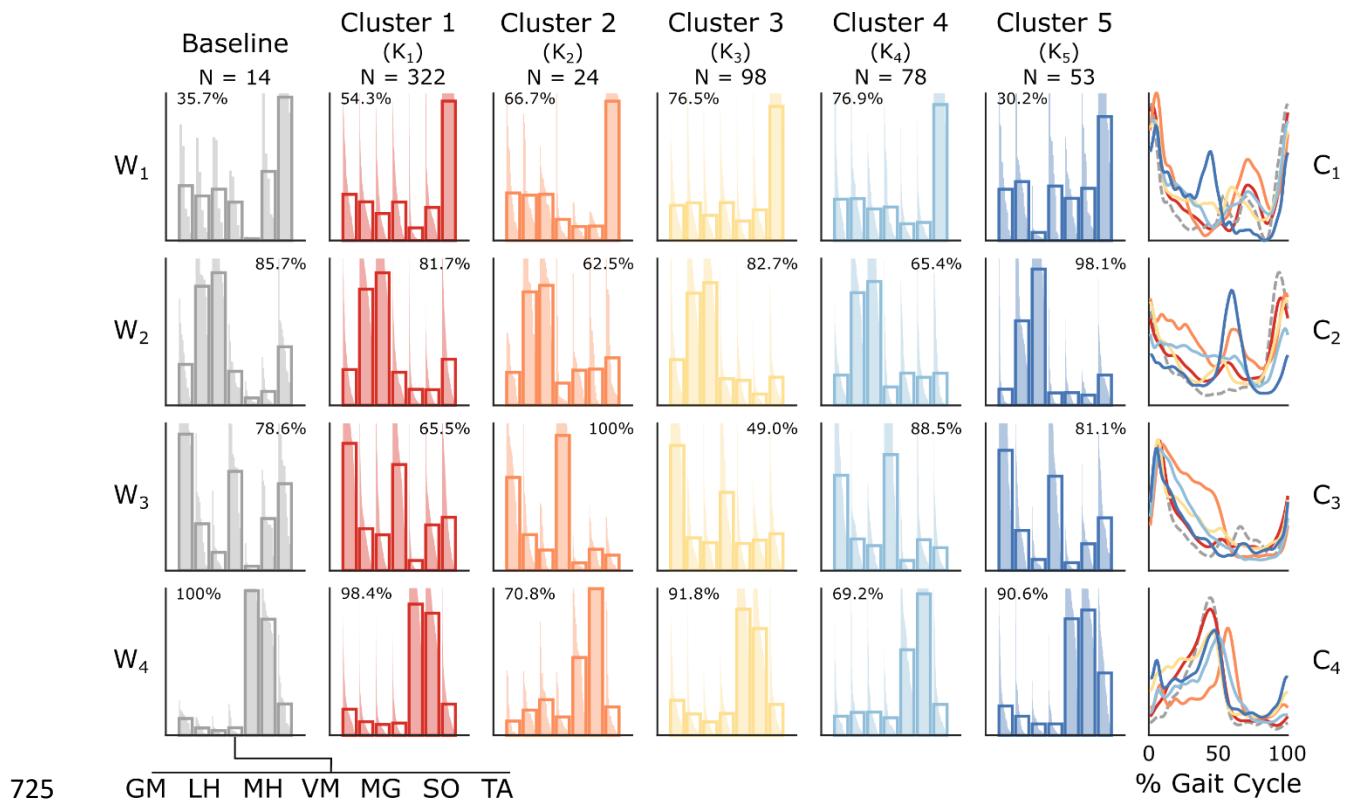
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711 *Figure 3: Average pelvis, hip, knee, and ankle kinematics for the five clusters identified by k-means*
712 *clustering ($K_1 - K_5$), representing common gait patterns attempted during exploration. The baseline*
713 *condition shows \pm ISD.*



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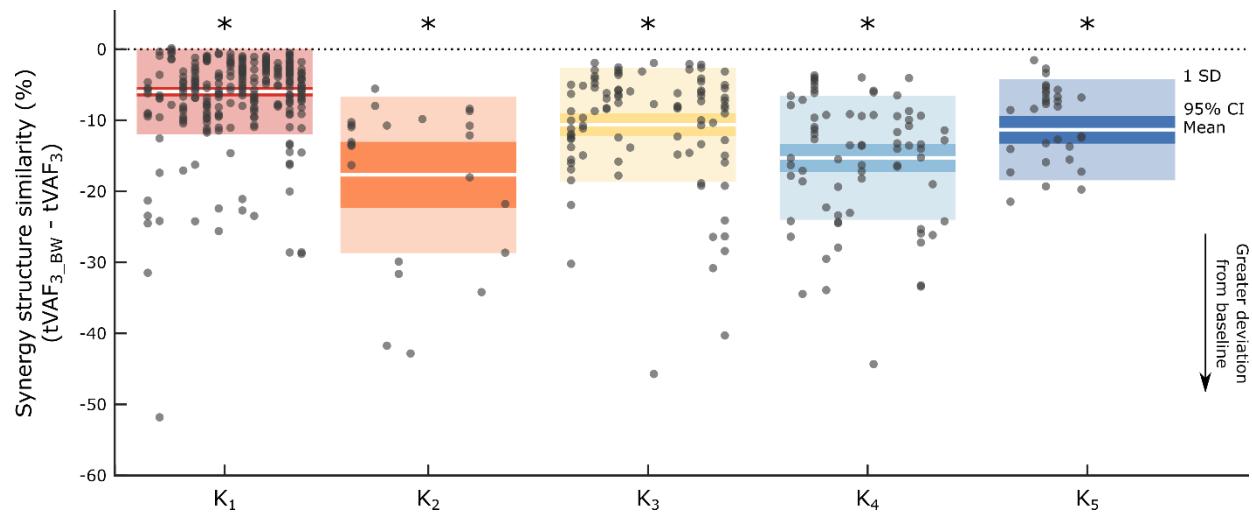
715 *Figure 4: The total variance accounted for by a one-synergy solution ($tVAF_1$) for each participant (P1-
716 P14). Each dot represents a unique gait pattern and is colored according to the cluster it was sorted into
717 (K₁ to K₅). For each participant, data is organized into two columns representing the feedback trials in
718 which participants were instructed to decrease (left) and increase (right) their $tVAF_1$. Baseline data is
719 presented as a mean \pm 1SD, representing the distribution of $tVAF_1$ values resulting from bootstrapping
720 using sets of five random strides (replicates = 200). Boxes represent the mean (black line), 95%
721 confidence interval (solid color), and standard deviation (shading) of $tVAF_1$ values for each cluster and
722 the baseline condition. Larger $tVAF_1$ values correspond to decreased motor control complexity. * denotes
723 significant difference between each group and the baseline condition and black bars indicate significant
724 inter-cluster differences ($\alpha = 0.05$).*



725 *Figure 5: Average synergy weights (W) and activations (C) for the three-synergy solution for baseline*
726 *walking and each cluster of kinematically-similar gait patterns (K_1 to K_5). K-means clustering was*
727 *performed for the $i = 3$ synergy solution across all unique gait patterns and yielded four unique*
728 *structures. Synergy weight plots reflect cluster-wise averages as well as weights for individual gait*
729 *patterns, sorted in descending order. Percentages reflect the number of gait patterns within each cluster*
730 *(i.e., K_1 to K_5) that used each synergy. Muscles: gluteus maximus (GM), lateral hamstring (LH), medial*
731 *hamstring (MH), vastus medialis (VM), medial gastrocnemius (MG), soleus (SO), and tibialis anterior*
732 *(TA).*

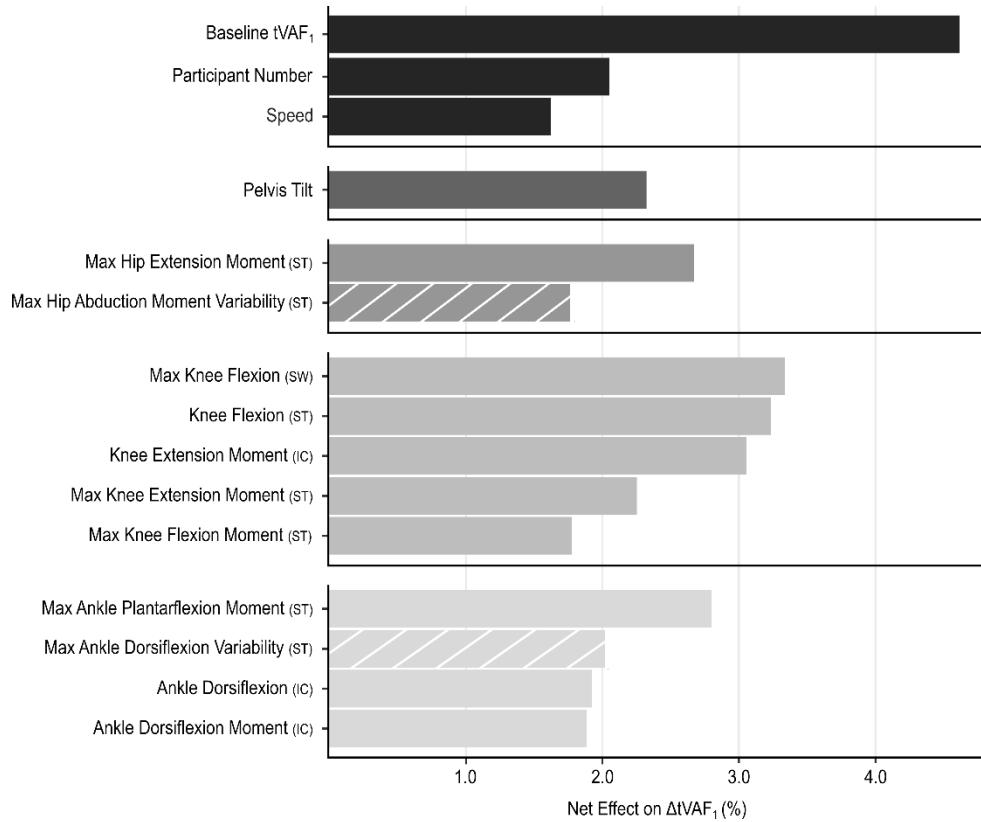
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737 *Figure 6: Similarity of the $i = 3$ synergy solution for each cluster (K_1 to K_5) as compared to baseline*
738 *walking. Baseline synergy weights were used to reconstruct the EMG data for all unique gait patterns*
739 *within each cluster. $tVAF_{3_BASE}$ represents the amount of variance accounted for by baseline walking*
740 *synergy weights and $tVAF_3$ represents the variance accounted for by weights extracted directly from*
741 *EMG data for each unique gait pattern. For each box, the white bars represent mean values, the solid-*
742 *colored blocks represent a 95% confidence interval, and shading shows ± 1 SD. Dots represent unique*
743 *gait patterns and are arranged in columns to represent individual participants (P1 to P14). Large*
744 *differences between $tVAF_{3_BASE}$ and $tVAF_3$ indicate that synergies during gait pattern exploration deviate*
745 *more from baseline walking. * denotes significant difference between each group and zero, indicating a*
746 *change in synergies from baseline walking.*

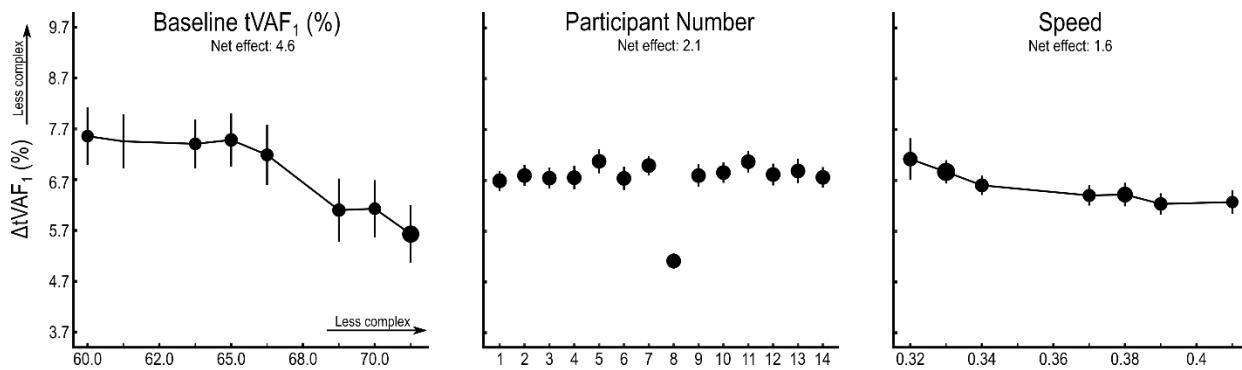


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748 *Figure 7: Net effects for the top fifteen predictors on $\Delta tVAF_1$ from the BART model. Net effects were*
749 *derived from the generated ALE plots and defined as the difference between the 95th and 5th percentile of*
750 *the response variable over the range of each predictor, when controlling for all other model covariates.*
751 *Cross-hatching indicates measures of stride-to-stride variability. Gait phases: Initial contact (IC), stance*
752 *(ST), and swing (SW). See Table 1 for all variables included in the BART model.*

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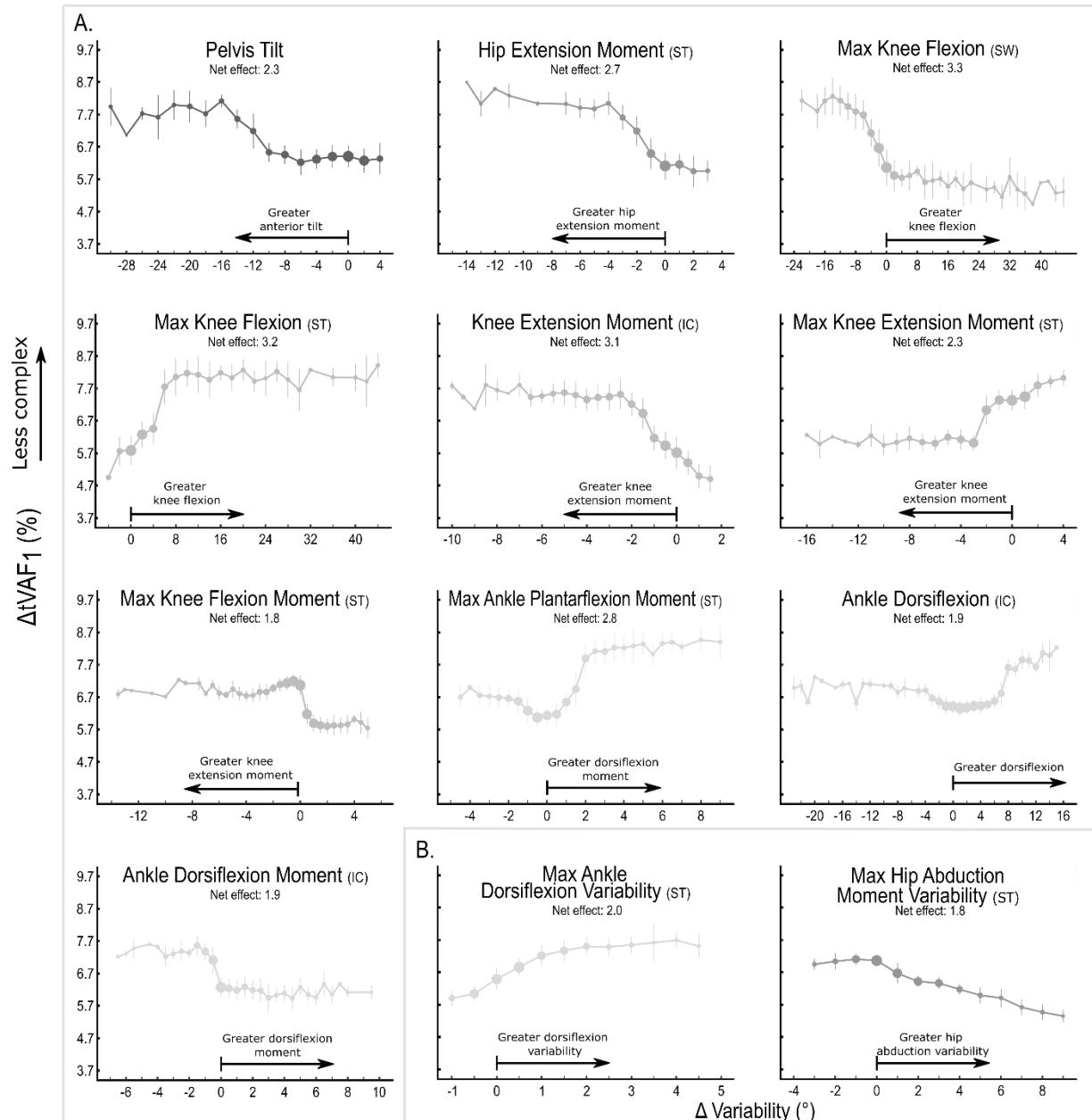
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756 *Figure 8: Accumulated local effect (ALE) plots for baseline synergy complexity, participant number, and*
757 *speed. Speed is normalized to participant leg length. Each plot depicts the effect of an individual*
758 *predictor on changes in synergy complexity from baseline, conditioned on all other predictors in the*
759 *model. Predictor data is separated into evenly spaced bins and the size of individual points represents the*
760 *number of samples in each bin. Larger values for tVAF₁ indicate less complex control. Net effects were*
761 *calculated as the difference between the 95th and 5th percentile of $\Delta tVAF_1$ over the range of each*
762 *predictor.*

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765 *Figure 9: Accumulated local effects of kinematic, kinetic, and variability measures on $\Delta tVAF_1$. Kinematic*
 766 *and kinetic measures (A) are presented as z-scores normalized to baseline walking, such that the x-axis*
 767 *depicts standard deviations away from baseline. Variability measures (B) are presented as degrees away*
 768 *from baseline variability. Vertical bars indicate ± 1 SD. Plots are cropped to display the middle 95% of*
 769 *the predictor data to remove extreme outliers. Predictor data was separated into evenly spaced bins and*
 770 *the size of individual points represent the number of samples in each bin. Gait phases: Initial contact*
 771 *(IC), stance (ST), and swing (SW).*

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