To cite: van der Stouwe AMM.

Tuitert I, Giotis I, et al. Next

aided classification tool for

hyperkinetic movement

bmjopen-2021-055068

Prepublication history for

this paper is available online.

To view these files, please visit

the journal online (http://dx.doi.

org/10.1136/bmjopen-2021-

Accepted 28 September 2021

Check for updates

Received 06 July 2021

© Author(s) (or their

employer(s)) 2021. Re-use

¹Department of Neurology.

University Medical Centre

Groningen, Groningen, The

²Expertise Centre Movement

Medical Center Groningen,

Groningen, The Netherlands

³ZiuZ Visual Intelligence BV.

⁴Department of Information

University of Utrecht, Utrecht,

and Computing Sciences,

Gorredijk, Groningen, The

Disorders Groningen, University

Groningen, University of

permitted under CC BY-NC. No

commercial re-use. See rights

and permissions. Published by

055068).

BM.J.

Netherlands

Netherlands

disorders. BMJ Open

move in movement disorders

(NEMO): developing a computer-

2021;11:e055068. doi:10.1136/

BMJ Open Next move in movement disorders (NEMO): developing a computer-aided classification tool for hyperkinetic movement disorders

A. M. Madelein van der Stouwe ,^{1,2} Inge Tuitert,^{1,2} Ioannis Giotis,³ Joost Calon,³ Rahul Gannamani,^{1,2} Jelle R. Dalenberg ,^{1,2} Sterre van der Veen,^{1,2} Marrit R. Klamer,^{1,2,3} Alex C. Telea,⁴ Marina A.J. Tijssen^{1,2}

ABSTRACT

Introduction Our aim is to develop a novel approach to hyperkinetic movement disorder classification, that combines clinical information, electromyography, accelerometry and video in a computer-aided classification tool. We see this as the next step towards rapid and accurate phenotype classification, the cornerstone of both the diagnostic and treatment process.

Methods and analysis The Next Move in Movement Disorders (NEMO) study is a cross-sectional study at Expertise Centre Movement Disorders Groningen, University Medical Centre Groningen. It comprises patients with single and mixed phenotype movement disorders. Single phenotype groups will first include dystonia, myoclonus and tremor, and then chorea, tics, ataxia and spasticity. Mixed phenotypes are myoclonus-dystonia, dystonic tremor, myoclonus ataxia and jerky/tremulous functional movement disorders. Groups will contain 20 patients, or 40 healthy participants. The gold standard for inclusion consists of interobserver agreement on the phenotype among three independent clinical experts. Electromyography, accelerometry and three-dimensional video data will be recorded during performance of a set of movement tasks, chosen by a team of specialists to elicit movement disorders. These data will serve as input for the machine learning algorithm. Labels for supervised learning are provided by the expert-based classification, allowing the algorithm to learn to predict what the output label should be when given new input data. Methods using manually engineered features based on existing clinical knowledge will be used, as well as deep learning methods which can detect relevant and possibly new features. Finally, we will employ visual analytics to visualise how the classification algorithm arrives at its decision.

Ethics and dissemination Ethical approval has been obtained from the relevant local ethics committee. The NEMO study is designed to pioneer the application of machine learning of movement disorders. We expect to publish articles in multiple related fields of research and patients will be informed of important results via patient associations and press releases.

INTRODUCTION

The hyperkinetic movement disorders myoclonus, dystonia, tremor, chorea and tics are

Strengths and limitations of this study

- This cross-sectional study addresses a real need in classification of hyperkinetic movement disorders and may revolutionise our clinical approach.
- Representative patients from the entire hyperkinetic movement disorders spectrum will participate, including myoclonus, dystonia, tremor, chorea and tics, as well as ataxia, spasticity and functional movement disorders.
- An extensive data set will be collected and used for analysis: clinical information and data from electromyography, accelerometry and three-dimensional video.
- The project is a collaboration between movement disorders specialists at an acknowledged expertise centre and computer scientists in the corporate and academic world.
- Calculation of the needed sample size is hampered by the absence of previous studies in this exact field.

all characterised by excessive, involuntary movements (table 1).¹ In a patient presenting with such a disorder, it is crucial to classify the movement disorder phenotype promptly and accurately, because the phenotype classification will guide clinical decisions on diagnostic testing and (symptomatic) treatment.¹ Current clinical algorithms on hyperkinetic movement disorder diagnosis, whether aimed at dystonia,² myoclonus³ or tremor,⁴ all start with the same step: make sure to establish the main movement disorder phenotype in the patient. The subsequent diagnostic steps consist of additional tests such as laboratory, imaging and genetic tests, which can greatly aid the patient's aetiological diagnosis. The same holds for treatment: decisions on which medication to prescribe or which target to choose for deep brain stimulation are directed by the main clinical movement disorder phenotype.

The Netherlands
Correspondence to

Dr A. M. Madelein van der Stouwe; a.m.m.van.der.stouwe@umcg.nl

Table 1 Hyperkinetic movement disorders and related disorders		
Disorder	Clinical characteristics	
Dystonia	Persistent or intermittent muscle contractions that lead to abnormal movements and/or postures.	
Myoclonus	Sudden, brief, shock-like movements.	
Tremor	mor Rhythmic, oscillatory movement of a body part.	
Chorea	Continuous, abrupt, unpredictable non-rhythmic jerky movements.	
Tics	Stereotyped, repeating, shock-like movements, which can be temporarily repressed. Patients feel an urge preceding the movements.	
Ataxia	Inability to perform smooth, coordinated, targeted movements.	
Spasticity	Variable combination of paresis, elevated reflexes and increased muscle tone.	

Given the importance of clinical phenotype recognition, it is unfortunate that correct classification is difficult in many patients. While each phenotype of movement disorder has its own clinical presentation, complex and mixed forms occur. Moreover, the big five hyperkinetic phenotypes (myoclonus, dystonia, tremor, chorea and tics) may share overlapping clinical features with ataxia, spasticity and functional movement disorders. Presently, the classification of involuntary movements is based on clinical definitions and therefore on expert opinion. However, research demonstrated large interobserver and intraobserver variability of clinical classification.^{5–8} This is a major problem that impairs correct phenotyping of patients which subsequently delays aetiological diagnosis, deliverance of tailored treatment and evaluation of treatment effects.

To improve the classification of hyperkinetic movement disorders we set-up the current study, Next Move in Movement Disorders (NEMO). Our aim is to combine electromyography (EMG), motion sensors and threedimensional (3D) video with machine learning to develop a computer-aided classification tool for hyperkinetic movement disorders, which will help healthcare professionals establish the movement disorders phenotype. The algorithm of this computer-aided classification tool will be trained with the input data from patients who have a hyperkinetic movement disorders phenotype on which a panel of blinded independent experts agree.

Currently, EMG and accelerometry are applied in clinical practice to support the clinical classification of movement disorders, particularly myoclonus, tremor and dystonia. This is done by assessment of features such as muscle activation pattern, burst duration and frequency. While not all applications are supported by extensive evidence, there is a long tradition and ample experience with these methods.⁹ Despite the frequently incorporated clinical application of video to discuss cases among panels of experts, the use of video to automatically detect, quantify and classify involuntary movements is still in its infancy. Some advances have recently been made in using video for assessment of tremor frequency in a research setting.¹⁰ While these first steps are encouraging, application in other movement disorders is still lacking. Moreover, until now, investigations of machine learning

in movement disorders have mainly focused on quantification of the hyperkinetic movements¹¹ and optimisation of adaptive deep brain stimulation.¹² Efforts to classify different types of hyperkinetic movement disorders have not yet been made, to our knowledge.

In the NEMO project, we combine EMG, accelerometry and video recordings to develop an innovative new approach to movement disorder classification. We see the development of a computer-aided classification tool as the next step towards swift and accurate phenotype classification of hyperkinetic movement disorders, the cornerstone of both the diagnostic and treatment process.

METHODS Study population

The NEMO study protocol comprises patients with both single and mixed phenotype movement disorders. Single phenotype groups will primarily include dystonia, myoclonus and tremor, and in the future chorea, tics, ataxia and spasticity will be added. Although spasticity is not typically classified as a movement disorder, this group is added for distinction from dystonia, as these two neurological conditions are sometimes confused with one another.⁶ Similarly, ataxia is not usually listed as a hyperkinetic movement disorder, but signs of ataxia can overlap with symptoms of hyperkinetic movement disorders, making it a relevant phenotype to include.¹³ Mixed phenotype groups will consist of myoclonus-dystonia, dystonic tremor, myoclonus ataxia and jerky/tremulous functional movement disorders. All patient groups will consist of 20 participants. In addition, a group of 40 healthy participants will be included. Patients will be carefully selected based on all clinical information (MRK, AMMvdS and MAJT): after participating in the experiment, their data will be reviewed by a panel of experts to assess their phenotype and whether they meet the gold standard for inclusion.

Participants will only be eligible for inclusion if they are at least 16 years old. Exclusion criteria consist of comorbid conditions that lead to impaired upper extremity function, and a silver allergy or implanted pacemaker due to incompatibility with the EMG device. Moreover, healthy participants cannot be first-degree relatives of patients with hyperkinetic movement disorders.

Recruitment

Patients will be selected from the University Medical Center Groningen (UMCG) hyperkinetic movement disorders database and recruited at the UMCG outpatient clinic. If patient inclusion using these two sources does not lead to the targeted number of patients, we will recruit at other hospitals and via patient associations. Healthy participants will also be recruited at the UMCG. Participants will receive written information about the study and will have the opportunity to ask the investigators questions beforehand. Prospective participants can also call on an independent neurologist, who is not involved in the NEMO study, if they have any remaining questions. All participants will give written informed consent before enrolment.

Study setting

Movement Disorders Groningen, based at the UMCG, is uniquely positioned for the conduction of the NEMO study, as we treat patients with a broad range of hyperkinetic movement disorders. Movement Disorders Groningen is acknowledged by the Dutch Federation of University Medical Centres (NFU) as an official centre of expertise, and it is part of the European Reference Network for rare diseases. Data analysis will be performed in collaboration with ZiuZ Visual Intelligence BV (IG and JC), and the departments of computing science at the University of Utrecht (ACT) and the University of Groningen.

Experimental set-up

Clinical information

Clinical information such as age at onset, family history, disease progression, medication use, symptom influencing factors and psychological reports will be acquired from the participants during their study visit. Moreover, the influence of their movement disorder on activities of daily life will be assessed using the Fahn-Tolosa-Marin Essential Tremor Rating Scale part C,¹⁴ with additional questions that originate from the Burke-Fahn-Marsden Dystonia Rating Scale,¹⁵ SARA Ataxia Scale,¹⁶ Unified Myoclonus Rating Scale,¹⁷ Shapiro Tourette's Syndrome Severity Scale¹⁸ and Abnormal Involuntary Movement Scale.¹⁹ Additionally, patients will be asked to indicate the perceived severity of their movement disorder on a Visual Analogue Scale.

Movement tasks

Data will be collected while participants are performing several movement tasks. These tasks are selected from the Fahn-Tolosa-Marin Essential Tremor Rating Scale,¹⁴ Burke-Fahn Marsden Dystonia Rating Scale,¹⁵ SARA Ataxia Scale,¹⁶ Unified Myoclonus Rating Scale,¹⁷ Shapiro Tourette's Syndrome Severity Scale¹⁸ and Abnormal Involuntary Movement Scale.¹⁹ All of these are validated scales, used both in clinical practice and in experimental settings to assess the severity of specific movement disorders. As such, they contain tasks that are designed to elicit the movement disorder in a patient, enabling assessment of severity. Selection of tasks was discussed in depth with seven movement disorders, neuropaediatric and neurorehabilitation specialists with particular experience in the field of dystonia, myoclonus, tremor, chorea, tics, ataxia or spasticity to ensure comprehensive coverage of all disorders. Table 2 lists the set of tasks to be included. A trained investigator will instruct the participant before the recording of each task. If the task is not performed as instructed or the data collection was suboptimal, the patient is asked to repeat the task. Version control of tasks is used to ensure that the correct version of the task is included in the data analysis.

Data collection

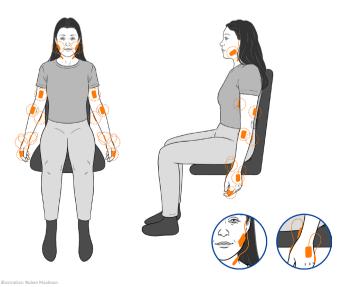
Data from EMG, motion sensors and 3D video will be collected simultaneously using custom made software (Visual Studio, Microsoft, USA). Surface EMG and motion sensor data will be recorded using the Trigno system (Delsys, Massachusetts, USA). Data collection frequencies of the EMG and the motion sensors are 2000 Hz and 150Hz, respectively. The Trigno Avanti sensors measure muscle activity, 3D accelerometry, 3D gyroscope and 3D magnetometry in each single wireless device. Ten Trigno Avanti sensors will be placed bilaterally on the participant's arms and neck, aiming for the musculus (m.) biceps, m. triceps, m. extensor carpi radialis, m. flexor carpi ulnaris and on both index fingers for accelerometry purposes only. Additionally, the Trigno Mini sensors measure muscle activity, using the sensor head and 3D accelerometry, using the sensor base, in each single wireless device. The six Trigno Mini sensor heads, registering EMG data, will be placed bilaterally aiming for the m. abductor pollicis, m. abductor digiti quinti and m. sternocleidomastoideus. The six Trigno Mini sensor bases, registering accelerometry, will be placed bilaterally at the inside of the forearm near the wrist, the back of the hand and the jaw. Figure 1 illustrates the placement of the EMG and motion sensors.

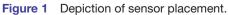
3D video data will be recorded using two types of 3D video cameras. A Realsense D435 camera (frame rate 30 fps; Intel, California, USA) will be used for whole body tracking; two Leap Motion cameras (frame rate 120 fps; Leap Motion, California, USA) will be employed to track the hands specifically. For the Realsense D435 camera, the Nuitrack API (3DiVi, California, USA) is used to extract joints from the depth frames. Aside from the depth frames, the Realsense D435 camera also collects two-dimensional (2D) colour frames at 30 fps. The Leap Motion camera uses an internal algorithm to extract joints from the depth frames. Figure 2 depicts which body parts are tracked by the cameras.

The EMG, motion sensors and 3D video data all have different sampling frequencies or frame rates. To enable synchronisation of modalities, a time stamp of each data sample point is saved during data acquisition.

Table 2 Included tasks			
Type of task	Task description	Duration/repetitions	
Rest	Arms at rest, in a relaxed position on the participant's lap and supinated hands	20-30s	
Postural tasks	Supinated outstretched arms and hands	20-30s	
	Pronated outstretched arms and relaxed wrists	20-30s	
	Pronated outstretched arms and hands	20-30s	
	Pronated outstretched arms and extended wrists	20-30s	
	Hands in front of chest: shoulders abducted at 90 degrees, fingers pointing towards each other	20-30s	
	Index finger in front of nose*	20-30s	
Dexterity	Spiral drawing*	1 Archimedes spiral	
	Writing*	1 sentence	
	Nine-hole pegboard test*	1 test	
	Diadochokinesis	20-30s	
Finger tapping	Finger tapping task: metronome cue* (3 and 5 Hz)	20s	
	Finger tapping task: self-paced*	20-30s	
	Four-finger tapping task*	20-30s	
Pointing tasks	Simple finger-to-nose manoeuvre*	5 repetitions	
	Point-to-nose test*	12 repetitions	
	Point-to-point test*	12 repetitions	
Mental distraction	Pronated outstretched arms and hands+'serial sevens' (100-7-7 etc)	20-30s	
Suppression	Pronated outstretched arms and hands+request to suppress all other movements	20-30s	
Speech	Repetition, spontaneous sentences, sustained vowel phonation	Not applicable	
Drinking	Drinking from a cup*	5 repetitions	
Gait	Standing	20-30s	
	Walking	20-30s	
	Tandem gait	20-30s	

*All unilateral tasks are subsequently performed with the right and left hand.



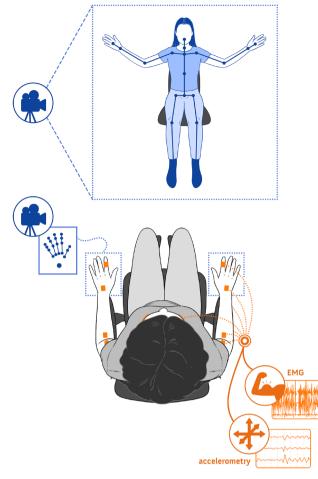


Sample size

For the current study, artificial intelligence experts expect that 20 subjects per patient group will be sufficient to develop proof-of-principle for a computer-aided diagnosis tool, because of the extensive amount of data that is collected per participant. Calculation of the needed sample size is hampered by the absence of previous studies in this exact field. While machine learning with, for example, MRI²⁰ and EEG²¹ data has previously been used to classify other diseases, both these data and disorders differ from the proposed study and are therefore not suitable to perform power analyses.

Data management

Study data will be handled confidentially and in accordance with the European personal data protection regulation (General Data Protection Regulation). A unique identification code will be assigned to each participant. All raw de-identified EMG, motion sensor and 3D video



illustraties: Ruben Maalma

Figure 2 Depiction of the study set-up and data collection. EMG, electromyography.

data will be stored at the research drive hosted and provided by the UMCG. Preprocessed data will be stored and analysed in the virtual research workspace, a secure virtualised desktop environment provided and hosted by the University of Groningen. De-identified clinical data are collected and managed using REDCap electronic data capture tools hosted at the UMCG.^{22 23}

Data analysis

Gold standard

The 'gold standard' for phenotype classification is defined as interobserver agreement among three experienced (inter)national clinical experts who will each assess the phenotype of a patient independently. Their evaluation will be based on the video recordings (RGB, 2D) of the participants performing all the different tasks described in table 2 during the experiment. Moreover, a sound fragment is recorded of the patient's speech for the experts' assessment. They will also be presented with information from the history and ancillary tests, to mimic the typical outpatient clinical situation as best as we can. The experts will be blind to the phenotype classification made by the investigators. All video data for the experts can be reviewed using the virtual research workspace and the relevant clinical information can be assessed in REDCap. Interobserver agreement among the clinical experts will be calculated using Fleiss' kappa (K).²⁴ We consider a Fleiss' K value above 0.80 (interpreted as excellent agreement) across all groups as sufficient. Moreover, for each individual participant, a minimum of two out of three experts should be in agreement. If a participant does not meet this requirement, their data will be excluded from further analysis.

Machine learning analyses

To develop the computer-aided classification tool, several machine learning techniques will be used. The clinical parameters, EMG, motion sensors and 3D video data will serve as input for the machine learning algorithm. The expert-based phenotype classification will serve as patient labels for supervised learning. In supervised machine learning, an algorithm is trained with multiple examples with a known output label (the expert-based phenotype classification), learning how to process the input data (clinical, EMG, motion sensor and video data) to reproduce the output label. The fully trained algorithm can then be given new input data, and is able to predict what the output label should be, making these types of algorithms well suited for classification.²⁵ Examples of machine learning techniques that will be investigated include deep learning methods such as long short-term memory networks²⁶ and methods based on manually engineered features such as random forest classifiers^{27 28} and learning vector quantisation^{29 30} using features from clinical neurophysiology research and practice (eg, frequency analysis or intermuscular coherence). From the above classes, deep learning methods have the advantage of computational scalability and the ability to automatically extract the relevant features for classification. In contrast, methods using manually engineered features allow one to control which aspects of the data are used by the classifier to reach its decisions and thus can be trained with fewer data to arrive at high accuracies. The information about how machine learning algorithms arrive at their output labels is often limited, which can mask confounding factors or simply diminish trust in the produced outputs. To address this problem, we will employ visual analytics to visualise how the classification algorithm arrives at its decision, and in particular which data aspects it uses to reach that decision.

ETHICS AND DISSEMINATION

The NEMO study protocol has been approved by the Medical Ethical Committee (METc 2018–444) of the UMCG and will be conducted according to the ethical standards of the Helsinki Declaration.

This study is designed to pioneer the application of machine learning in the classification of movement disorders, and allows for comprehensive data collection of clinical information, EMG and motion sensor data and 3D video. Because of its scope, we expect to publish multiple articles in the fields of clinical neurology, particularly movement disorders, clinical neurophysiology, artificial intelligence and visual analytics. Moreover, patients will be informed of important study results via the different patient associations, press releases, the website www.move mentdisordersgroningen.com and at the Movement Disorders Groningen biannual Patient Day.

Patient and public involvement

Members of the Movement Disorders Groningen Patient Council are updated regularly about the progress of the NEMO study and advise on online patient information. The help of several patient associations is enlisted for the recruitment of participants. Dissemination to patients and the general public is described above.

DISCUSSION

The NEMO study seeks to develop a computer-aided classification tool for hyperkinetic movement disorders. Such a tool would aid doctors in establishing the movement disorders phenotype. As adequate phenotype classification is the foundation of clinical decision-making and treatment of hyperkinetic movement disorders, improving this process would ultimately result in swifter and more accurate diagnoses and treatment for patients. To this end, we will investigate patients with myoclonus, dystonia, tremor, chorea, tics and functional jerky/tremulous movement disorders, as well as the related disorders ataxia and spasticity.

Our study has several strengths and some possible limitations. A major strength of this study is that it is conducted at a leading centre of expertise for movement disorders, Movement Disorders Groningen at the UMCG, The Netherlands. Experts in the field of all hyperkinetic movement disorders are present for collaboration and patient recruitment. For example, several experts assisted in selection of the appropriate tasks that patients will perform during the experiment, ensuring proper investigation of all relevant movement disorders. Also, the centre has a large patient database, which is a substantial advantage as some movement disorders are rare. Moreover, we have good rapport with our centre's patient council, who can advise on recruitment and dissemination of results. Disease rarity might still limit our speed of inclusion, a potential limitation, but given the advantages linked to our centre of expertise as well as the collaborations that exist with other centres, we are confident we will include enough patients in each group.

A second significant strength is the collection and analysis of extensive data: both clinical information and data from EMG, motion sensors and 3D video. In the analysis of these data, we can make use of the expertise of several members in our team. For instance, we have ample experience with the analysis of EMG and motion sensor data in movement disorders patients at the department of clinical neurophysiology.^{31–34} Similarly, ZiuZ Visual Intelligence BV, a major collaborator in this project, has a proven track

record of developing and deploying systems that collect and analyse visual data in different sectors and thus the required expertise for designing and implementing the computer-aided diagnosis tool as well as the data acquisition software for NEMO. In addition, academic input on the topic of machine learning and visual analytics is available from the University of Utrecht^{35–38} and the University of Groningen.^{39–42}

Significance

The aim of this study is to move the classification of movement disorder phenotypes forward, to improve the diagnostic process and optimise treatment in patients with hyperkinetic movement disorders. This could be achieved by the development of an all-encompassing computer-aided diagnostic tool for all hyperkinetic movement disorder phenotypes, which is the ultimate goal. Such a tool would be a game changer, helping neurologists to establish the phenotype in their patients with more certainty, so they can swiftly move on to other investigations into aetiology and to start the correct treatment.

In the process, it is likely that several new diagnostic biomarkers for hyperkinetic movement disorders will be discovered. These biomarkers can have their bearing on the current clinical practice quite quickly, particularly if they are in the fields of ECG or accelerometry because these techniques are already widely applied in movement disorders diagnostics.

Additionally, the tool that is being developed in the current study could be used to monitor disease severity and progression. This is of relevance for the monitoring of patients during treatment, for instance before and after deep brain stimulation. Another application could be patient assessment in clinical trials.

Study status

Data collection is in progress.

Twitter A. M. Madelein van der Stouwe @maddyvdstouwe

Contributors Conception of the study: MAJdKT and ACT. Design of the experiment: AMMvdS, IT, IG, JC, RG, SvdV, MRK, ACT and MAJdKT. Writing of the article: AMMvdS and IT. Review and critique: IG, JC, RG, JD, SvdV, MRK, ACT and MAJdKT.

Funding This work was supported by the European Fund for Regional Development of the European Union in collaboration with the province of Fryslan (grant number 01492947), and a ZonMW TOP Grant (grant number 91218013).

Competing interests Authors IG, JC and MRK are employed by ZiuZ Visual Intelligence BV, a company with an interest in the development of software tools to aid medical care. Presently, there is no commercial stake in this project.

Patient and public involvement Patients and/or the public were involved in the design, or conduct, or reporting, or dissemination plans of this research. Refer to the Methods section for further details.

Patient consent for publication Not applicable.

Provenance and peer review Not commissioned; peer reviewed for ethical and funding approval prior to submission.

Open access This is an open access article distributed in accordance with the Creative Commons Attribution Non Commercial (CC BY-NC 4.0) license, which permits others to distribute, remix, adapt, build upon this work non-commercially, and license their derivative works on different terms, provided the original work is properly cited, appropriate credit is given, any changes made indicated, and the use is non-commercial. See: http://creativecommons.org/licenses/by-nc/4.0/.

Open access

ORCID iDs

A. M. Madelein van der Stouwe http://orcid.org/0000-0001-9497-2019 Jelle R. Dalenberg http://orcid.org/0000-0001-8580-5358

REFERENCES

- Abdo WF, van de Warrenburg BPC, Burn DJ, et al. The clinical approach to movement disorders. Nat Rev Neurol 2010;6:29–37.
- 2 van Egmond ME, Kuiper A, Eggink H, et al. Dystonia in children and adolescents: a systematic review and a new diagnostic algorithm. J Neurol Neurosurg Psychiatry 2015;86:774–81.
- 3 Zutt R, van Egmond ME, Elting JW, et al. A novel diagnostic approach to patients with myoclonus. Nat Rev Neurol 2015;11:687–97.
- 4 van de Wardt J, van der Stouwe AMM, Dirkx M, et al. Systematic clinical approach for diagnosing upper limb tremor. J Neurol Neurosurg Psychiatry 2020;91:822–30.
- 5 Defazio G, Abbruzzese G, Livrea P, et al. Epidemiology of primary dystonia. Lancet Neurol 2004;3:673–8.
- 6 Eggink H, Kremer D, Brouwer OF, et al. Spasticity, dyskinesia and ataxia in cerebral palsy: are we sure we can differentiate them? Eur J Paediatr Neurol 2017;21:703–6.
- 7 Beghi E, Regio V, Papantonio A, et al. Reliability of clinical diagnosis of dystonia. *Neuroepidemiology* 2014;43:213–9.
- 8 van der Salm SMA, de Haan RJ, Cath DC, et al. The eye of the beholder: inter-rater agreement among experts on psychogenic jerky movement disorders. J Neurol Neurosurg Psychiatry 2013;84:742–7.
- 9 van der Veen S, Klamer MR, Elting JWJ, et al. The diagnostic value of clinical neurophysiology in hyperkinetic movement disorders: a systematic review. Parkinsonism Relat Disord 2021;89:176–85.
- Williams S, Fang H, Relton SD, et al. Accuracy of smartphone video for contactless measurement of hand tremor frequency. *Mov Disord Clin Pract* 2021;8:69–75.
- 11 Zhan A, Mohan S, Tarolli C, et al. Using Smartphones and machine learning to quantify parkinson disease severity: the mobile Parkinson disease score. JAMA Neurol 2018;75:876–80.
- 12 Habets JGV, Heijmans M, Kuijf ML, et al. An update on adaptive deep brain stimulation in parkinson's disease. *Mov Disord* 2018;33:1834–43.
- 13 Brandsma R, Verschuuren-Bemelmans CC, Amrom D, et al. A clinical diagnostic algorithm for early onset cerebellar ataxia. Eur J Paediatr Neurol 2019;23:692–706.
- 14 Fahn S, Tolosa E, Marin C. Clinical rating scale for tremor., 1988: vol 2, 271–80.
- 15 Burke RE, Fahn S, Marsden CD, et al. Validity and reliability of a rating scale for the primary torsion dystonias. *Neurology* 1985;35:73–7.
- 16 Schmitz-Hübsch T, du Montcel ST, Baliko L, et al. Scale for the assessment and rating of ataxia: development of a new clinical scale. *Neurology* 2006;66:1717–20.
- 17 Frucht SJ, Leurgans SE, Hallett M, *et al.* The unified myoclonus rating scale. *Adv Neurol* 2002;89:361–76.
- 18 Shapiro A, Shapiro E, Young J. Gilles de la tourette syndrome, 1988.
- 19 Guy W. ECDEU assessment manual for psychopharmacology, 1976.
- 20 Boudraa AO, Dehak SM, Zhu YM, et al. Automated segmentation of multiple sclerosis lesions in multispectral MR imaging using fuzzy clustering. Comput Biol Med 2000;30:23–40.

- 21 Aslan K, Bozdemir H, Şahin C, et al. A radial basis function neural network model for classification of epilepsy using EEG signals. J Med Syst 2008;32:403–8.
- 22 Harris PA, Taylor R, Thielke R, et al. Research electronic data capture (REDCap)--a metadata-driven methodology and workflow process for providing translational research informatics support. J Biomed Inform 2009;42:377–81.
- 23 Harris PA, Taylor R, Minor BL, *et al.* The redcap consortium: building an international community of software platform partners. *J Biomed Inform* 2019;95:103208.
- 24 Fleiss JL. Measuring nominal scale agreement among many raters. *Psychol Bull* 1971;76:378–82.
- 25 Auger SD, Jacobs BM, Dobson R. machine learning and artificial intelligence: a neurologist's guide. *Pract Neurol* 2020;21:4–11.
- 26 Hochreiter S, Schmidhuber J. Long short-term memory. Neural Comput 1997;9:1735–80.
- 27 Breiman L. Random forests. *Mach Learn* 2001;45:5–32.
- 28 Kulkarni VY. Random forest classifiers :a survey and future research directions. 36, 2013.
- 29 Schneider P, Biehl M, Hammer B. Adaptive relevance matrices in learning vector quantization. *Neural Comput* 2009;21:3532–61.
- 30 Biehl M. Biomedical applications of prototype based classifiers and relevance learning. In: *Lecture notes in computer science (including subseries lecture notes in artificial intelligence and lecture notes in bioinformatics*. . LNCS. Springer Verlag, 2017: Vol 10252. 3–23.
- 31 Zutt R, Elting JW, van Zijl JC, et al. Electrophysiologic testing AIDS diagnosis and subtyping of myoclonus. *Neurology* 2018;90:e647–57.
- 32 van der Stouwe AMM, Elting JW, van der Hoeven JH, *et al.* How typical are 'typical' tremor characteristics? sensitivity and specificity of five tremor phenomena. *Parkinsonism Relat Disord* 2016;30:23–8.
- 33 van der Stouwe AMM, Conway BA, Elting JW, et al. Usefulness of intermuscular coherence and cumulant analysis in the diagnosis of postural tremor. *Clin Neurophysiol* 2015;126:1564–9.
- 34 Kramer G, Van der Stouwe AMM, Maurits NM, et al. Wavelet coherence analysis: a new approach to distinguish organic and functional tremor types. *Clin Neurophysiol* 2018;129:13–20.
- 35 Schneider P, Biehl M, Hammer B. Distance learning in discriminative vector quantization. *Neural Comput* 2009;21:2942–69.
- 36 Bunte K, Schneider P, Hammer B, et al. Limited RANK matrix learning, discriminative dimension reduction and visualization. *Neural Netw* 2012;26:159–73.
- 37 Schneider P, Bunte K, Stiekema H, *et al.* Regularization in matrix relevance learning. *IEEE Trans Neural Netw* 2010;21:831–40.
- 38 Bunte K, Biehl M, Hammer B. A general framework for dimensionality-reducing data visualization mapping. *Neural Comput* 2012:24:771–804.
- 39 Rauber PE, Fadel SG, Falcao AX, et al. Visualizing the hidden activity of artificial neural networks. *IEEE Trans Vis Comput Graph* 2017;23:101–10.
- 40 Rauber PE, Falcão AX, Telea AC. Visualizing time-dependent data using dynamic t-SNE. *EuroVis* 2016.
- 41 Vernier EF, Comba JLD, Telea AC. Guided stable dynamic projections. *Computer Graphics Forum* 2021;40:87–98.
- 42 Espadoto M, Martins RM, Kerren A, *et al.* Toward a quantitative survey of dimension reduction techniques. *IEEE Trans Vis Comput* Graph 2021;27:2153–73.