

Individual Detection of Patients with Parkinson Disease using Support Vector Machine Analysis of Diffusion Tensor Imaging Data: Initial Results¹

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Introduction

Parkinson's Disease (PD) is the second most common neurodegenerative disorder after Alzheimer's disease. Nowadays, brain MR imaging is routinely performed as a diagnostic tool but its use is mostly limited to the exclusion of other pathologies.

Based on the assumption that PD is linked to systemic brain modifications and that some recent investigations in neurodegenerative disorders²⁻⁴ put in light a better sensitivity of white matter DTI tract-based spatial statistics (TBSS)⁵ analysis versus gray matter VBM⁶, we decided to perform a group level analysis as well as a pattern recognition analysis in order to detect PD patients among subjects with various forms of Parkinsonism at the individual level using support vector machine (SVM) technology.

Methods

- 40 consecutive patients with parkinsonism suggestive of PD who had (i) DTI at 3T without motion artefacts, (ii) brain ¹²³I-ioflupane SPECT (DaTScan) as reference, (iii) extensive neurological testing including followup 17 PD subjects (age range, 67.8 ± 6.7 years; 9 women) and 23 other subjects (age range, 67.2 ± 9.7 years; 7 women) and (iiii) the absence of morphologic findings on brain MR imaging.
- TBSS group level analysis.
- Individual-level analysis including SVM classification.

References

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Results

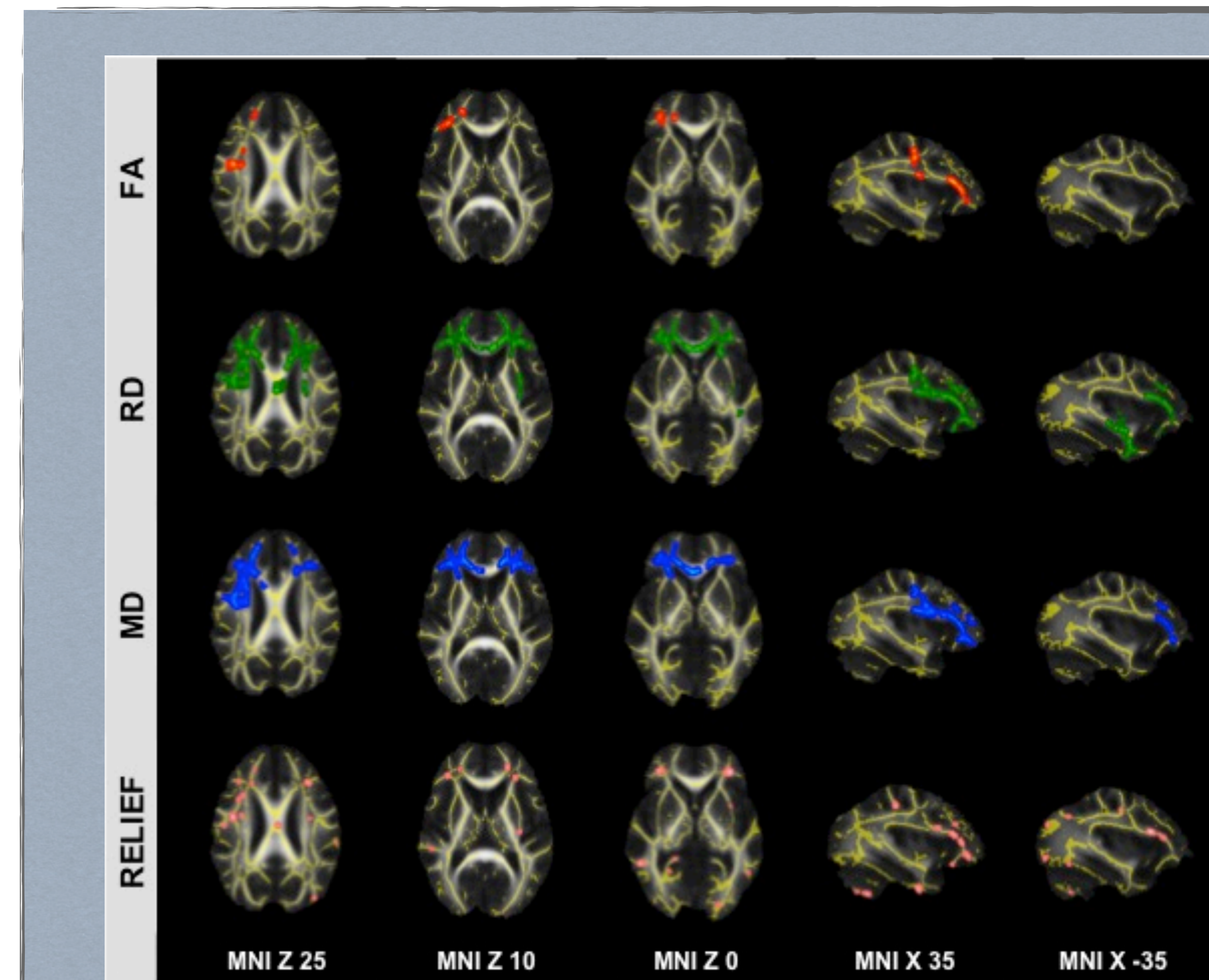


Figure 1: TBSS analysis in PD. Patients with idiopathic PD versus Other had a spatially consistent increase in FA (red) and a decrease in RD (green) and MD (blue) in a bilateral network predominantly in the right frontal white matter. The spatially consistent difference in LD was just below threshold (not illustrated). Gray indicates mean FA value; yellow, average skeleton. FA, RD, and MD clusters are TFCE corrected for multiple comparisons at $P < 0.05$. Supra-threshold voxels are enlarged by using TBSS fill (part of FSL) for illustrative purposes.

SVM Individual Classification Analysis

SVM analysis of FA provided a correct classification between PD versus Other with accuracies of up to 97.50 ± 7.54%. The spatial distribution of the most discriminative voxels (features) overlapped substantially with the results of the group-level TBSS analysis as illustrated in Fig 1 and Table 2.

TBSS Group Differences

The PD group compared with the Other group had a significant increase in fractional anisotropy (FA) and a corresponding significant decrease in radial diffusivity (RD) and mean diffusivity (MD), in particular in the right frontal white matter (Fig 1 and On-line Table). The level of significance of FA was slightly lower compared with RD and MD.

Longitudinal (LD) had spatially overlapping changes, which were just below threshold (not illustrated).

The inverse comparisons yielded no supra-threshold clusters.

Table 2: Individual SVM classification of PD based on DTI FA TBSS

17 PD, 23 Other: Chance Rate of Classification Accuracy, 57.5% (23/40)^a

No. of Features	Accuracy	TP Rate	FP Rate	TN Rate	FN Rate
100	97.50 (7.54)	0.94 (0.19)	0.00 (0.00)	1.00 (0.00)	0.06 (0.19)
250	95.50 (10.29)	0.90 (0.25)	0.00 (0.00)	1.00 (0.00)	0.11 (0.25)
500	96.25 (9.65)	0.91 (0.24)	0.00 (0.00)	1.00 (0.00)	0.09 (0.24)
750	97.25 (7.86)	0.94 (0.20)	0.00 (0.00)	1.00 (0.00)	0.07 (0.20)
1000	96.25 (9.65)	0.92 (0.23)	0.00 (0.00)	1.00 (0.00)	0.09 (0.23)

Note:—TP indicates true-positive; FP, false-positive; TN, true-negative; FN, false-negative.
^a Accuracy, TP, FP, TN, and FN rates for individual classifications using an SVM classifier with the indicated number of selected features for the individual classification of PD versus Other. Note that the accuracy is calculated as the average accuracy of 10 repetitions using 10-fold cross-validation (average and SD).

Conclusions

We propose the current study as the initial results that show the feasibility of performing SVM individual classification of DTI data in PD, which may merit future prospective and larger scale follow-up studies.

