

Research Report

Video-Based Automated Assessment of Movement Parameters Consistent with MDS-UPDRS III in Parkinson's Disease

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Abstract.

Background: Among motor symptoms of Parkinson's disease (PD), including rigidity and resting tremor, bradykinesia is a mandatory feature to define the parkinsonian syndrome. MDS-UPDRS III is the worldwide reference scale to evaluate the parkinsonian motor impairment, especially bradykinesia. However, MDS-UPDRS III is an agent-based score making reproducible measurements and follow-up challenging.

Objective: Using a deep learning approach, we developed a tool to compute an objective score of bradykinesia based on the guidelines of the gold-standard MDS-UPDRS III.

Methods: We adapted and applied two deep learning algorithms to detect a two-dimensional (2D) skeleton of the hand composed of 21 predefined points, and transposed it into a three-dimensional (3D) skeleton for a large database of videos of parkinsonian patients performing MDS-UPDRS III protocols acquired in the Movement Disorder unit of Avicenne University Hospital.

Results: We developed a 2D and 3D automated analysis tool to study the evolution of several key parameters during the protocol repetitions of the MDS-UPDRS III. Scores from 2D automated analysis showed a significant correlation with gold-standard ratings of MDS-UPDRS III, measured with coefficients of determination for the tapping (0.609) and hand movements (0.701) protocols using decision tree algorithms. The individual correlations of the different parameters measured with MDS-UPDRS III scores carry meaningful information and are consistent with MDS-UPDRS III guidelines.

Conclusion: We developed a deep learning-based tool to precisely analyze movement parameters allowing to reliably score bradykinesia for parkinsonian patients in a MDS-UPDRS manner.

Keywords: Bradykinesia, deep learning, Parkinson's disease, MDS-UPDRS III

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INTRODUCTION

The diagnosis of Parkinson’s disease (PD) is based on the presence of a parkinsonian syndrome, i.e., the association of rigidity and/or rest tremor to bradykinesia, this latter being mandatory for the diagnosis [1]. Bradykinesia is defined as a motor slowness associated with a decrease in the amplitude and/or speed of movement [2–4]. Currently, the evaluation of motor impairment of PD is based on part III (motor) of the Movement Disorder Society-Unified Parkinson’s Disease Rating Scale (MDS-UPDRS) composed of 18 items rated from 0 (normal) to 4 (severe), which assess the severity of bradykinesia, rigidity and tremor [5, 6]. This score is used worldwide for patient follow-up in outpatient clinics but also in clinical research and more specifically in therapeutic trials. However, the semi-quantitative assessment of parkinsonian symptoms by the MDS-UPDRS III score suffers from a certain subjectivity and the reproducibility of the measurements arising from assessors is questionable especially in case of non-parkinsonian expertise [7–13]. This may contribute to the difficulty in the follow-up of PD patients and also to induce some biases in clinical research due to the multiplicity of assessors and clinical centers, and variability across longitudinal iterative visits.

Although digital tools have been recently developed in an attempt to improve bradykinesia scoring by providing quantitative measures [14–16], they exhibit limited use in practice because they rely on wearable sensors on patients [17, 18] in a dedicated room or need specific conditions [15, 18–20]. More recent studies aimed to circumvent these material issues by developing video-based only tools [12, 21–24]. First, machine learning-based automatic rating for rest tremor and finger tapping was investigated to measure bradykinesia [12]. The authors showed that it was feasible, reliable when compared to movement disorders specialist ratings, and more accurate than a non-trained one. The analysis was based on the periodicity of the repeating task videotaped, i.e., the frequency of the movement, to predict a severity score, but the outcomes did not fully reflect the MDS-UPDRS guidelines in which decreases in speed and amplitude of movement over time were also considered [12]. The global PD severity was assessed by applying deep learning approach to automatically score the severity level by compiling 7 of the 18 items of the MDS-UPDRS III [22]. MDS-UPDRS III bradykinesia was investigated for upper limbs tasks but with the constraint of fixed webcam record-

ings and with different assessment guidelines from MDS-UPDRS [21, 22]. In the same way, Guo et al. [25] demonstrated that a computer-vision approach to obtain finger tapping 3D acquisition is feasible in patients with PD. Such approaches have also been developed to successfully study gait of parkinsonian patients [23, 24]. Although these studies bring interesting approaches, tools still need to be developed to establish metrics closer to the MDS-UPDRS III guidelines for a better assessment over time of movement parameters of bradykinesia.

In this emerging context of computer-based tools and given the importance of bradykinesia evaluation in the PD diagnosis, we developed a tool based on hand-pose estimation and movement analysis to compute an objective score of bradykinesia with multiple medically-relevant parameters using a deep learning approach in accordance with MDS-UPDRS III guidelines [26].

MATERIALS AND METHODS

Participants and videos

We included only PD patients according to MDS clinical criteria [1], with the exception of one MSA patient [27] and another one with undetermined atypical parkinsonian syndrome. All patients were consecutively seen in an outpatient clinic by a movement disorder specialist (BD) at Avicenne University Hospital between June 2019 and December 2020. All patients were video-recorded, using standard smartphones (rescaled to 720×1280 pixels with 30 or 60 fps), for the items 3.4, 3.5, and 3.6 of the MDS-UPDRS III corresponding to the assessment of finger tapping, hand movements, and pronation-supination movements of the hands, respectively.

The videos were then stored in a secure database, according to the French data protection authority (Commission Nationale de l’Informatique et des Libertés) recommendations. All participants gave their written informed consents for videos and their analyses. The study was approved by the local ethics committee (CLEA-2019-83) and registered in ClinicalTrials.gov (NCT04974034).

MDS-UPDRS III ratings of the video recordings

Two movement disorders specialists certified for MDS-UPDRS (BD, MM) and three neurologists trained for movement disorders, but not certified for MDS-UPDRS scoring (CD, QS and BG), rated all the

131 videos for both hands according to the items 3.4, 3.5,
132 and 3.6 of the MDS-UPDRS III [5]. Assessors were
133 blinded to the ON or OFF state of patients.

134 The python package penguin was used to compute
135 the Intraclass Correlation Coefficient (ICC), with
136 options `nan_policy = 'omit'`. ICC values below 0.50,
137 between 0.50 and 0.75, between 0.75 and 0.90 and
138 above 0.90 are considered as poor, moderate, good,
139 and excellent, respectively.

140 *Hand-pose estimation using deep learning* 141 *algorithms*

142 We analyzed videos using two deep learning algo-
143 rithms in order to extract interesting temporal features
144 of the movement. We used a first network, DeepLab-
145 Cut [26], to extract predefined points of interest
146 in 2D from images. Using an associated software
147 (<https://www.mackenziemathislab.org/deeplabcut>),
148 we roughly labelled 5 images per video with 22
149 different points (5 for each finger, one for the wrist,
150 one for the center of the palm) and the network
151 was trained to detect the 22 joints [28] using the
152 DeepLabCut toolbox. After the 2D hand coordinates
153 detection, several algorithms (see next section)
154 were applied to filter and smooth the movement
155 trajectories. We defined a bounding box around the
156 patient's hand from the 2D coordinates obtained with
157 DeepLabCut for each timestep, and each frame was
158 cropped along this box. The cropped image was pro-
159 cessed through a second network, HandGraphCNN
160 [29], which without specific training predicted 2D
161 and 3D positions of the 21 points (all the previous
162 ones, except center of palm). We then processed and
163 analyzed all trajectories (2D DeepLabCut, 2D & 3D
164 HandGraphCNN) to study the temporal evolution
165 of the movement for each protocol. We realized
166 training and inference using Python 3.X, tensorflow
167 and Pytorch on a GPU Nvidia Geforce GTX 1080 Ti.
168 The deep-learning algorithms inferred the positions
169 of the hand points frame by frame, and the detection
170 was achieved correctly for a large majority of the
171 frames. However, sometimes the hand position was
172 not correctly inferred for a single frame, usually at
173 unrealistic coordinates. We detected and excluded
174 such outliers in our analysis.

175 *Processing of 2D and 3D coordinates*

176 The 2D trajectories extracted from DeepLabCut
177 went through different post-processing processes: (i)
178 using an iterative algorithm, outlier points, where

179 the speed of movement was above a defined thresh-
180 old, were deleted; (ii) using the score maps for
181 each point given by DeepLabCut, we took out all
182 coordinates for which the probability was below a
183 defined threshold (fixed at $p=0.8$); (iii) the trace
184 was smoothed using a Savitzky-Golay filter (with
185 parameters `window_length = 11`, `polyorder = 5`) from
186 the *scipy* library. For (i) and (ii), we used linear inter-
187 polation to infer the coordinates of deleted points,
188 with *numpy interp* function.

189 For the 2D and 3D coordinates from Hand-
190 GraphCNN, we reproduced step (i) and (iii) of the
191 post-processing algorithm.

192 *Analysis of bradykinesia protocols using 2D and* 193 *3D coordinates*

194 Using the 2D and 3D trajectories, we computed
195 different metrics, specific for each protocol in order
196 to evaluate bradykinesia symptoms as recommended
197 by the MDS-UPDRS III guidelines.

198 Finger tapping (using 2D and 3D hand-pose esti-
199 mation): distance between the thumb and index
200 fingers tips.

201 Hand movements (using 2D and 3D hand-pose
202 estimation): averaged distance between each finger
203 tip and the wrist point.

204 Pronation-supination movements of the hands
205 (using only 3D hand-pose estimation since we
206 need to measure angle): azimuthal angle from
207 spherical coordinates of the tip of the thumb (com-
208 puted with z-axis being the line between wrist and
209 start of the third finger).

210 Each parameter was evaluated for all timesteps,
211 smoothed using a Savitzky-Golay filter (with param-
212 eters `window_length = 9`, `polyorder = 3`) and then nor-
213 malized between 0 and 1. Speed (in each direction)
214 was also computed, smoothed using a Savitzky-
215 Golay filter (with parameters `window_length = 5`,
216 `polyorder = 3`) and normalized between -1 and 1 .

The temporal evolution of the parameters was peri-
odic because the protocols consisted of 10 repetitions
of each movement. We used an algorithm based on
scipy.signal.find_peaks function to detect the 10 rep-
etitions. Note that we restricted the analysis to the
9 repetitions that were clearly detectable and which
did not depend on the initial position of the hand,
since the aim was to study the whole dynamics of
each sweep. The frequency F of the repetitions was
measured. These 9 sweeps were analyzed individ-
ually and the following properties were computed:

(i) duration of the sweep (noted $T_{sweep}[i]$ for each sweep i); (ii) amplitude of the sweep (maximum minus minimum values) (noted A_{sweep}); (iii) speed (amplitude divided by duration, noted S_{sweep}). The three first/last sweeps properties were compared to study if the movement was slowed down or altered in any way during the protocol, i.e., we computed

$$\Delta X_{sweep} = \frac{\text{mean}(X_{sweep}[i], i = 7, 8, 9)}{\text{mean}(X_{sweep}[i], i = 1, 2, 3)} - 1$$

A fatigue parameter was also computed,

$$L = \min(A_{sweep}[i], i = 1, \dots, 9) / \max(A_{sweep}[i], i = 1, \dots, 9) - 1,$$

L represents the maximal change in amplitude during the whole protocol, i.e., all the repetitions from 1 to 9.

To compute the deviation from a periodic trajectory, we fitted the following function on each trace, using the *lmfit* package:

$$f(t) = A_c + t \times \Delta A_c - \frac{A_m + t \times A_m}{(1 + \exp(\gamma \sin((w_0 + t \times w_1) \times t + D) - c))^\alpha}$$

From these fitted parameters, we computed:

- The period variation, which represents the change in period computed from the fit parameters,

$$\Delta T_{fit} = \frac{w_0}{w_0 + T \times w_1} - 1$$

where T is the duration of the whole 9 sweeps.

- The amplitude variation during the 9 sweeps, which represents the change in amplitude computed from the fit parameters,

$$\Delta A_{fit} = \frac{\Delta A_m}{A_m} \times T$$

We used 7 parameters (F , ΔT_{sweep} , ΔA_{sweep} , ΔS_{sweep} , ΔT_{fit} , ΔA_{fit} , L) for correlation with MDS-UPDRS III scores.

Videos were discarded for three reasons: (i) the algorithms DeepLabCut or HandGraphCNN failed to perform hand-pose estimation, the automated analysis failed (ii) to detect 9 sweeps or (iii) do the fitting procedure. Overall, 67% (64/96) of the videos for the tapping finger, 83% (78/94) for the hand movements,

43% (35/82) for the pronation-supination movements were used for further analysis.

Impact of the measured parameters for MDS-UPDRS scoring using statistical learning algorithms

To evaluate the impact of the measured parameters for MDS-UPDRS scoring, three classical algorithms of machine learning were tested: (i) the linear regression, (ii) the decision tree with `max_depth=2` and (iii) the decision tree with `max_depth=3`, all from the *scikit-learn* toolbox. We trained these algorithms on the task of predicting the averaged MDS-UPDRS III score, from the 7 previously defined parameters, for each protocol.

The coefficient of determination after training was computed with correct labels. Concurrently, we estimated the distribution of coefficients of determination obtained when training the algorithm on shuffled MDS-UPDRS III scores: we computed the mean and standard deviation on 100 different shuffles, and approximated them with a normal distribution. Using this control distribution, we estimated the probability $p_{shuffle}$ that trained networks with shuffled scores had higher coefficients of determination than the one obtained with the correct MDS-UPDRS scores. We used $p_{shuffle}$ to study the significance of the predictions (* $p_{shuffle} < 0.05$, ** $p_{shuffle} < 0.005$, *** $p_{shuffle} < 0.0005$).

Individual correlations were computed between the averaged MDS-UPDRS III scores and the 7 different parameters computed in the previous section, using *scipy.stats* `linregress` and `spearmanr` functions. Slopes and p -values are extracted for each case. Results with $p < 0.05$ were considered statistically significant (* $p < 0.05$, ** $p < 0.005$, *** $p < 0.0005$).

RESULTS

Demographic features

Among the 36 parkinsonian patients, 34 patients had a confirmed PD diagnosis according to MDS clinical criteria, 2 of them having a genetic form [1] (Supplementary Table 1). Patients assessment was carried out without considering the last dopamine intake in order to record a panel of patients in a wide range of parkinsonian state. We also recruited 11 healthy individuals, without any known neurological condition, to assess bradykinesia in a priori non-parkinsonian subjects.

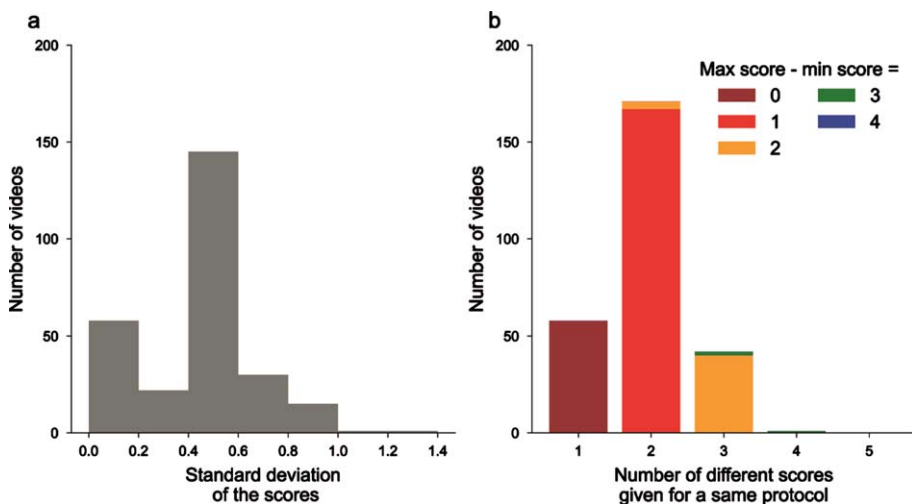


Fig. 1. Unreliability of MDS-UPDRS scores across neurologists' ratings. a) Histogram of the standard deviations (computed for each video, on the 5 different MDS-UPRS scores); b) videos ranked by the number of different MDS-UPDRS scores given by the physicians, colors for the difference between maximum and minimum scores.

Unreliability of MDS-UPDRS III scoring for bradykinesia

Five neurologists were asked to blindly score $N=272$ videos from PD and non-PD subjects performing with both hands the 3.4, 3.5, and 3.6 items of the MDS-UPDRS III. Comparing the different scores and analyzing their distribution for each video led to the conclusion that in the majority cases, the five examiners did not give the same scores. We also computed the standard deviation of the scores for each video and plotted its distribution (Fig. 1a). We observed that 58 videos had a null standard deviation, and as such were scored similarly by all five neurologists. When considering the whole dataset, MDS-UPDRS scores had a mean standard deviation of 0.409.

We also gathered videos as a function of the number of different scores that were given by the neurologists, and then in each class on the difference between the maximum and minimum scores (Fig. 1b). 58 videos had the same ratings, 171 with 2 different scores, 42 with 3 different scores and even 1 with 4 different scores given for the same videos (on a total of 5 possible scores from 0 to 4). Overall, this confirms the existence of inter-rater variability for MDS-UPDRS III scoring.

Finally, to assess the reproducibility of measurements between MDS-UPDRS III raters, we measured the ICC. The ICC score was 0.792 (95% confidence interval [0.76, 0.82]) which is defined as good between 0.75 and 0.9, but still showed that the assess-

sors do not exactly rate videos in the same manner. We did not observe difference between the variability of finger tapping, hand movements and pronation-supinations (0.836 [0.79, 0.88]; 0.774 [0.71, 0.83]; 0.763 [0.69, 0.83], respectively).

Extractions of relevant parameters for MDS-UPDRS III scores using deep learning

Using deep learning algorithms (DeepLabCut and HandGraphCNN), we detected 21 important points describing the hands of the patients, and extrapolated their coordinates in 2D and 3D as a function of time (see Materials and Methods). After complex steps of post-processing and analysis, one metric is extracted for each protocol and represented across time. Examples of extracted data in a single patient are presented in Figs. 2–4 for finger tapping, hand movements, and pronation-supination movements of the hands, respectively. For each figure, we present snapshots from the initial videos, with the different skeletons extracted using the two deep learning algorithms (Figs. 2a–4a, blue DeepLabCut 2D, red HandGraphCNN 2D, green HandGraphCNN 3D). From the trajectories of the hand joints, we computed protocol specific metrics (see Materials and Methods) during the whole protocol. As the protocol is composed of 10 repetitions of the same hand movement, for each metric 9 sweeps (delimited by the 10 plain circles on each trajectory, see Figs. 2b1–4b1, left) were detected and then analyzed with the evolution of movement during the protocols. The shapes from

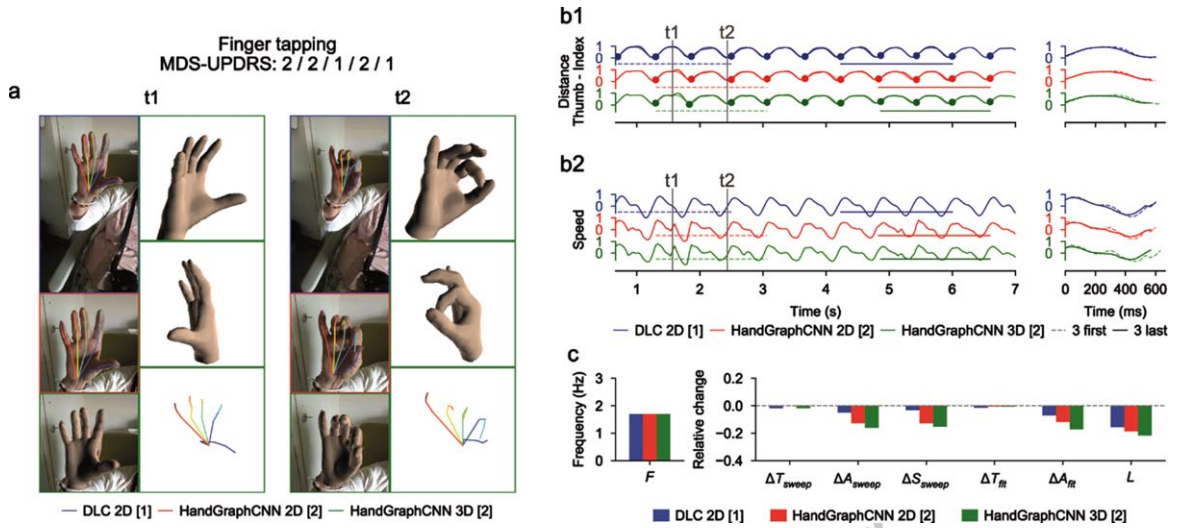


Fig. 2. Video analysis using deep learning algorithms for tapping protocol (associated Supplementary Movie 1). a) Snapshots from the initial videos, with the different skeletons extracted using the two deep learning algorithms (blue DeepLabCut 2D, red HandGraphCNN 2D, green HandGraphCNN 3D). b1, left) Evolution of the distance between the thumb and pinky fingers tips with 9 sweeps (delimited by the 10 plain circles on each trajectory). b1, right) First three sweeps (dotted lines) compared to the last three ones (plain lines). b2) same as (b1) for the speed. c) Frequency F , fatigue L , relative variation of the period ΔT_{sweep} , amplitude ΔA_{sweep} , speed ΔS_{sweep} across the protocol, more complex variation coefficients, ΔT_{fit} and ΔA_{fit} based on the fit of a periodic function.

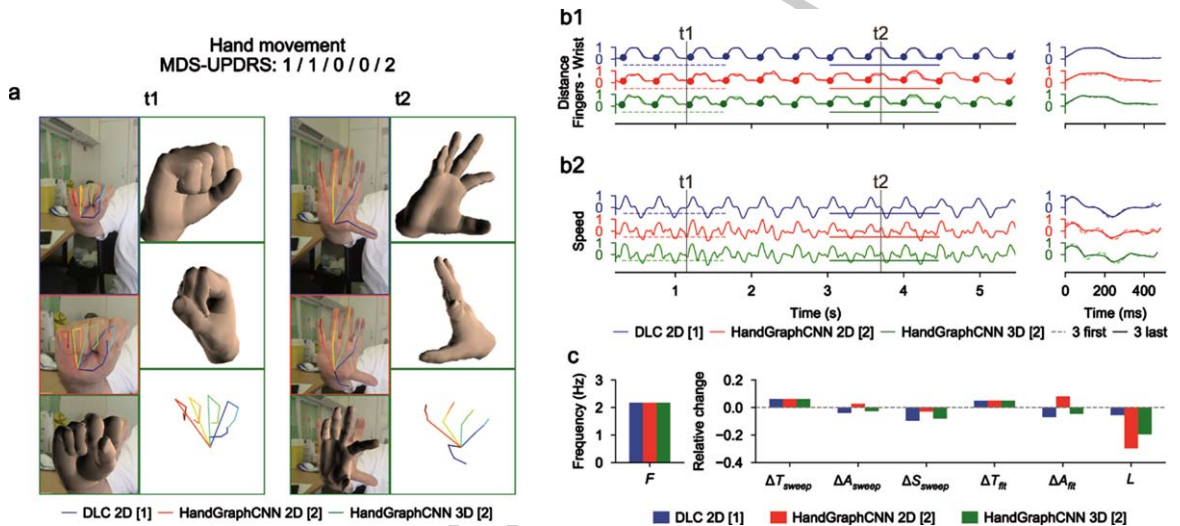


Fig. 3. Video analysis using deep learning algorithms for hand movements protocol (associated Supplementary Movie 2). a) Snapshots from the initial videos, with the different skeletons extracted using the two deep learning algorithms (blue DeepLabCut 2D, red HandGraphCNN 2D, green HandGraphCNN 3D). b1, left) Evolution of the averaged distance between each finger tip and the wrist point with 9 sweeps (delimited by the 10 plain circles on each trajectory). b1, right) First three sweeps (dotted lines) compared to the last three ones (plain lines). b2) same as (b1) for the speed. c) Frequency F , fatigue L , relative variation of the period ΔT_{sweep} , amplitude ΔA_{sweep} , speed ΔS_{sweep} across the protocol, more complex variation coefficients, ΔT_{fit} and ΔA_{fit} based on the fit of a periodic function.

344 the first three sweeps (dotted lines) were compared
345 to the last three ones (plain lines) (Figs. 2b1–4b1,
346 right). Similarly, we computed the associated speed
347 in Figs. 2b2–4b2. The frequency F , fatigue L , relative
348 variation of the period ΔT_{sweep} , amplitude ΔA_{sweep}

and speed ΔS_{sweep} across the protocol were com-
puted and presented in Figs. 2c/2c/4c. More complex
variation coefficients, ΔT_{fit} and ΔA_{fit} based on
the fit of a periodic function were also obtained
(Methods) and represented in Figs. 2c/2c/4c. This

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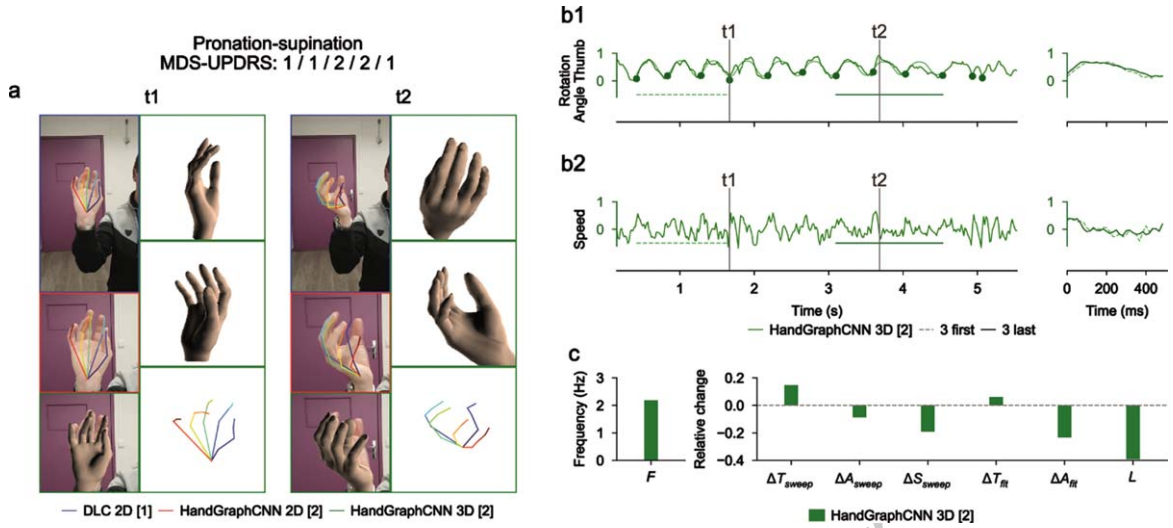


Fig. 4. Video analysis using deep learning algorithms for pronation-supination protocol (associated Supplementary Movie 3). a) Snapshots from the initial videos, with the different skeletons extracted using the two deep learning algorithms (blue DeepLabCut 2D, red HandGraphCNN 2D, green HandGraphCNN 3D). b1, left) Evolution of the azimuthal angle from spherical coordinates of the tip of the thumb with 9 sweeps (delimited by the 10 plain circles on each trajectory). b1, right) First three sweeps (dotted lines) compared to the last three ones (plain lines). b2) same as (b1) for the speed. c) Frequency F , fatigue L , relative variation of the period ΔT_{sweep} , amplitude ΔA_{sweep} , speed ΔS_{sweep} across the protocol, more complex variation coefficients, ΔT_{fit} and ΔA_{fit} based on the fit of a periodic function.

innovative analysis of several parameters over time allowed for accurate application of the MDS-UPDRS III item scoring guidelines.

For the finger tapping protocol (Fig. 2 and Supplementary Movie 1), we computed the distance between the tips of the thumb and the index fingers as the primary metric, and the associated speed. We observed on this example that the trajectories were almost periodic. The analysis showed that, with the three different detection algorithms (Fig. 2: DeepLabCut 2D (blue), HandGraphCNN 2D (red) and HandGraphCNN 3D (green)), the parkinsonian patient reduced the amplitude $\Delta A_{sweep}/\Delta A_{fit}$ of its movement during the protocol, and also reduced its speed ΔS_{sweep} . The period $\Delta T_{sweep}/\Delta T_{fit}$ stayed constant during the experiment. This agrees with the slightly positive MDS-UPDRS III scores (2/2/1/2/1) given by the neurologists.

For the hand movements (Fig. 3 and Supplementary Movie 2), we computed the mean distance between the finger tips and the wrist. We observed a periodic trajectory, with an increase of the period $\Delta T_{sweep}/\Delta T_{fit}$, and consequently a reduction of the speed ΔS_{sweep} . Since period is more difficult to assess than amplitude, it might explain why the raters gave so different scores (1/1/0/0/2).

For the pronation-supination movements of the hands (Fig. 4 and Supplementary Movie 3), an angle

was computed from the 3D representation. The trajectories were periodic enough to enable detection of the sweeps. Because of the irregularity of the trajectory, it is more accurate to use measures from fits to a periodic function. There was a drop of the amplitude of the movement ΔA_{fit} , consistent with the MDS-UPDRS III scores (1/1/2/2/1).

Overall, these three examples highlight the potential of the current analysis to accurately quantify hand movements over time during MDS-UPDRS III protocols.

Impact of the measured parameters on MDS-UPDRS using statistical analysis

We trained three algorithms (linear regression, decision tree with $\text{max_depth}=2$, decision tree with $\text{max_depth}=3$) to predict the averaged MDS-UPDRS III score from the previously defined 7 variables for each video. The results are presented in Table 1 with the coefficients of determination obtained for algorithms trained with correct MDS-UPDRS scores, the averaged coefficients of determination over 100 different randomly shuffled datasets and the $p_{shuffle}$ value used for significance (see Methods for its definition). For the finger tapping and hand movements protocols, all 3 algorithms predicted significantly better the correct MDS-UPDRS III score than algo-

Table 1

Regression algorithms trained to match MDS-UPDRS scores based on the parameters detected by the automated analysis

	Finger tapping (DeepLabCut 2D)	Hand movements (DeepLabCut 2D)	Pronation-supination (HandGraphCNN 3D)
Linear regression	0.309 (shuffled: 0.115) $p_{shuffle} = 4.23e-04^{***}$	0.427 (shuffled: 0.087) $p_{shuffle} = 4.83e-11^{***}$	0.286 (shuffled: 0.194) $p_{shuffle} = 1.27e-01$ n.s
Decision tree with max_depth = 2	0.461 (shuffled: 0.265) $p_{shuffle} = 6.65e-04^{**}$	0.598 (shuffled: 0.252) $p_{shuffle} = 2.72e-09^{***}$	0.377 (shuffled: 0.405) $p_{shuffle} = 6.20e-01$ n.s
Decision tree with max_depth = 3	0.609 (shuffled: 0.403) $p_{shuffle} = 1.76e-02^*$	0.701 (shuffled: 0.403) $p_{shuffle} = 3.95e-04^{***}$	0.630 (shuffled: 0.602) $p_{shuffle} = 4.05e-01$ n.s

Coefficients of determination are presented for training with the correct scores, the averaged over 100 trainings with shuffled scores, and the probability $p_{shuffle}$ used to test significance (see Materials and Methods).

408 rithms trained with shuffled scores (Table 1). For
409 the pronation-supination movements of the hands pro-
410 tocol, algorithms failed to predict the correct scores
411 since algorithms trained with the correct scores
412 performed as well as the ones trained with shuf-
413 fled scores ($p_{shuffle} > 0.1$). Using more complex
414 algorithms (Linear regression < Decision tree with
415 max_depth = 2 < Decision tree with max_depth = 3)
416 led to higher coefficient of determinations for
417 each protocol, but to decreases in significance
418 (Table 1).

419 In conclusion, the different parameters computed
420 using our automated analysis include pertinent infor-
421 mation for 3.4 and 3.5 MDS-UPDRS scoring.

422 Individual correlations with MDS-UPDRS III

423 Figure 5 showed correlations of the measured
424 parameters with the averaged MDS-UPDRS III
425 scores. For the finger tapping protocol (top), metrics
426 extracted from the 2D DeepLabCut coordinates are
427 shown here and their correlation with MDS-UPDRS
428 III scores is presented. The variation of amplitude
429 (for both empirical measure and fit), speed and fatigue
430 are significantly correlated with the MDS-UPDRS III
431 scores, with a negative slope, as expected from MDS-
432 UPDRS III guidelines. The other metrics (frequency
433 and period variations) are not significantly correlated.
434 Similar results are observed for the hand movements
435 protocol (center), with higher significant correlations
436 for amplitude (both types of measure), speed and
437 fatigue. Moreover, the period variation measured with
438 the fit is also negatively and significantly correlated.
439 This analysis did not reach significance for pronation-
440 supination movements of the hands (bottom). Over-
441 all, measured parameters related to MDS-UPDRS
442 guidelines are directly correlated with MDS-UPDRS
443 scores.

444 DISCUSSION

445 In this study, we created an innovative tool able
446 to reproduce clinical observations obtained with
447 MDS-UPDRS scores with a precise and objective
448 quantification and an adequate evaluation of the
449 MDS-UPDRS III scores for finger tapping and hand
450 movements. We obtained an accurate extraction of
451 important temporal parameters of MDS-UPDRS III
452 hand tasks, in an automated way only starting from
453 standard videos.

454 The 7 parameters tested here, exploring speed,
455 frequency, amplitude, duration and fatigue were mea-
456 sured using the automated analysis for finger tapping
457 and hand movements protocols, estimated the MDS-
458 UPDRS III used worldwide for PD motor symptoms
459 evaluation with high coefficients of determination.
460 More importantly, predictions using different statisti-
461 cal learning regression algorithms were significantly
462 higher than algorithms trained with shuffled datasets.
463 We concluded that the 7 parameters computed in the
464 analysis presented here contained enough informa-
465 tion for MDS-UPDRS estimation for finger tapping
466 and hand movement protocols. Hand movements
467 probably gave better correlations because we com-
468 puted the averaged distance for all fingers, compared
469 to the finger tapping metric which only relied on two
470 points, and therefore was more prone to undesired
471 variations. For pronation-supination movements of
472 the hands, our methods did not lead to statistically
473 significant predictions. The non-significant correla-
474 tion of pronation-supination with MDS-UPDRS III
475 might be due to the vertical plane of video recording
476 and a horizontal one would be more appropriate and
477 should be considered for further studies. This might
478 also be explained by the low number of samples to
479 reach a significant response. Indeed, less than half of
480 the videos with pronation-supination movements of
481 the hands passed all the tests to be included in the

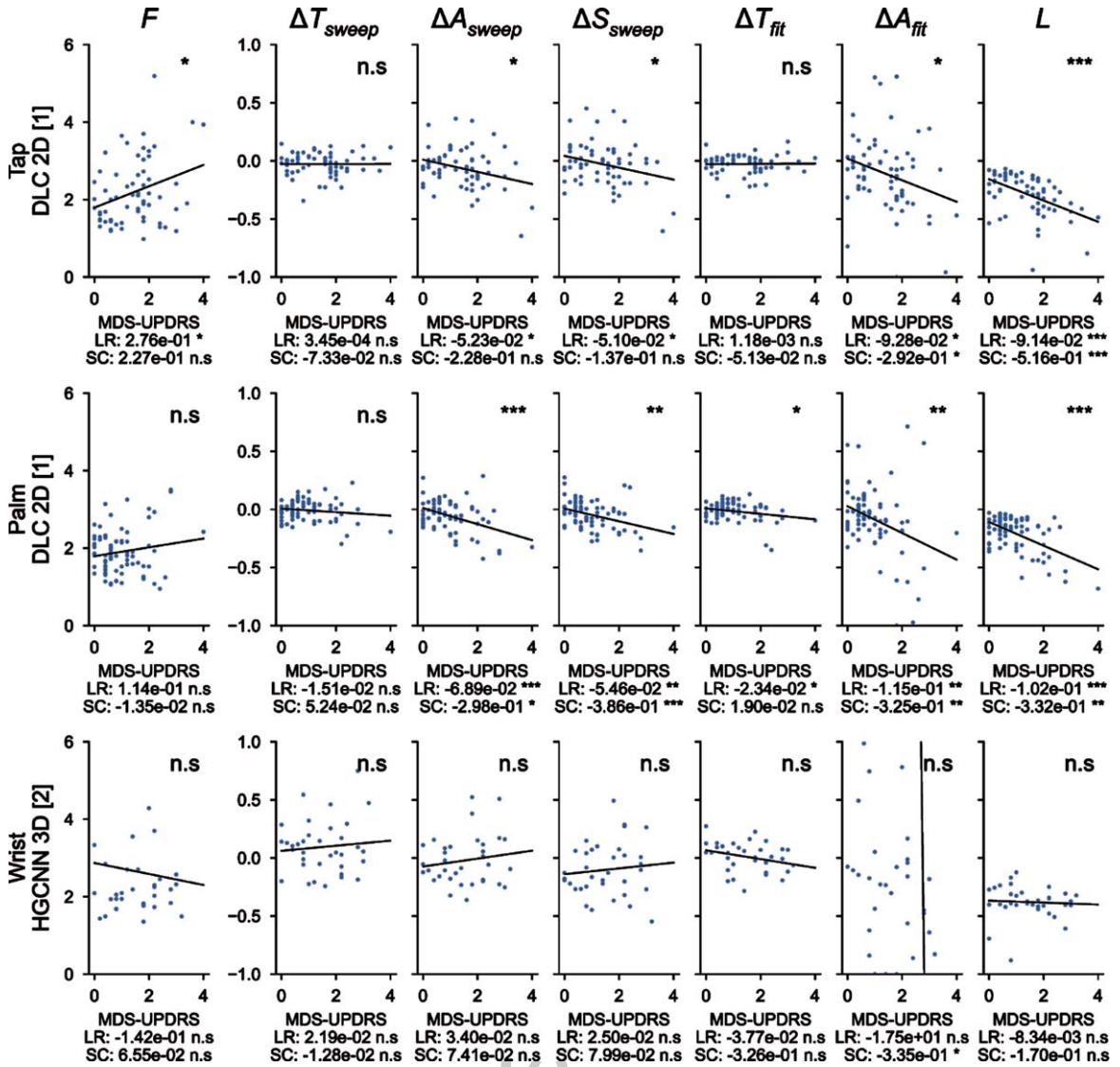


Fig. 5. Correlations between the 7 metrics computed with deep learning algorithms and averaged MDS-UPDRS scores. F , frequency of the repetitions; ΔT_{sweep} , comparison of the duration of the three first/last sweeps; ΔA_{sweep} , comparison of the amplitude of the three first/last sweeps; ΔS_{sweep} , comparison of the speed of the three first/last sweeps; ΔT_{fit} , period variation, which represents the change in period computed from the fit parameters; ΔA_{fit} , amplitude variation, which represents the change in amplitude computed from the fit parameters; L , fatigue parameter which represents the maximal change in amplitude during the whole protocol. ns: not significant; * <0.05 , ** <0.005 , *** <0.0005 for the significance of the p -values. LR, linear regression; SC, Spearman correlation.

482 prediction analysis. Moreover, we computed individ-
 483 ual correlations for the three protocols, and showed
 484 that the variation of amplitudes, speed and fatigue
 485 were significantly correlated with the MDS-UPDRS
 486 III scores. Importantly, the parameters measured in
 487 this analysis were consistent with MDS-UPDRS III
 488 guidelines for scoring. For instance, despite some
 489 studies analyzed more than 10 repetitions [20–22],
 490 we limited the repetitions to 10 because we wanted
 491 to strictly follow the MDS-UPDRS III recommenda-

492 tions. The 2D and 3D models of the hand recorded
 493 by video allow precise quantification of multiple
 494 parameters such as speed, amplitude and rate of the
 495 movement during MDS-UPDRS III evaluation. Thus,
 496 the composite parameters analyzed here are the same
 497 as those used for assessing bradykinesia during a
 498 medical consultation. Bradykinesia is a complex phe-
 499 nomenon of slowness of movement that cannot be
 500 seen only as a simple decrease of the movement rate
 501 [12], and therefore more complex properties need to

502 be considered in the scoring. The video-based method
503 described in our study by-passes the subjective mea-
504 surements of these parameters.

505 As already stated in the literature [7–12], we
506 also highlight here that MDS-UPDRS III scoring
507 is physician-dependent, and as such is a less reli-
508 able parameter than an automated and quantifiable
509 assessment tool to evaluate bradykinesia. Here, we
510 video-taped patients in real-life situations without
511 specific preparation and environment. Moreover,
512 patients were not selected regarding the phenotype or
513 the severity of the parkinsonian syndrome, thus pro-
514 viding a representative panel of real-life parkinsonian
515 patients. Importantly, by labelling only 5 frames per
516 videos we demonstrate that, even with scarce labels,
517 the network performs accurately. Overall, the time
518 needed to label 5 frames is quite small (1-2 minutes
519 for an expert) while the operation would last at least
520 an hour to individually label each frame, leading to a
521 great gain in term of time for a result almost as precise
522 as manual labelling.

523 In the expanding field of telemedicine, the present
524 tool appears of much interest. As an example,
525 COVID-19 pandemic has been striking evidence that
526 remote evaluation of neurological patients is needed,
527 this being even more crucial with chronic diseases
528 [30–32]. It has been shown that remote assessment
529 of MDS-UPDRS III is feasible via videoconferenc-
530 ing, except for specific items such as rigidity, or
531 postural instability, which are not as important as
532 bradykinesia for general evaluation of the disease [33,
533 34]. Indeed, some studies, which restricted bradyki-
534 nesia assessment to the upper limb motor tasks of
535 the MDS-UPDRS III, showed that upper limb motor
536 performance was a predictive feature of PD onset
537 and progression [21, 35]. Research on remote eval-
538 uation of PD patients has been mainly focused on
539 the use of technological devices [14, 15, 36, 37].
540 Such techniques require specific setup, which dif-
541 fers from MDS-UPDRS III tasks (e.g., touching
542 tactile screen of tapping, holding the phone during
543 pronation-supination). Recording video while clinically
544 assessing the MDS-UPDRS III is easy, and there-
545 fore any videos recorded, while strictly following
546 MDS-UPDRS III instructions, can be analyzed and
547 quantified by our system. Also, it is well known that
548 PD patients' symptoms, such as tremor and motor
549 functions, vary upon emotional load [13, 38]. Thus,
550 by quantifying bradykinesia from homemade videos,
551 in a less stressful environment than hospital or medi-
552 cal consultation, clinicians could have a more reliable
553 assessment of their patient's condition on a daily life

554 basis. It is noteworthy that this deep-learning analy-
555 sis of parkinsonian movement could be extended to
556 other body parts (e.g., feet movements, hypomimia,
557 posture) and therefore most of the MDS-UPDRS III
558 procedures [23, 24]. Thus, a wide range of movement
559 disorders, such as tremor or chorea, could be eligible
560 to this strategy of evaluation [16, 39].

561 This study has several limitations. First, it is a
562 monocentric study, which needs to be extended to
563 other centers. Nevertheless, our PD patient popula-
564 tion was similar to epidemiological data found in the
565 medical literature. Two of the 36 patients had atyp-
566 ical parkinsonian syndrome and were still included
567 since we were interested in analyzing parkinsonian
568 movements regardless of the underlying parkinson-
569 ian pathology. Indeed, although there are specific
570 scales for the assessment of degenerative parkinson-
571 ian syndromes (such as the UMSARS for MSA),
572 clinicians also apply in real life the MDS-UPDRS III
573 to atypical parkinsonian syndromes for the assess-
574 ment of the parkinsonian symptoms, especially in
575 outpatient clinic/consultation. Secondly, the MDS-
576 UPDRS III is considered as the international and
577 consensual scale which is a reference in the field
578 of parkinsonian symptoms. The main problem with
579 its use can arise from the assessor who rates the
580 MDS-UPDRS III as well as how the guidelines are
581 respected, often resulting in an inter- and intra-rater
582 variability. In our study, to reduce the risk of errors in
583 rating and to maximize the validity of the human rat-
584 ing, we have consciously asked several raters working
585 in a Movement disorders unit to rate the videos. We
586 demonstrated that it is feasible to objectively capture
587 and measure the parameters used in the guidelines
588 to rate the MDS-UPDRS III and then to get rid of
589 inter- and intra-rater variability. Thirdly, regarding
590 pronation-supination movements of the hands, the
591 results were not significant despite a trend for the
592 amplitude parameter. This could be explained by the
593 low number of patients but also the complexity of
594 extracting 3D coordinates. Frontal video recording
595 of the hand pronation-supination movements with
596 the forearm horizontal and not vertical would cer-
597 tainly facilitate the hand rotation analysis. Further
598 prospective analyses with more patients are needed
599 to implement this specific assessment. Forthcoming
600 development of this software will allow analysis of
601 other movement disorders and other body parts.

602 In conclusion, using a deep-learning approach, we
603 provided a quantitative measurement of bradykine-
604 sia that prevents inter-operator variability. We have
605 reached an unprecedented level of precision and sim-

plicity for its assessment. This precision, even at distance, could help non-movement disorder specialists to rate bradykinesia of their patients accurately. It would also be useful for specialists and non-specialists, to monitor bradykinesia of patients at distance, with video recordings provided directly by the patient or caregivers, with appropriate instructions as indicated in the MDS-UPDRS III.

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CONFLICT OF INTEREST

GV, CD, QS, MM, BG, LV declare no conflicts of interest. BD received research support from Orkyn, Merz-Pharma and Contrat de Recherche Clinique 2021 (CRC 2021). No sponsorship was obtained for this study.

SUPPLEMENTARY MATERIAL

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