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Encoding of both reaching and grasping kinematics in dorsal and ventral premotor cortices

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33 **Abstract**

34 Classically, it has been hypothesized that reach-to-grasp movements arise from
35 two discrete parieto-frontal cortical networks. As part of these networks, the dorsal
36 premotor cortex has been implicated in the control of reaching movements of the arm,
37 while the ventral premotor cortex has been associated with the control of grasping
38 movements of the hand. Recent studies have shown that such a strict delineation of
39 function along anatomical boundaries is unlikely partly because reaching to different
40 locations can alter distal hand kinematics and grasping different objects can affect
41 kinematics of the proximal arm; Here, we used chronically implanted multi-electrode
42 arrays to record unit spiking activity in both PMd and PMv simultaneously while rhesus
43 macaques engaged in a reach-to-grasp task. Generalized linear models were used to
44 predict the spiking activity of cells in both areas as a function of different kinematic
45 parameters, as well as spike history. To account for the influence of reaching on hand
46 kinematics and vice-versa, we applied demixed principal component analysis to define
47 kinematics synergies that maximized variance across either different object locations or
48 grip types. We found that single cells in both PMd and PMv encode the kinematics of
49 both reaching and grasping synergies suggesting that this classical division of reach
50 and grasp in PMd and PMv, respectively, does not accurately reflect the encoding
51 preferences of cells in those areas.

52 **Significance statement**

53 For reach-to-grasp movements, the dorsal premotor cortex (PMd) has been implicated
54 in the control of reaching movements of the arm, while the ventral premotor cortex
55 (PMv) has been associated with the control of grasping movements of the hand. We
56 recorded unit spiking activity in PMd and PMv simultaneously while macaques
57 performed a reach-to-grasp task. We modeled the spiking activity of neurons as a
58 function of kinematic parameters and spike history. We applied demixed principal
59 component analysis to define kinematics synergies. We found that single units in both
60 PMd and PMv encode the kinematics of both reaching and grasping synergies
61 suggesting that the division of reach and grasp in PMd and PMv, respectively, can't be
62 made based on their encoding properties.

63
64 **Introduction**

65 Reaching to grasp is a fundamental, ethologically relevant primate behavior
66 (Kaas et al., 2013). Several cortical and sub-cortical structures are involved in the
67 generation of these movements (Fattori et al., 2004; Grafton, 2010; Kaas et al., 2013).
68 Recently, much emphasis has been placed on understanding the behavior of single
69 cells in primary motor cortex (MI), one of the main sources of cortical output to the
70 spinal cord, during reach to grasp movements (Saleh et al., 2010, 2012, Mollazadeh et
71 al., 2011, 2014). The motor cortex, however, does not exhibit a clear hierarchical
72 organization with MI as the sole source of corticospinal output (He et al., 1993; Dum
73 and Strick, 2005). Rather, MI is part of a network of cortical areas involved in the
74 production of reach to grasp movements. Two other cortical areas that are densely

connected horizontally with MI are dorsal and ventral premotor cortex (PMd and PMv, respectively) with these two areas providing a large proportion of cortical inputs to MI (Dum and Strick, 2005; Dea et al., 2016).

The dual channels hypothesis has posited that reach to grasp movements arise from the temporal coordination of activity in independent reaching and grasping brain networks (Jeannerod, 1988; Karl and Whishaw, 2013). In this framework, PMd, which makes anatomical connections to superior parietal lobe, specifically area V6A (Fattori et al., 2010, 2012), has been considered part of a dorsomedial parietofrontal reaching network involving movements of the arm, while PMv, which makes anatomical connections to inferior parietal lobe, specifically AIP and VIP (Luppino et al., 1999; Borra et al., 2008), has been associated with a dorsolateral parietofrontal grasping network and concerned with distal aspects of the limb, namely, the hand (Dum and Strick, 2005; Hoshi and Tanji, 2007; Grafton, 2010; Dea et al., 2016). However, PM single unit spiking activities have been shown to encode both grip type and target position/reach direction during reach-to-grasp movements in both PMd and PMv (Lehmann and Scherberger, 2013; Stark et al., 2007). Those studies considered grip type and target position/reach direction as categorical variables. What is unknown is whether the encoding of kinematic trajectories of reaching and grasping is spatially segregated across the premotor cortex.

Here, we directly tested whether representations of reaching and grasping kinematics were strictly segregated in PMd and PMv. To address this, however, we had to account for the fact that reaching for objects in different locations influenced not only proximal limb kinematics of the arm but also distal kinematics of the hand. Likewise,

98 grasping different objects affected both hand and arm kinematics. Thus, we applied a
 99 novel dimensionality reduction technique to dissociate anatomical representations (i.e.
 100 based on encoding specific joints) from functional representations (i.e. encoding reach-
 101 related or grasp-related kinematic synergies). We found no evidence of a strict
 102 segregation of reaching and grasping activity in PMd or PMv. Instead, we observed that
 103 both areas contained complete anatomical representations of the upper limb and
 104 functional representations of both reaching and grasping.

105 **Methods**

106 *Neurophysiology*

107 All surgical and experimental procedures were approved by the University of
 108 Chicago Animal Care and Use Committee and conformed to the principles outlined in
 109 the Guide for the Care and Use of Laboratory Animals (NIH publication no 86-23
 110 revised 1985). Two rhesus macaques (*macaca mulatta*) were implanted with 96-
 111 electrode Utah arrays in the dorsal and ventral premotor cortex contralateral to their
 112 working arm. Electrodes were 1.5 mm in length except for the array in PMd of animal J
 113 (1.0 mm). Neural spiking activity from the electrodes were amplified with a gain of
 114 5000, band-pass filtered between 0.3 Hz and 7.5 kHz, and recorded digitally (14 bit
 115 resolution) using a Cerebus acquisition system (Blackrock Microsystems, Salt Lake
 116 City, UT). A threshold was set above the noise floor on each channel (5.5 standard
 117 deviations) and every time this threshold was crossed a 1.6 ms sample (sampled at 30
 118 kHz) was recorded as a putative spike waveform. These threshold crossings were
 119 subsequently sorted offline using a semi-manual clustering procedure (Offline Sorter,
 120 Plexon Inc., Dallas, TX, RRID:SCR_000012). Only sorted waveforms with a signal-to-

121 noise ratio (SNR) greater than 3 were used in subsequent analyses. Here, signal to
 122 noise ratio is defined as the peak minus the trough of the average waveform divided by
 123 the average standard deviation of the waveform across time.

124 *Behavioral Task*

125 Two male rhesus macaques were trained to perform a reach to grasp task with
 126 their left hand while their heads were fixed. A robot (RV-1A-S11 6-axis robot;
 127 rixan.com) presented each animal with objects (Fig. 1A) to grasp at four different spatial
 128 locations (Fig. 1B) in its peripersonal space. The set of objects consisted of geometric
 129 shapes that were designed to evoke a variety of different hand conformations when
 130 grasped. Object and location pairings were randomly varied on a trial-by-trial basis to
 131 minimize the amount of motor planning. Additionally, the animal's vision was occluded
 132 between trials with a pneumatically controlled screen. Once the vision screen was
 133 opened, the animal made a reach to grasp movement to the target and held his grip for
 134 at least 500 ms in order to receive a juice reward. During each movement, the animal
 135 was able to see his arm and hand. Each object was attached to a shaft whose ending
 136 was attached to a cube with a magnet that allowed the experimenter to quickly affix a
 137 different object to the end-effector of the robot on each trial.

138 The precise set of objects that were presented to animal L consisted of 5
 139 different objects (Cylinder, Small Disc, Key, Large Disc, Ring) in different orientations
 140 for a total of 11 different grasp conformations, or grips, (Cylinder Horizontal, Cylinder
 141 Out, Cylinder Vertical, Small Disc Horizontal, Small Disc Out, Small Disc Vertical, Key,
 142 Large Disc Horizontal, Large Disc Vertical, Ring Horizontal, Ring Vertical). On average,
 143 the animal completed 21 repetitions of each grip/location pairing, although some

144 combinations were less frequent than others due to the random sampling of objects,
 145 with the minimum number of repetitions being 5. Animal J was presented with three
 146 objects (Cylinder, Small Disc, Ring) to evoke four different grips (Cylinder Out, Small
 147 Disc Out, Ring Horizontal, Ring Vertical). On average, animal J completed 36
 148 repetitions of each grip/location pairing, and all combinations were sampled at least 15
 149 times. Data from each monkey was recorded from one session each.

150 *Motion capture and inverse kinematics*

151 A 10-camera motion capture system (Vicon Motion Tracking System, Oxford,
 152 UK) detected and recorded the 3-D position of retro reflective markers glued to the
 153 animal. A total of 30 markers were placed on the animal's dorsal hand and dorsolateral
 154 arm enabling the tracking of 21 degrees of freedom in the arm and hand (listed below).
 155 The motion capture system was also used to monitor the position of the object and
 156 vision block screen. The time series of 3D marker positions were lowpass filtered bi-
 157 directionally (4th order Butterworth filter 15 Hz cutoff). All filtering and calculations are
 158 done in Matlab (Mathworks, Natick, MA, RRID:SCR_001622) unless otherwise noted.
 159 To measure grasp aperture, we computed the Euclidean distance between markers
 160 attached to the distal most aspects of the thumb, and index finger (D1 and D2,
 161 respectively).

162 To compute wrist speed, we numerically differentiated the positional data of three
 163 markers on the animal's wrist. After numerical differentiation, wrist velocities were again
 164 lowpass filtered (using the same filter design as before). Wrist speed was defined as
 165 the average tangential velocity of the three wrist markers. Movement onset, derived
 166 from wrist speed, was defined as the first moment that wrist speed exceeded 10% of the

167 maximum wrist speed. Reaction times were subsequently defined as the difference in
168 time between movement onset and release of the vision block.

169 We used open-source software (Delp et al., 2007, Opensim, RRID:SCR_002683)
170 to infer joint angles from marker positional data. We estimated joint angles for 21
171 degrees of freedom in the arm and hand including: humerus flexion/extension, humerus
172 abduction/adduction, humerus rotation, elbow flexion/extension, wrist
173 pronation/supination, wrist abduction/adduction, wrist flexion/extension, 1 carpal-
174 metacarpal (CMC) flexion/extension, 1 CMC abduction/adduction, 1 metacarpal-
175 phalangeal (MCP) flexion/extension, 1 MCP abduction/adduction, 2 MCP
176 abduction/adduction, 2 MCP flexion/extension, 2 proximal interphalangeal (PIP)
177 flexion/extension, 3 MCP flexion/extension, 3 MCP abduction/adduction, 4 MCP
178 flexion/extension, 4 MCP abduction/adduction, 4 PIP flexion/extension, 5 MCP
179 flexion/extension, 5 MCP abduction/adduction, and 5 CMC flexion/extension,. Inverse
180 kinematics was bi-directionally, lowpass filtered with a 6 Hz cutoff.

181 *Demixed principal components analysis*

182 We used a novel dimensionality reduction technique, demixed principal
183 component analysis (dPCA), to identify kinematic synergies associated with reaching,
184 grasping, the interaction of reach and grasp, or synergies that were common across all
185 experimental conditions (Brendel et al., 2011; Kobak et al., 2016). This analysis was
186 motivated by two observations about the kinematics. First, on a given trial, there are
187 often strong correlations between the kinematics of different joints suggesting that most
188 of the variance in the kinematics may be described by a few kinematic synergies, often
189 estimated using principal component analysis (Santello and Soechting, 1998; Mason et

190 al., 2001). Second, the correlation patterns across joints during reach-to-grasp
 191 movements depend upon the exact nature of the movement; thus, different
 192 experimental conditions may produce different correlation patterns (Todorov and
 193 Ghahramani, 2004). Briefly, the dPCA algorithm attempts to find linear combinations of
 194 kinematic variables that explain the most variance, yet also vary only with a given
 195 experimental condition.

196 We present a mathematical explanation of dPCA that closely mirrors the
 197 description in (Kobak et al., 2016). Suppose that every trial, e (of E total trials), has an
 198 associated location, l (out of L possible locations), and grip o (out of O possible grips).
 199 Also, suppose that on every trial we have recorded the activity of J kinematic features at
 200 T different time points. We computed the trial-average kinematic trajectory of every
 201 kinematic feature, j , for each grip and location pairing denoted $R_{jlo}(t)$. We then gathered
 202 all the trial-averaged kinematic trajectories into a single matrix, \mathbf{X} of size $J \times LOT$. We
 203 subsequently demeaned \mathbf{X} such that the average of any kinematic feature over all
 204 locations, grips, and times was 0. It has been previously shown that for such a matrix,
 205 \mathbf{X} , it can be decomposed into independent parts called marginalizations that satisfy:

$$\mathbf{X} = \mathbf{X}_t + \mathbf{X}_{tl} + \mathbf{X}_{to} + \mathbf{X}_{lto},$$

206 where \mathbf{X}_t denotes the time varying, but location and grip-invariant part of \mathbf{X} , obtained by
 207 averaging \mathbf{X} over all locations and grips, i.e.,

$$\mathbf{X}_t = \frac{1}{LO} \sum_{l=1}^L \sum_{o=1}^O \mathbf{X}_{tlo}.$$

208 The location dependent term, \mathbf{X}_{tl} , is obtained by averaging $\mathbf{X} - \mathbf{X}_t$ over all grips, and
 209 similarly, the grip dependent term, \mathbf{X}_{to} is obtained by averaging $\mathbf{X} - \mathbf{X}_t$ over all locations.
 210 Finally, the location-grip interaction, \mathbf{X}_{lto} is obtained by computing $\mathbf{X}_{lto} = \mathbf{X} - \mathbf{X}_t - \mathbf{X}_{tl} - \mathbf{X}_{to}$.

211 It is important to note that every marginalization of \mathbf{X} has the same dimension as \mathbf{X} , i.e.

212 $J \times LOT$.

213 Using the decomposition of \mathbf{X} into its marginalizations, dPCA aims to find
 214 directions in \mathbf{R}^J that explain as much variance as possible, with the additional constraint
 215 that this variance should come from only one marginalization. Unlike standard PCA,
 216 dPCA relaxes the assumption that all the directions must be strictly orthogonal.
 217 Formulated as a constrained optimization problem, dPCA seeks to find a direction ϕ in
 218 \mathbf{R}^J , to minimize the following objective:

$$L = \sum_{\phi} \|\mathbf{X}_{\phi} - F_{\phi} D_{\phi} \mathbf{X}\|_F^2,$$

219 where F_{ϕ} is an encoder matrix with q columns, D_{ϕ} is a decoder matrix with q rows, and
 220 $\|\cdot\|_F^2$ denotes the Frobenius norm of a matrix. An efficient algorithm was recently
 221 proposed to solve this objective function (Kobak et al., 2016).

222 In order to compute the cumulative variance explained by the first p dPCs, we
 223 cannot simply add variances because the dPCs are not strictly orthogonal. Instead, the
 224 cumulative variance explained by the first p dPCs is given by:

$$\frac{\|\mathbf{X}\|_F^2 - \|\mathbf{X} - [f_{1:p}][d_{1:p}]' \mathbf{X}\|_F^2}{\|\mathbf{X}\|_F^2},$$

225 where $[f_{1:p}]$ is a matrix of the first p encoding vectors, $[d_{1:p}]$ is a matrix of the first p
 226 decoding vectors, and $[\cdot]'$ is a transpose of a matrix. We emphasize that the
 227 cumulative proportion of variance explained by each component is based off of the trial
 228 averaged data in the matrix, \mathbf{X} .

229 *Generalized linear model analysis*

230 *Input features*

231 We attempted to predict the instantaneous probability of spiking in a given single unit
 232 from many different combinations of extrinsic covariates (i.e. kinematics and
 233 experimental factors) and intrinsic covariates (i.e. neural signals).

234 *Extrinsic covariates*

235 We used the position and velocity of 21 joints in the arm and hand at various time lags
 236 as input features to our encoding model. We used 8 different time lags from -156 ms
 237 (i.e. spikes lag kinematics by 156 ms) to 208 ms (i.e. spikes lead kinematics by 208 ms)
 238 in 52 ms steps. In total, there were 168 (21 joints x 8 lags) positions and velocities for a
 239 total of 336 kinematic features. Other extrinsic features included kinematic synergies
 240 identified via dPCA. Additional categorical variables indicating the reach location and
 241 grip on a given trial were defined; inter-trial intervals were coded as one category in
 242 these variables.

243 *Intrinsic covariates*

244 In addition to extrinsic features, intrinsic features such as a given neuron's own spike
 245 history and the spike history of other neurons may also contribute to a neuron's firing
 246 rate. To account for the fact that spike history may be relevant in different ways at
 247 different time scales, we filtered binary spike trains with raised cosine basis functions of
 248 the form:

$$y(t) = 0.5 \cos(a \log(t + c) - \phi) + 0.5,$$

249 for t such that $a \log(t + c) \in [\phi - \pi, \phi + \pi]$, and 0 elsewhere (Saleh et al., 2012). We
 250 specified three different values of ϕ , the temporal peak of the cosine curve, to account
 251 for short (16 ms), medium (44 ms), and long (108 ms) time scale spike history effects.

252 Each basis vector was convolved with the binary spike train of a given neuron, thus
 253 giving rise to 3 spike history vectors for every neuron.

254 *Logistic regression*

255 We used logistic regression to predict the probability that a neuron fired a spike in a
 256 small time window (4 ms) based on previously described input features.

257 Mathematically, this model may be expressed as:

$$\log \left[\frac{p_n(t)}{1 - p_n(t)} \right] = \beta_0 + \sum_{m=1}^M \sum_{k=1}^K \beta_{k,m} X_k(t + \tau_m) + \sum_{j=1}^J \beta_j X_j(t),$$

258 where $p_n(t)$ is the probability that neuron n fires a spike at time t , β_0 represents the
 259 baseline probability that the cell will spike, $X_k(t - \tau_l)$ is the value of the k^{th} (of K) extrinsic
 260 feature at time $t - \tau_m$, where τ_m is the m^{th} (of M) lead or lag time (ranging from 52, 104,
 261 and 156 ms leading and 52, 104, 156, 208 ms lagging) against the spike time at t , and
 262 $X_j(t)$ is the value of the j^{th} (of J) intrinsic feature at time t with corresponding weights $\beta_{k,m}$
 263 and β_j , respectively.

264 *Data segmentation and assessing goodness of fit*

265 Model goodness of fit (GoF) was quantified using the area under the receiver operating
 266 characteristic curve (AUROC) (Hatsopoulos et al., 2007; Saleh et al., 2010, 2012;
 267 Truccolo et al., 2010). We assessed goodness of fit using 10 folds of cross-validated
 268 test data, and, unless otherwise noted, all reported AUROC values are taken to be the
 269 median across cross-validation folds. Each training fold was comprised of at least 329
 270 experimental trials and inter-trial activity between those trials. Because the folds were
 271 based on experimental trials rather than data points, the number of test and training

272 data points varied across folds, however, at least 531,747 and 1,541,496 data points
 273 were in every fold in animals J and L, respectively.

274 **Results**

275 *The kinematics of unconstrained reaching to grasp*

276 We used a marker-based infrared motion tracking system combined with an
 277 advanced biomechanical model of the primate upper limb (Delp et al., 2007) to infer the
 278 kinematics of 21 joints in the arm and hand as two rhesus macaques engaged in an
 279 unconstrained reach-to-grasp task. We presented multiple grips at different locations in
 280 the animal's workspace designed to elicit a diversity of prehensile movements that
 281 included both precision and power grips (Fig. 1).

282 Differences in kinematics across experimental conditions were evident at the
 283 level of individual joints (Fig. 2). From visual inspection of Fig. 2A, it can be seen that
 284 the different object locations elicited different patterns of arm movement and the
 285 different grip types elicited different patterns of hand movement. Less obvious,
 286 however, is that the object to be grasped also influenced arm kinematics and, to a
 287 lesser extent, the position of the object in space influenced hand kinematics (Fig. 2B).
 288 These differences at the level of single joints were also manifested in multi-joint
 289 features. The average temporal profiles of grasp apertures were qualitatively different
 290 across grips (Fig. 2C) and quantitatively, the maximum grasp aperture over time was
 291 significantly different across the four grasping conditions (Kruskal Wallis test, animal J:
 292 X^2 with 3 d.f. = 330.84, $p < 0.01$; animal L: X^2 with 3 d.f. 199.91, $p < 0.01$). Similarly, we
 293 observed differences in the peak wrist speed across object locations (Kruskal Wallis
 294 test, animal J: X^2 with 3 d.f. = 87.16, $p < 0.01$; animal L: X^2 with 3 d.f. 272.19, $p < 0.01$).

295 To control for the fact that both reaching and grasping affect kinematics in the
 296 entirety of the upper limb, we employed a novel dimensionality reduction technique,
 297 demixed principal component analysis (dPCA), to identify functionally defined kinematic
 298 synergies. Each kinematic synergy (i.e. each dPC) could be either reach-related,
 299 grasp-related, condition-independent (i.e. common to all experimental conditions), or an
 300 interaction between reach and grasp (i.e. specific to a particular experimental condition).
 301 As with other studies that have applied dimensionality reduction techniques to kinematic
 302 data (Mason et al., 2001; Mollazadeh et al., 2014; Schaffelhofer et al., 2015), we found
 303 that a handful of functional kinematic synergies described the majority of the variability
 304 in the kinematics. In both animals, we found 8 components were sufficient to explain
 305 90% of the variance in joint angles (Fig. 3A), while 10 and 9 components were needed
 306 to explain 90% of the variance in joint angular velocities in animals J and L, respectively
 307 (result not shown). In both animals, we observed that a substantial proportion of the
 308 kinematic variance was common across all experimental conditions, and that grasping
 309 components explained more variance than reaching components (Fig. 3B). Examples
 310 of the kinematic synergies in animal L are shown in Fig. 3C. The temporal evolution of a
 311 subset of these kinematic synergies as a function of time relative to the movement
 312 onset (MO) is shown in Fig. 4 for both animals. Although the convexity for animal J is
 313 flipped, dPC1's for both animals are condition independent and they appear to resemble
 314 aperture trajectories and clearly indicate behavioral differences between the two
 315 animals: animal J made much faster reaching than animal L. dpc1 CI for animal J
 316 appears to represent the invariant hand speed profile. Other grasp or reach specific
 317 dPC's show object and location specific components such that trajectories associated

318 with different locations formed clustered bundles that were distinct for different objects
 319 (dPC3 G for animal J and dPC2 G and dPC3 G for animal L), and trajectories
 320 associated with different objects formed bundles that were distinct for different locations
 321 (dPC4 R for animal J and dPC4 R for animal L).

322 *Encoding of extrinsic features in premotor cortex*

323 One session for each monkey was performed to collect single unit spiking activity
 324 from PMd and PMv arrays. We obtained populations of task related units: 51 and 34
 325 units from PMd and PMv respectively for monkey J and 29 and 22 for monkey L
 326 respectively. Figure 5 illustrates one example each for PMd and PMv single unit
 327 activities across different objects and different locations.

328 We used generalized linear models (GLM) to develop encoding models of
 329 premotor cortical single unit activity and compared the performance of these models
 330 that explicitly accounted for the moment-by-moment kinematic features against a more
 331 abstract, set-related representation of behavior. In this first class of model, we used the
 332 detailed kinematics of 21 joints in the arm and hand at a variety of different time lags to
 333 predict spiking activity (see Methods for details). In contrast, the second class of model,
 334 the set-related model, assumed a single, constant firing rate for each experimental
 335 condition. To assess model performance, the area under the receiver operating
 336 characteristic curve (AUROC) was computed for each model.

337 Across all animals, brain areas, and types of model, we found that the median
 338 AUROC values across cells were significantly greater than the chance level of the
 339 AUROC, 0.5 (Fig. 6A; Wilcoxon signed rank test, all $p < 0.01$). In both animals, we
 340 observed that kinematics better predicted activity in PMd as compared to PMv (Fig. 6A;

341 Mann-Whitney *U*-test, animal J: $Z = 2.85$, $p < 0.004$; animal L: $Z = 4.12$, $p < 0.01$). The
 342 set related models also predicted activity in PMd better than PMv in animal L (Fig. 6A; Z
 343 $= 4.75$, $p < 0.01$) but not in animal J ($Z = 1.59$, $p > 0.05$). In both animals and brain
 344 areas, we found that encoding models based on kinematics better predicted spiking
 345 activity than the set-related models (Fig. 6B; Wilcoxon signed rank test, animal J PMd: Z
 346 $= 5.74$, $p < 0.01$; PMv: $Z = 4.98$, $p < 0.01$; animal L PMd: $Z = 3.45$, $p < 0.01$; PMv: $Z =$
 347 4.11 , $p < 0.01$). We computed AUROC for each kinematic lead/lag, but there was no
 348 clear indication of an optimal lead/lag time for either of the two cortical areas (results
 349 now shown).

350 Having established that whole arm kinematics were predictive of spiking activity
 351 in PMd and PMv, we subsequently considered which aspects of the kinematics were
 352 important. The dual channels hypothesis posits that PMd is concerned with the
 353 proximal aspect of the limb while PMv is related to the distal aspect. We tested the
 354 extent to which anatomical segments of the limb were represented in both areas.
 355 Encoding models were fit using only arm kinematics, (i.e. shoulder and elbow
 356 kinematics), or only hand kinematics (i.e. wrist and fingers). We found evidence that
 357 both arm and hand kinematics were predictive of spiking in each area as the median
 358 AUROC across neurons was significantly greater than 0.5 (Fig. 7A; Wilcoxon signed
 359 rank test, all $p < 0.01$). But, this finding, in isolation, does not provide strong evidence
 360 against the dual channels hypothesis. For example, a cell that truly encodes only arm
 361 kinematics may appear to also encode hand kinematics simply because of correlations
 362 between arm and hand kinematics.

363 We performed a control analysis to ensure that any putative encoding
364 relationship was not a product of kinematic correlations. In this analysis, we subtracted
365 off either arm or hand AUROC from the AUROC of the full model containing both arm
366 and hand kinematics (Fig. 7B). Returning to the previous example, if a cell encodes
367 only arm kinematics, then the difference between the full model AUROC and the arm
368 AUROC should be 0. However, the difference between the full model AUROC and the
369 hand AUROC should be positive because this example cell truly encodes arm
370 kinematics, which the full model contains. In real data, cells that had a positive
371 difference between full model AUROC and hand model AUROC were classified as
372 encoding arm kinematics. Similarly, cells with positive differences between full model
373 AUROC and arm model AUROC were classified as encoding hand kinematics. These
374 classifications were not mutually exclusive. We found that both arm and hand
375 kinematics were effective predictors of single premotor cortical neuron activity. We
376 observed (Table 1) that 82% PMd and 79% PMv cells for monkey J and 100% PMd and
377 95% PMv cells for monkey L encoded arm kinematics, 56% PMd and 53% PMv cells for
378 monkey J and 75% PMd and 76% PMv cells for monkey L encoded hand kinematics,
379 while 42% PMd and 41% PMv cells for monkey J and 75% PMd and 71% PMv cells for
380 monkey L encoded both arm and hand kinematics respectively. The dual channels
381 hypothesis would predict that cells encode either arm kinematics or hand kinematics,
382 i.e., that they were anti-correlated. However, the proportion of cells that encoded both
383 arm and hand kinematics simultaneously was not significantly different than would be
384 expected if both arm and hand kinematics were encoded statistically independently

385 (binomial test, $B(0.497, 121) = 62$, $p > 0.05$), i.e., the presence of encoded arm
 386 kinematics does not affect the encoding of hand kinematics and vice versa.

387 We subsequently tested whether the arm and hand were preferentially
 388 represented in PMd and PMv, respectively by subtracting the AUROC of the hand-only
 389 from the arm-only models. We found that arm kinematics was preferentially
 390 represented in the PMd of both animals because the median AUROC differences were
 391 significantly greater than 0 (Fig. 7C; Wilcoxon signed rank test, animal J: $Z = 2.51$, $p <$
 392 0.02 ; L: $Z = 3.35$, $p < 0.01$). Arm kinematics was also preferentially represented in the
 393 PMv of animal J ($Z = 2.57$, $p < 0.01$), but not in animal L ($Z = -0.41$, $p > 0.05$).

394 In the previous analysis, we showed that both PMd and PMv encode both arm
 395 and hand kinematics. While this finding is seemingly in contradiction to the dual
 396 channels hypothesis, one possibility is that the hand representation in PMd is in the
 397 service of reaching and the arm representation in PMv is in the service of grasping.
 398 Thus, we explicitly tested if there is any preferential encoding of single unit activities of
 399 PMd and PMv neurons using functional representations of prehensile movements
 400 revealed by dPCA. Specifically, we fit encoding models using only reaching
 401 components, or only grasping components of the dPCA analysis. We found that both
 402 reach and grasp kinematic synergies predicted spiking activity in both PMd and PMv
 403 (Fig. 8A; Wilcoxon signed rank test, all $p < 0.01$).

404 Again, we performed a control analysis to verify that both reaching and grasping
 405 kinematic synergies were encoded in the spiking activity of premotor cortical cells. We
 406 assessed the extent to which a full model containing both reach and grasp synergies
 407 outperformed a model containing only reach, or only grasp (Fig. 8B). Here, we found

408 (Table 2) that 69% PMd and 56% PMv cells for monkey J and 93% PMd and 91% PMv
 409 cells for monkey L encoded reaching kinematics while 88% PMd and 65% PMv cells for
 410 monkey J and 97% PMd and 91% PMv cells for monkey L encoded grasping. The
 411 proportion of cells that simultaneously encoded both reach and grasp (monkey J: 67%
 412 PMd and 53% PMv cells and monkey L: PMd 90% PMd and 91% PMv cells
 413 respectively) was significantly more frequent than would be expected under chance if
 414 reach and grasp were independent (binomial test, $B_{0.628, 136} = 98$, $p < 0.01$). Although
 415 many cells encoded both reach and grasp simultaneously, subtracting the AUROC of
 416 the reach and grasp models revealed there was a preferential representation of
 417 reaching in the PMv of animal J and PMd of animal L because, as a population, the
 418 difference in AUROC between the reach only and grasp only models was significantly
 419 different from 0 (Fig. 8C, Wilcoxon signed rank test, J: $Z = 3.58$, $p < 0.01$; L: $Z = 3.75$, p
 420 < 0.01).

421 Finally, we considered the relationship between the anatomical and functional
 422 kinematic features encoded by each cell. To this end, we compared the difference of
 423 arm and hand model AUROCs to the difference of reach and grasp model AUROCs.
 424 Under the classical model, we might assume that PMd cells preferentially encode the
 425 kinematics of the arm and prefer the kinematics of reaching, and correspondingly, PMv
 426 cells prefer the hand and grasping (Fig. 9A). Such a simplistic relationship, however,
 427 does not appear to capture the patterns we observed in our data because we observed
 428 several cells that preferred both hand and reach or arm and grasp in both PMd and PMv
 429 (Fig. 9B). These results indicate that anatomical encoding preferences are not clearly
 430 related to functional encoding preferences as we failed to reject the null hypothesis that

functional and anatomical encoding preferences were statistically independent (chi-squared test of independence, χ^2 with 1 d.f. = 3.03, $p > 0.05$).

We additionally considered the strength of anatomical and functional representations in each area. Specifically, we sought to ask if anatomical or functional relationships were more strongly encoded in a given area. To do so, we computed the logarithm of the ratio of the arm AUROC and reach AUROC (Fig. 9C), or the hand AUROC and the grasp AUROC (Fig. 9D). If this number is negative, that would indicate that functional kinematic synergies predict spiking activity better than the anatomical features; however, if this number is positive, then the anatomical features predict spiking activity better than functional kinematic synergies. In both PMd and PMv, we found that anatomical features were more strongly encoded as compared to functional synergies because the logs of AUROC ratios were positive (Wilcoxon signed rank test, all $p < 0.01$). However, the degree to which anatomical features were preferred differed between areas. We found that anatomical features were significantly more strongly represented in the activity of PMd cells as compared to PMv (Fig. 9C-D; Mann Whitney *U*-test, arm / reach: $Z = 3.87$, $p < 0.01$, hand / grasp: $Z = 2.64$, $p < 0.01$).

Encoding of intrinsic features in premotor cortex

Since extrinsic features were not the sole predictors of spiking activity, we also explored how intrinsic features are encoded in premotor cortex. We compared the encoding performance of the full kinematic model to the full intrinsic, i.e. spike history model. The spike-history model contained information about the spike histories of all other simultaneously recorded neurons at a variety of temporal scales (Fig. 10; see methods for details). We found that spike history was an effective predictor of spiking

activity (Fig. 10A; Wilcoxon signed rank test, animal J PMd: $W = 91$, $p < 0.01$; PMv: $W = 78$, $p < 0.01$; animal L PMd: $Z = 4.7$, $p < 0.01$ PMv: $Z = 4.1$, $p < 0.01$). In animal J, we found spike history predicted PMd activity significantly better than PMv (Mann-Whitney U -test, $Z = 2.20$, $p < 0.03$), but there was no difference in animal L across areas ($Z = -0.33$, $p > 0.05$). Also, spike history better predicted spiking activity than the kinematics in all areas (Fig. 10B, all Bonferroni corrected $p < 0.011$).

We wanted to identify which temporal scale of intrinsic activities led to the best encoding performance. We fit models using each of the three different temporal scales and found that the longest temporal scale (108 ms) led to the best encoding performance in both animals irrespective of cortical area (Fig. 10C). We subsequently wondered if the information at each temporal scale was totally redundant, or, if using spike history at multiple temporal scales led to superior predictive ability. Accordingly, we compared the AUROC at the best temporal scale for each cell and compared it to the AUROC of the full spike history model including all temporal scales (Fig. 10D). In the PMd of animal J, we observed that there was no significant difference between the AUROC of the full spike history model and the model at the best lag (Wilcoxon rank sum test, PMd: $W = 69$, $p < 0.11$). In contrast, spike history at multiple spatial scales better predicted spiking activity in the PMv of both animals (J: $W = 69$, $p < 0.02$, L: $Z = 4.11$, $p < 0.01$) and the PMd of animal L ($Z = 3.30$, $p < 0.01$).

Discussion

Demixed principal component analysis

476 Here, we applied a novel dimensionality reduction technique, demixed principal
 477 component analysis (Kobak et al., 2016), to identify kinematic synergies associated with
 478 either reaching or grasping during an unconstrained prehensile movement task. In both
 479 animals, we observed that a few functionally defined kinematic synergies accounted for
 480 most of the variance in the data. Moreover, reaching and grasping synergies each was
 481 explained a large proportion of the kinematic variance. In contrast, comparatively little
 482 variance was due to interactions between reaching and grasping conditions, supporting
 483 the decomposition of kinematics into reaching and grasping sub-movements
 484 (Jeannerod, 1984; Haggard and Wing, 1995).

485 Although this method provides a promising avenue for gaining insight into
 486 prehensile movements, there are some limitations to the approach. Chiefly, the
 487 relationship between the demixed kinematic synergies and muscle synergies currently
 488 remains unknown. Using a similar approach to our current method, instantaneous
 489 lengths of 50 musculotendon units were estimated, and it was found that 8 PCA
 490 components were sufficient to explain more than 95% of variance while at least 11
 491 components were needed in the joint angle space to capture the same amount of
 492 variance for monkeys performing reach-to-grasp (Schaffelhofer et al., 2015). Recent
 493 work has argued that the encoding preferences of primary motor cortical neurons reflect
 494 muscle synergies evoked by intracortical microstimulation (Overduin et al., 2014). In
 495 complementary work, it was demonstrated that decoding joint kinematics from units on
 496 electrodes that evoked movements of that joint when stimulated led to better decoding
 497 performance as compared to a simple random allocation (Best et al., 2014). The
 498 kinematic synergies identified by dPCA, however, arose from our experimental design.

499 In future work, we wish to address the similarity between kinematic synergies inferred
500 using dPCA, and kinematic or muscle synergies characterized via ICMS.

501 At a more technical level, dPCA is limited in that it is based on trial-averaged
502 data and fails to account for trial-to-trial variation within an experimental condition. An
503 additional limitation is that it does not support the use of continuous variables as
504 conditions. That is, object locations were not treated as positions in Euclidean space,
505 but rather discrete covariates. Finally, on individual trials, the scores of the dPCs often
506 exhibited strong moment-by-moment correlations suggesting that even though reach
507 and grasp were decoupled across conditions, within a condition there may remain some
508 strong correlation.

509 *Implications for cortical control of reach to grasp*

510 Here, we used GLMs to develop encoding models of spiking activity in premotor
511 cortex based on intrinsic and extrinsic features. We found evidence that challenges the
512 dogmatic view of premotor cortical organization centered around discrete reaching
513 (dorsomedial) and grasping (dorsolateral) pathways. In particular, we demonstrated that
514 there was a complete anatomical representation of the upper limb in PMd. This finding
515 is consistent with previous studies that used intracortical-microstimulation to study
516 motoric representations in PMd and further work that showed firing rates of PMd
517 neurons are modulated by changes in grasping condition (Raos et al., 2003, 2004;
518 Fattori et al., 2010). Additional work has demonstrated that whole arm kinematics can
519 be decoded from PMd ensemble activity (Bansal et al., 2012). Our work complements
520 that study by showing that single cells encode complex, whole-arm kinematics.
521 Furthermore, our finding is consistent with studies on single units from area V6A, which

522 is a part of the dorsomedial reaching network, illustrating that the dorsomedial pathway
 523 is also involved in hand preshaping and grip formation and may play a central role for all
 524 phases of reach-to-grasp movements (Fattori et al., 2010, 2012). Similarly, in PMv, we
 525 also found a complete representation of the upper limb, although, in absolute terms, it
 526 was weaker than in PMd. This result is consistent with previous reports that ICMS of
 527 PMv occasionally evoked arm movements (Godschalk et al., 1981, 1985; Hocherman
 528 and Wise, 1991; Stark et al., 2007).

529 Kinematically, we demonstrated that arm kinematics were affected by grasping
 530 condition and hand kinematics were affected by reaching condition, so, while we
 531 demonstrated that whole arm representations were present in both PMd and PMv, that
 532 was insufficient to argue against our null hypothesis that reach and grasp were
 533 processed in independent pathways. We used a novel dimensionality reduction
 534 technique to dissociate reaching from grasping and found that single neurons in PMd
 535 and PMv encoded both reaching and grasping kinematic synergies. Moreover, across
 536 the entire population of recorded cells, we found no relationship between anatomical
 537 and functional representations. That is, the anatomical preferences of a cell (i.e.
 538 whether it was biased towards arm or hand kinematics) were not related to its functional
 539 preferences (i.e. whether it preferred reach or grasp synergies).

540 *Differences between PMd and PMv*

541 In this work, we compared the encoding properties of neurons simultaneously
 542 recorded in the dorsal and ventral premotor cortex of two rhesus macaques. By using
 543 this simultaneous recording paradigm, our results were not confounded by potential
 544 behavioral differences across sessions.

545 In both animals, we observed that movement kinematics was more strongly
 546 represented in PMd as compared to PMv. One potential explanation for this finding is
 547 that PMd is seemingly more directly related to motor output. Pyramidal cells in layer V
 548 of PMd comprise approximately 10% of fibers from the frontal lobe in the corticospinal
 549 tract (Dum and Strick, 1991), and intracortical microstimulation of PMd is known to elicit
 550 arm movements (Weinrich and Wise, 1982; Weinrich et al., 1984). Consistent with this
 551 interpretation, we found that spiking activity in PMd was generally better predicted by
 552 anatomical features rather than functional kinematic synergies. PMv too sends
 553 projections to spinal cord, but it largely projects to the upper cervical spinal segments
 554 (He et al., 1993). Recent work using ICMS has shown that PMv acts on spinal cord
 555 primarily through facilitation of MI (Cerri et al., 2003; Shimazu et al., 2004).

556 In both animals, we observed that while single cells in PMd encode whole arm
 557 kinematics, they exhibit a preferential encoding of arm kinematics. This is consistent
 558 with the classical studies of PMd implicating it in the control of reaching movements
 559 (Weinrich and Wise, 1982; Weinrich et al., 1984; Kurata and Tanji, 1986; Riehle and
 560 Requin, 1989; Pesaran et al., 2006).

561 In PMv, however, we observed no consistent bias for either arm or hand
 562 kinematics or grasp or reach kinematics despite some studies illustrated causal
 563 influence of PMv lesion to a grasp deficit (Fogassi et al., 2001). Additionally, in PMv, we
 564 found that spike history at multiple time scales was a better predictor of spiking activity
 565 than a single timescale. This suggests that PMv may be more influenced by several
 566 internal processes operating at different timescales, and may also explain why
 567 kinematics was not strongly represented in PMv. An alternative possibility is that these

types of unconstrained movements are not the most effective drivers of activity in PMv, but rather a different class of movement altogether may be represented (Wise, 2006; Lehmann and Scherberger, 2013; Bonini et al., 2014). It should be noted that the PMv arrays were placed on or close to the convexity of the inferior arm of arcuate sulcus. Thus, we were not sampling from neurons in the sulcus, particularly F5p, and were likely recording from neurons in F4 as well as F5 divisions of PMv (Matelli et al., 1985).

Inter-animal differences

We observed some differences in behavior between the two animals. Animal J had faster reaction times than animal L, and, generally, executed faster movements that were qualitatively more stereotyped. Although the movements were executed at different speeds and with different reaction times, we observed many similarities in the dPCA analysis across the two animals suggesting that the kinematic synergies we identified were robust across a naturalistic range of self-selected movement speeds. In future work, it could be tested if the same set of dPCA synergies predicts movement kinematics at a variety of speeds by having an animal engage in a reach-to-grasp task with an explicit speed cue.

An additional difference in the kinematics between the two animals was their resting posture. In animal L, resting grasp aperture was proportionately much larger than in animal J. Indeed, the grasp aperture was maximal at the rest position for some trials in animal L. In contrast, animal J maintained a more closed aperture during the rest epoch. This finding may explain why the temporal profiles of the largest dPC in each animal are different. The temporal profile of the largest condition independent (CI)

dPC in animal J was a bell-shaped curve, reflecting that the hand opened and closed during the reach to grasp movement. In contrast, the temporal profile of the largest CI dPC in animal L was a sigmoidal curve corresponding to the fact that his hand started opened and gradually conformed to the shape of the object to be grasped. Finally, the animals adopted qualitatively different strategies to successfully grasp the ring object. Grasp aperture of the ring object in animal J followed a biphasic profile that was absent in animal L.

At the neural level, there were also several differences between the two animals. We observed that arm (as compared to hand) kinematics was preferentially represented in PMv of animal J, but not L. One potential explanation of this finding is based on array placement. Although an attempt was made to place the arrays in approximately the same location in both animals, a large blood vessel constrained our placement of the PMv array in animal J. In both animals, we saw a preferential representation of the arm in PMd. However, animal J's PMv array was inserted much closer to its PMd array because of the aforementioned blood vessel as compared to animal L. Accordingly, the arm representation in the PMv of J may be stronger because it is closer to PMd.

In summary, we employed GLM framework to characterize encoding properties of single neurons in PMd and PMv with kinematic inputs that were computed using a novel dimensionality reduction technique (dPCA) to dissociate anatomical representations of arm and hand from functional representations of reach and grasp. We showed that kinematics was more prominently encoded in PMd responses as compared to PMv. More importantly, there was no consistent preference for arm versus hand or for reach versus grasp kinematics in either PMd or PMv. By considering the

614 encoding of kinematic trajectories, our findings extend but are consistent with the results
 615 in a previous study (Stark and Abeles, 2007) that demonstrated that the encoding
 616 properties of PMd and PMv neurons often show mixing between reach and grasp which
 617 suggests that these two cortical areas may functionally serve as a neural substrate for
 618 coordination between reach and grasp during prehension.

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770 Figure Legends

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772 **Figure 1 Methods. A.** Animals were trained to grasp a variety of geometric shapes
773 including (left to right) ring (vertical), small disc (out), key, small Disc (horizontal),
774 cylinder (horizontal). These drawings depict static hand conformation during grasping
775 of these objects. **B.** Target locations in relation to the starting position. Aerial view (Top)
776 and side view (bottom). Blue circle denotes the starting position and white circles
777 denote target locations respectively. Red numbers in or near the target circles indicate
778 the location numbers, 1-4. **C.** Drawing of animal J in rest position. Pink dots correspond

to the approximate placement of the infrared markers. **D.** Depiction of the arm in the same position as **C.** in the Opensim software environment. **E-F.** Placement of electrode arrays in animals L and J, respectively.

Figure 2 Trial-averaged kinematics of the arm, wrist and hand. **A.** Within-condition trial-averaged kinematics for three degrees of freedom in the arm (left column), wrist (center column) and hand (right column) of animal J. Line style indicates object location while color indicates grip types. **B.** We found that the object can significantly affect arm kinematics (top). Here, we computed the average humerus abduction over all object locations when grasping Cyl o (purple trace) and Ring h (green trace). Shaded area indicates ± 2 standard errors of the mean (S.E.M.) Similarly, reach location can affect hand kinematics (bottom). Flexion in the metacarpophalangeal joint of the second digit (2mcp flexion) was averaged over all objects and is shown for two object locations. **C.** We computed the grasp aperture, defined as the distance between the distal-most thumb and index finger markers and plotted it relative to movement onset in both animals.

Figure 3 Demixed principal component analysis. **A.** Scree plot of demixed PCA. The cumulative percentage of explained variance for a given number of dPCs is shown for joint angles. Each individual component varies only along one task parameter. It can be either reach related (R, red points), grasp related (G, green points), condition independent (CI, gold points), or an interaction between reach and grasp condition (I, blue points). Marker shape (either square or triangle) indicates animal. Note that the scree plots are highly similar across animals. **B.** Percentage of variance explained by the different task conditions in joint angle data. **C.** Visualization of kinematic synergies revealed by the demixed PCA. The posture of the upper limb is shown at various projections along the largest condition independent synergy (top row), the largest grasp synergy (middle row), and the largest reach synergy (bottom row). Data are from animal J.

Figure 4 Trial averaged dPCA trajectories relative to movement onset. The temporal profile of the first four dPCs is shown for the various object locations and grips. Conventions are the same as Fig. 2. To the left of each dPC is an indication of whether it is a condition-independent (CI), grasp (G), or reach (R). The components are ordered from top to bottom in terms of variance explained; that is, dPC1 explained more variance than all others.

Figure 5 Exemplary single unit spiking activities of neurons in PMd (A) and PMv (B) for monkey L reaching to grasp different objects at different locations. Each raster plot corresponds to a particular object and location combination while each PSTH was computed for either one object over all locations or one location over all objects. Time 0 corresponds to movement onset.

Figure 6 Encoding models that include kinematics predict spiking activity better than set-related models. **A.** We fit encoding models using either set-related activity or kinematics to each neuron in our sample and measured the area under the receiver

operating characteristic curve (AUROC) across 10 folds of cross validated data. Here, each point corresponds to the median AUROC across folds as a function of animal (triangles or squares) and model type. Cells in PMd are blue, while cells in PMv are red. The vertical line next to each point-set indicates the interquartile range while the median is indicated with the horizontal line. **B.** Distribution of differences between kinematic and set-related models. The lines to the immediate right of each point-set follow the same conventions as **A.** Kinematic models predicted spiking activity better than set-related activity models in PMd and PMv in both animals (significance codes: *** $p < 0.0005$, see main text for additional statistics).

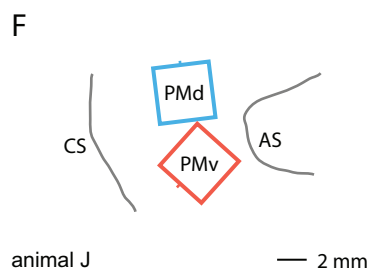
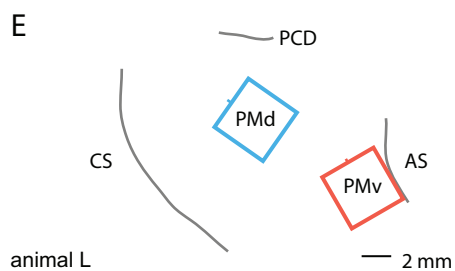
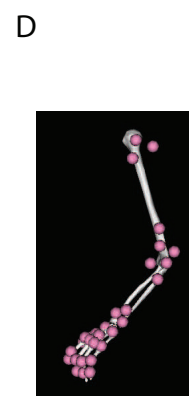
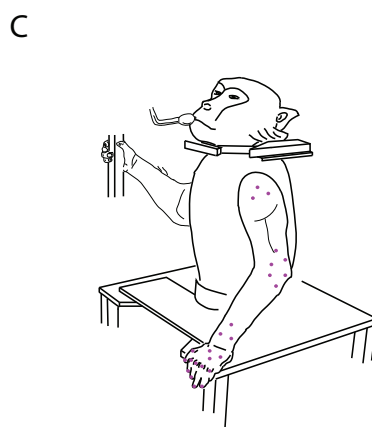
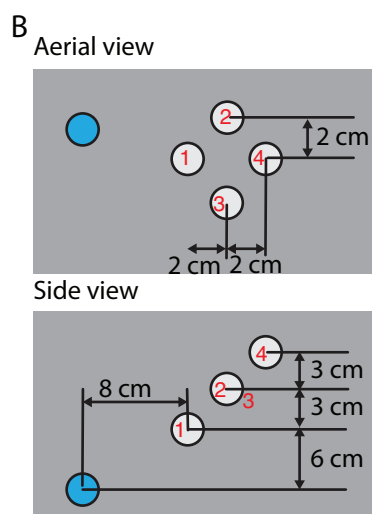
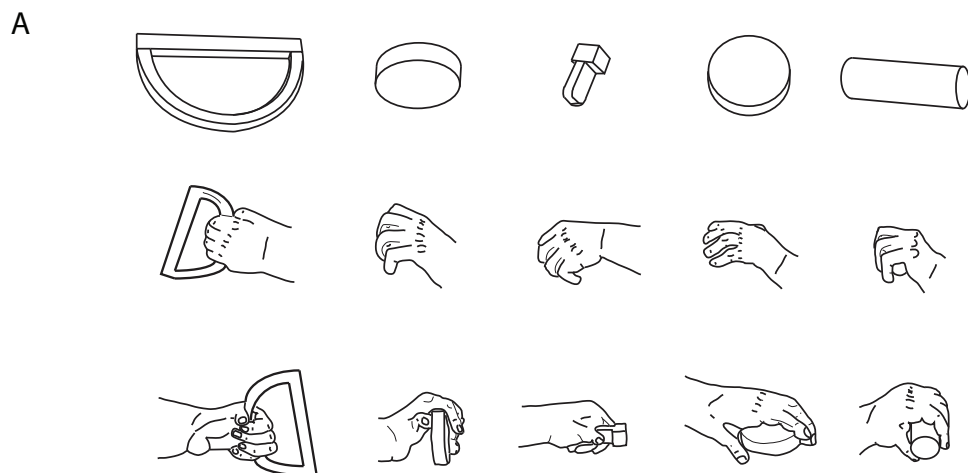
Figure 7 Comparing encoding performance of models based on an anatomical division of the upper limb. **A.** We fit GLMs using either arm or hand (including wrist) kinematics. For each model type and animal, the distribution of median AUROCs across folds is shown as raw data (left column; each point corresponds to a cell) and interquartile range + median (vertical and horizontal lines, respectively). **B.** Median AUROCs were compared between the full model that included all kinematic terms, and reduced models that contained only arm, or only hand kinematics. **C.** Distribution of differences in arm and hand AUROCs. We found that arm kinematics were preferentially encoded by PMd in both animals and also in PMv in animal J because a significant proportion of cells preferred arm kinematics in those two areas (significance codes: * $p < 0.05$, ** $p < 0.005$; see main text for additional statistics).

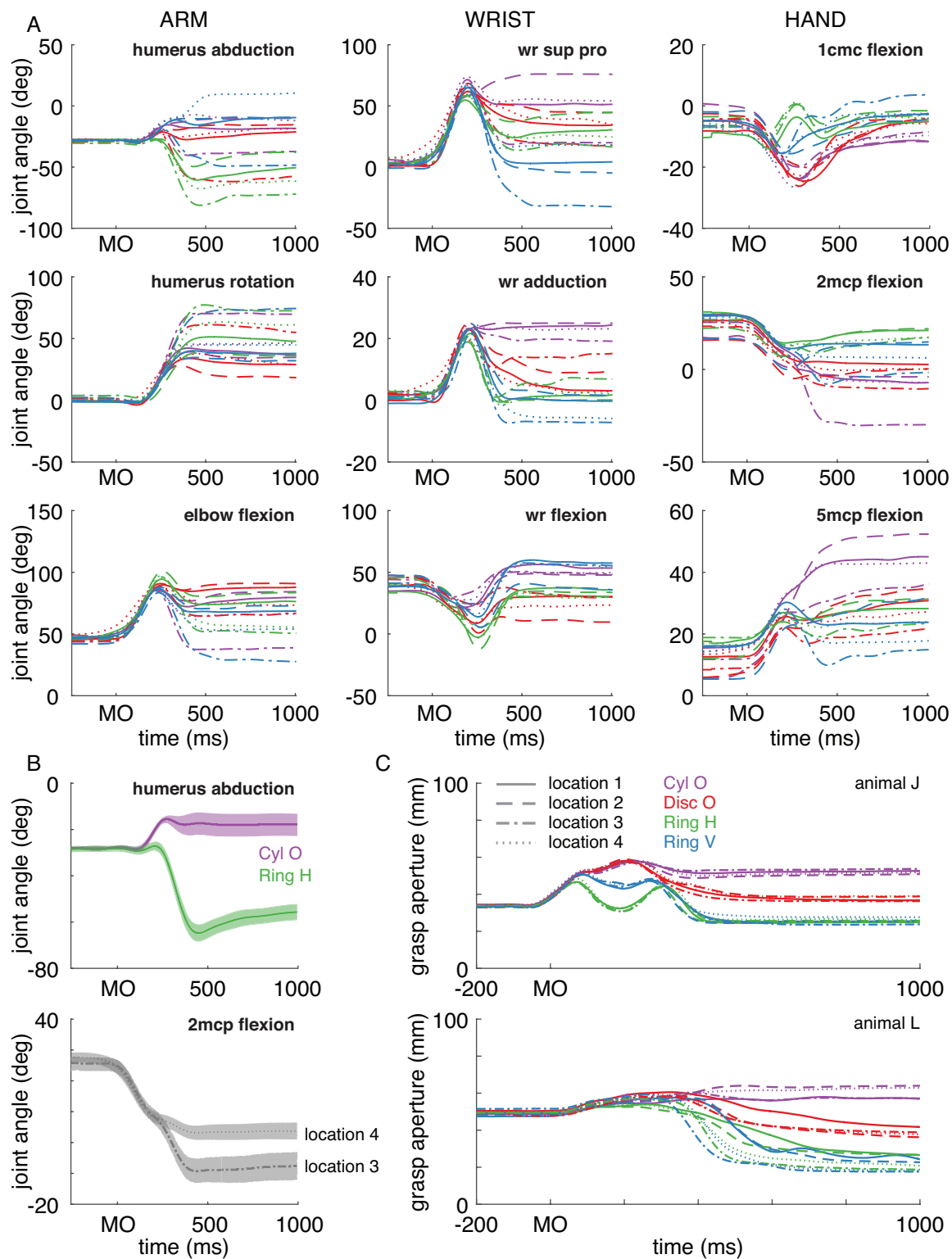
Figure 8 Comparing encoding performance of models based on a functional division of the upper limb. **A.** We fit GLMs using either reach or grasp kinematic synergies. For each model type and animal, the distribution of median AUROCs across folds is shown as raw data (left column) and interquartile range + median (vertical and horizontal lines, respectively). **B.** Median AUROCs were compared between the full model that included both reach and grasp synergies, and reduced models that contained only reach or only grasp synergies. **C.** Distribution of differences in reach and grasp AUROCs. We found that reach kinematic synergies were preferentially encoded by PMv in animal J and PMd in animal L (significance codes: *** $p < 0.0005$ see text for additional statistics).

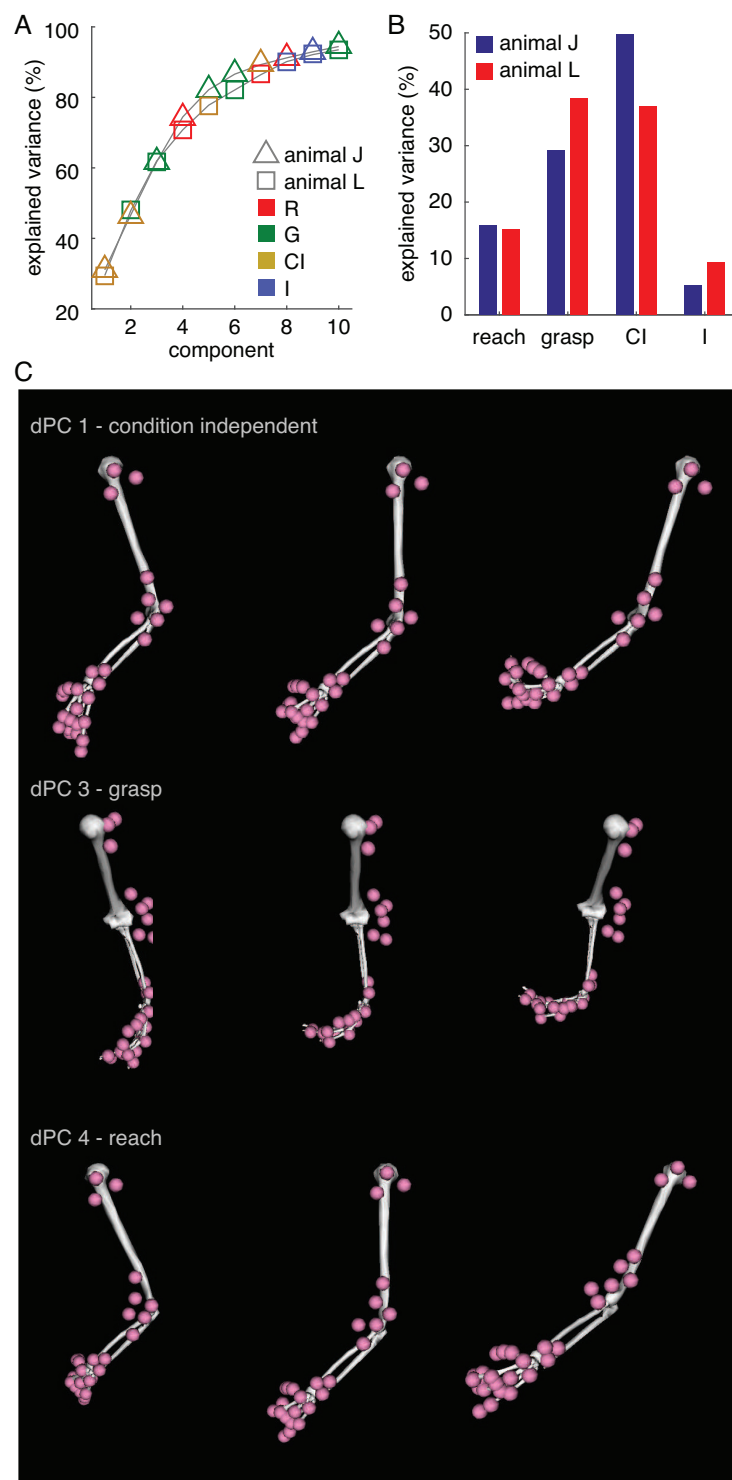
Figure 9 Comparing anatomical and functional representations in premotor cortex. We compared the differences of arm and hand AUROCs to the differences of reach and grasp AUROCs to determine if there was a systematic relationship such that cells that preferentially encoded arm kinematics likewise encoded reaching synergies, and similarly, cells that preferentially encoded hand kinematics likewise encoded grasping synergies. **A.** Expected results based on the dual channels hypothesis. PMd would be concerned exclusively with arm kinematics and reaching, while PMv would be concerned with hand kinematics and grasping. There would be a segregation of both function and anatomy between PMd and PMv. **B.** Actual results based on observed data. A chi-squared test of independence revealed no significant relationship between functional and anatomical encoding preferences. **C.** Comparing the relative encoding performance of anatomical and functional models. Here, we show the distribution of the logarithm of the ratio of arm and reach model AUROCs. If the log of this ratio is positive, then the arm model had a higher AUROC than the reach model, and vice versa for negative values. We found that both PMd and PMv are better predicted by

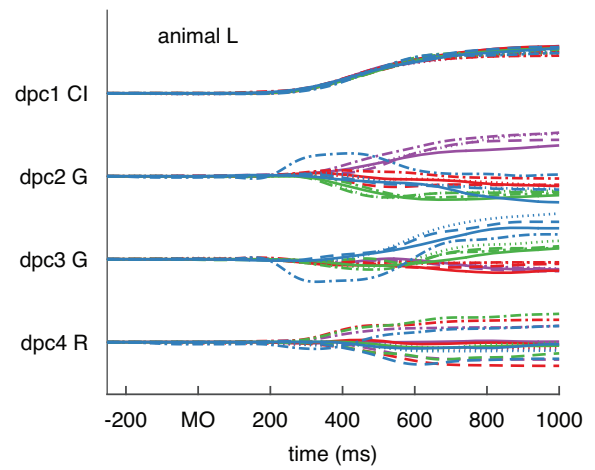
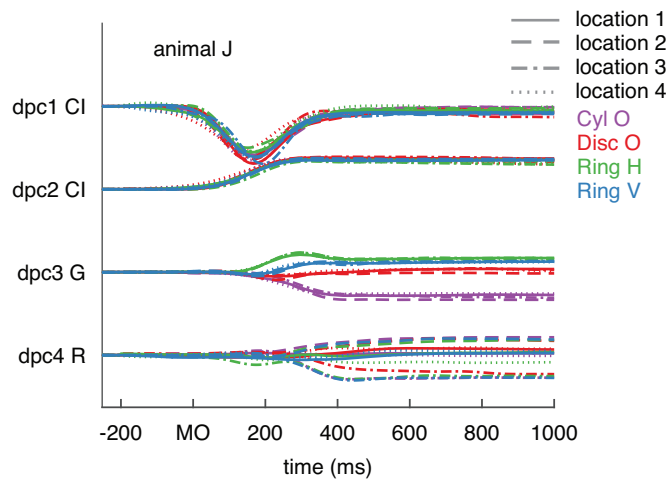
872 anatomical features rather than functional features, but this preference is stronger in
873 PMd (see main text for additional statistics). **D.** Same conventions as panel **C.** but
874 using hand and grasp models.

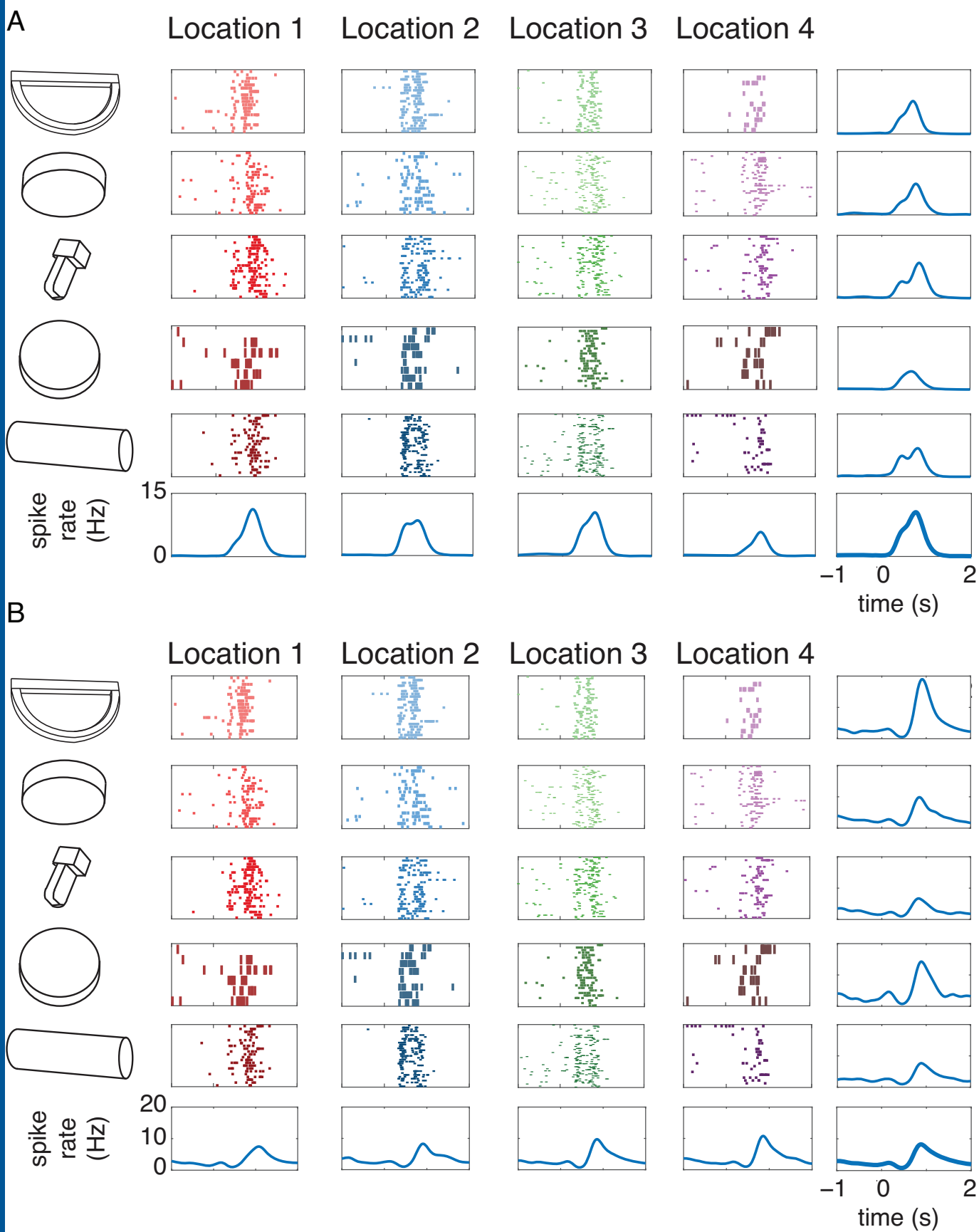
875
876 **Figure 10 Encoding properties of intrinsic features and comparing spike history**
877 **with kinematics in premotor cortex.** **A.** We found that spike history from all cells at
878 multiple temporal scales effectively predicted spiking activity (see main text for
879 statistics). Each point corresponds to a cell from a given animal (shape) and cortical
880 area (color). Lines next to each column of data indicate interquartile range and median.
881 **B.** We compared the performance of the spike history model to the full kinematic model
882 and observed that generally, spike history was a more effective predictor of spiking
883 activity than kinematics. **C.** Median AUROC across cross validation folds as a function
884 of basis function, cortical area, and animal. In both areas, the longest spike history term
885 had the best encoding performance. **D.** Distribution of the differences in AUROC
886 between the full spike history model and a model based on the best single basis
887 function (lag). To the right of each data column are the median and interquartile ranges
888 of the data.

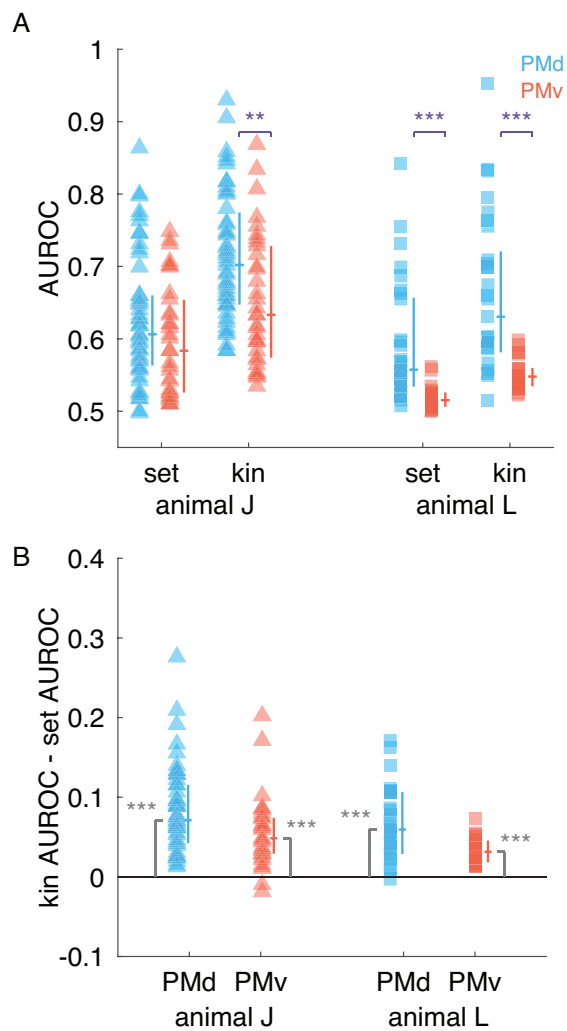


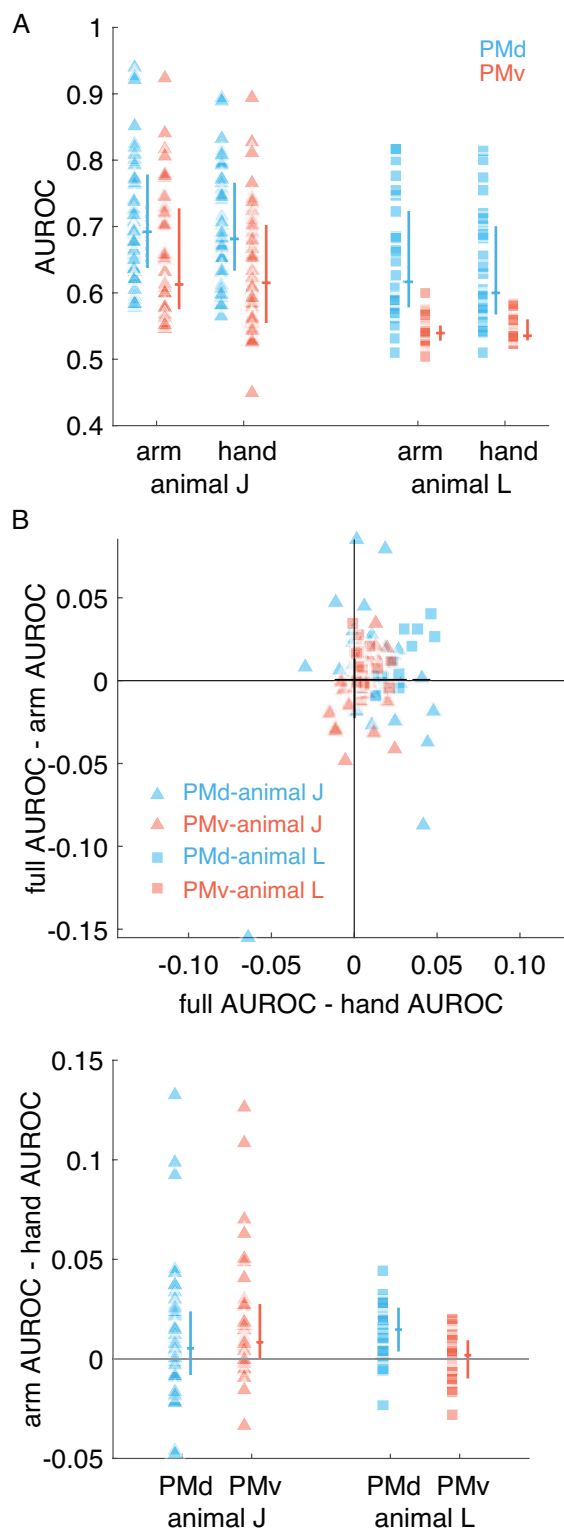


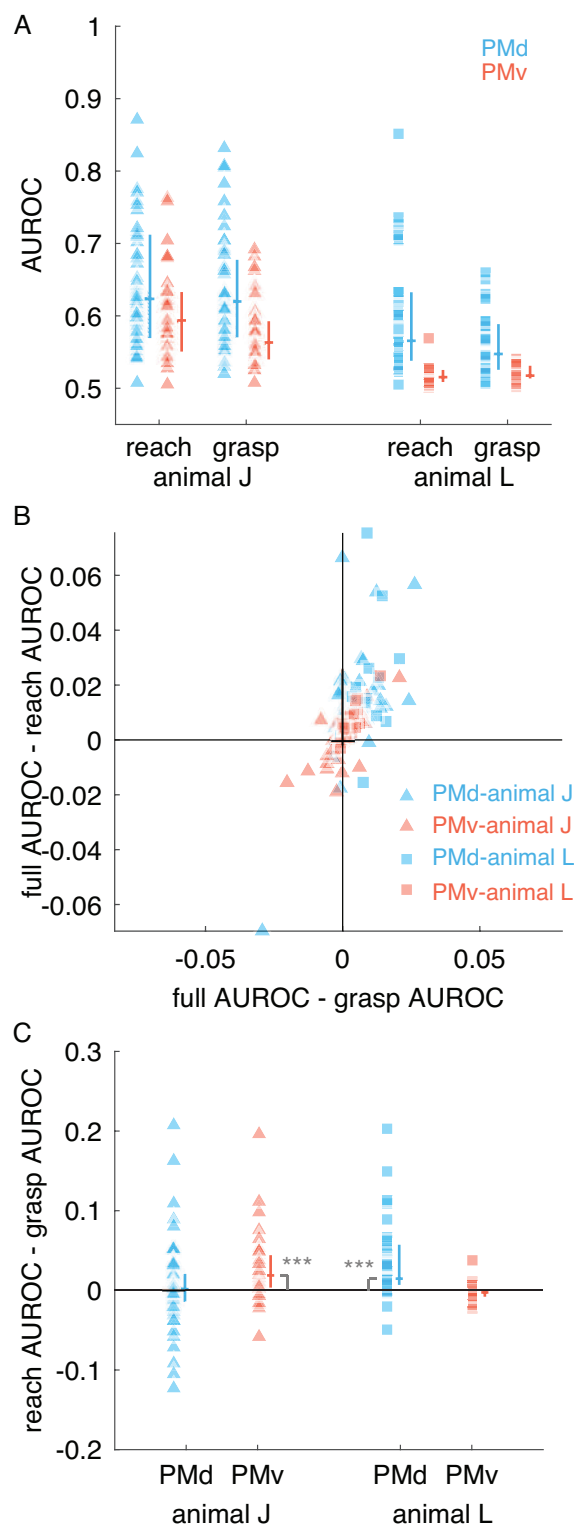


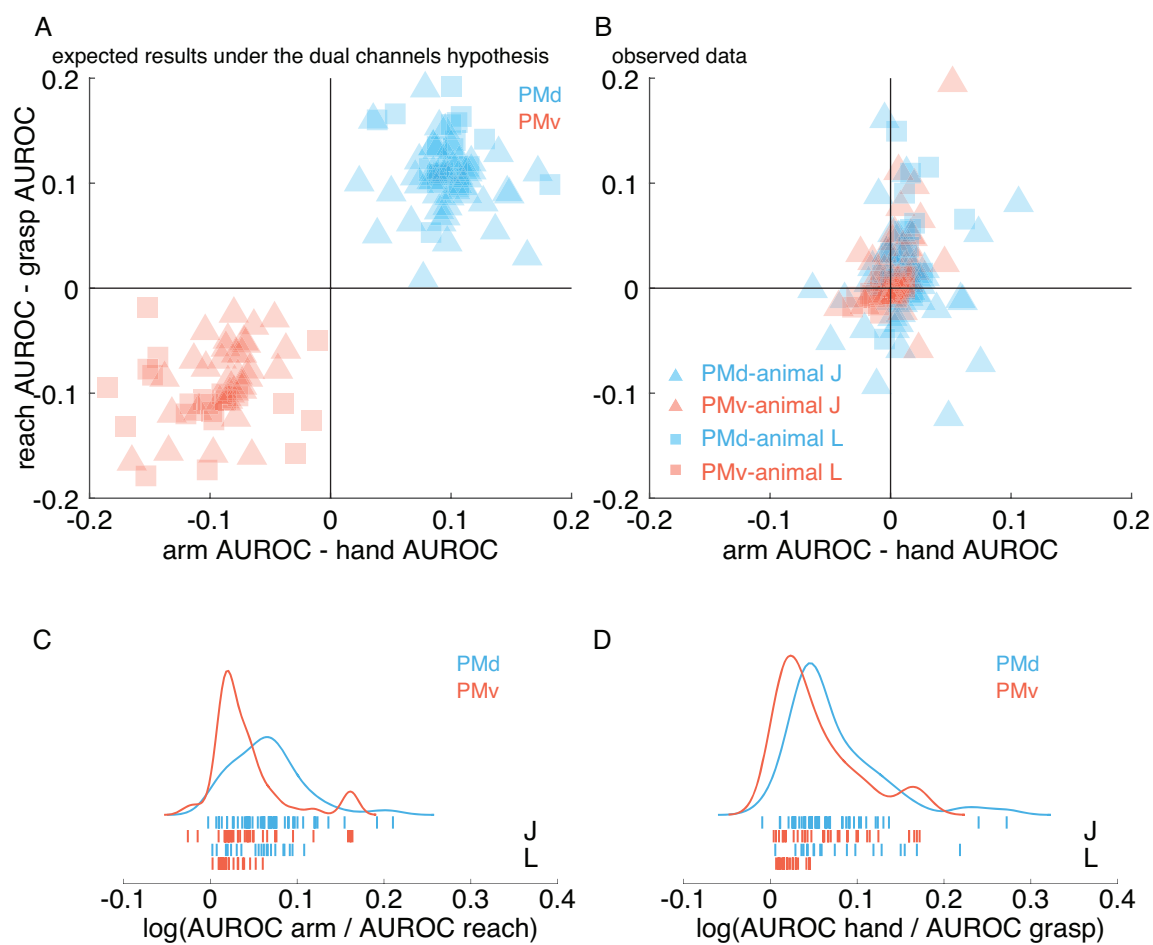


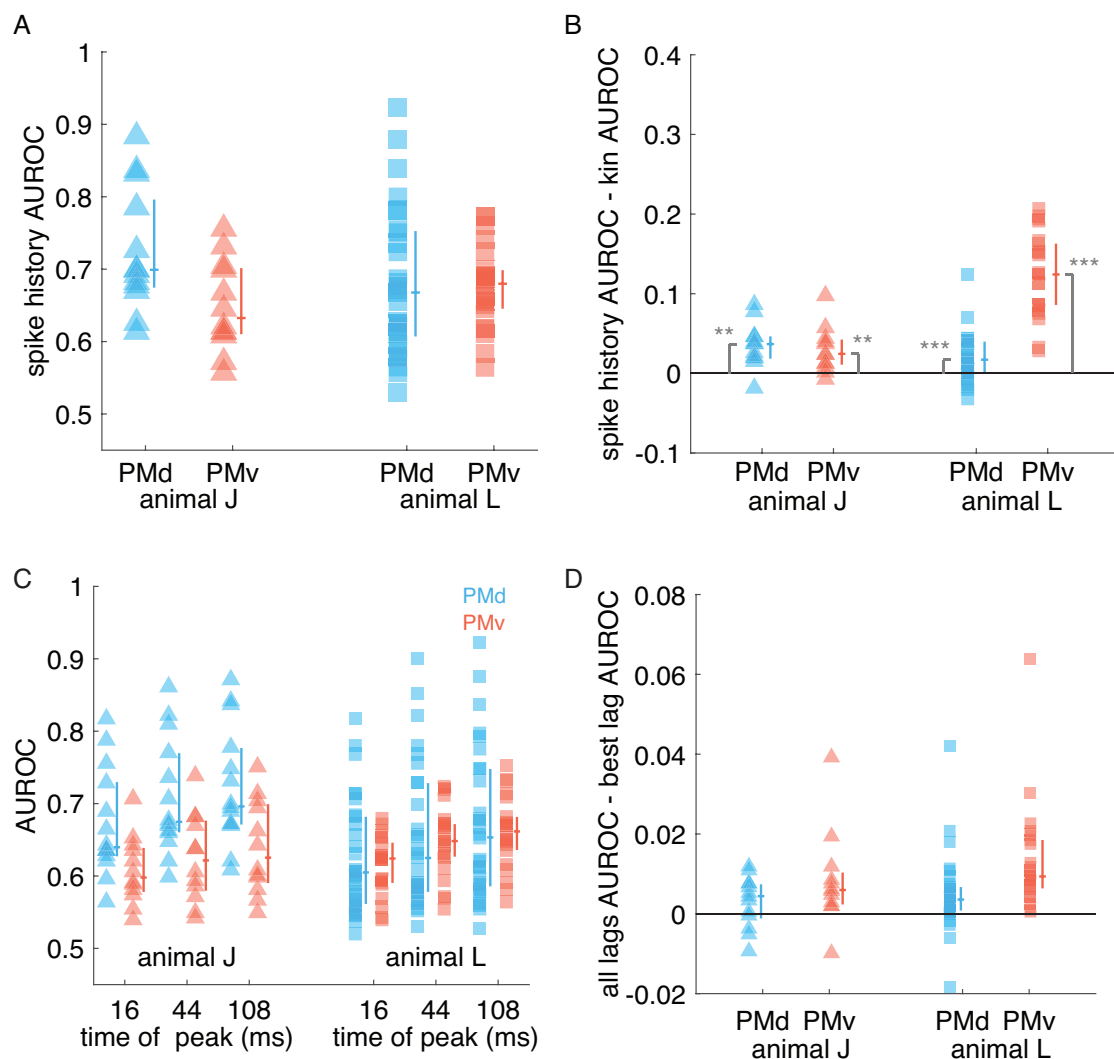












1 Tables

2

	PMd		PMv	
	animal J	animal L	animal J	animal L
Arm	82% (41/50)	100% (16/16)	79% (27/34)	95% (20/21)
Hand	56% (28/50)	75% (12/16)	53% (18/34)	76% (16/21)
Arm+Hand	42% (21/50)	75% (12/16)	41% (14/34)	71% (15/21)

3
4 Table 1. Percentage (number of neurons/total number) of neurons encoding arm, hand,
5 or both arm & hand kinematics by area and animal.
6
7

	PMd		PMv	
	animal J	animal L	animal J	animal L
Reaching	69% (35/51)	93% (27/29)	56% (19/34)	91% (20/22)
Grasping	88% (45/51)	97% (28/29)	65% (22/34)	91% (20/22)
Reaching+Grasping	67% (34/51)	90% (26/29)	53% (18/34)	91% (20/22)

8
9
10 Table 2. Percentage (number of neurons/total number) of neurons encoding reaching,
11 grasping, or both reaching & grasping kinematics by area and animal.
12