

Summary of MEFOPA

Mendelian forms of Parkinson's Disease (PD) are a group of rare diseases which recapitulate many of the clinical, pathological and biochemical features of the common sporadic form of the disorder, and are therefore seen as model diseases which allow to study the underlying pathogenic molecular mechanisms and pathways. It is the understanding of PD pathways that is the prerequisite for the development of novel disease modifying and neuroprotective treatments. It is likely that those treatments will benefit all PD patients, as it becomes increasingly clear that the gene products and their interaction partners identified in rare Mendelian variants also play a crucial role in the sporadic disease.

Mendelian forms of PD can be divided into two groups: autosomal-dominant forms caused by mutations in the genes for α -synuclein (SNCA) and for leucine rich repeat kinase 2 (LRRK2), and autosomal-recessive forms caused by mutations in the genes for parkin (PRKN), Pten-induced kinase-1 (PINK-1), and the oncogene DJ1.

The mutations in the dominant genes are thought to cause PD by a gain-of-function mechanism. SNCA- as well as most cases of LRRK2-related PD are pathologically characterized by aggregates of α -synuclein ("Lewy-pathology"). It is therefore appropriate to assume that the pathogenic mechanisms of SNCA- and LRRK2-related PD interconnect. Research on the pathogenesis of those forms of Mendelian PD is coordinated in subproject 1.

Autosomal-recessive mutations in the genes for parkin, PINK-1 and DJ1 are believed to exert their pathogenic effect due to loss of some essential protective function. There is converging evidence suggesting that dysfunction of all recessive PD genes contribute to a state of increased cellular stress due to mitochondrial dysfunction and increased burden of radical oxygen species (ROS) and that at least two of the recessive PD genes, parkin and PINK1, operate within one single pathway. The molecular underpinnings of recessive forms of Mendelian PD will therefore also be investigated together, in subproject 2.

The findings in the basic science work-packages of the first two subprojects will directly feed into subproject 3, where a European registry and biobank for patients with rare Mendelian forms of PD will be established in order to advance investigations of biomarkers.

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Meetings

- 1) Preparatory meeting, Frankfurt, 05. October 2009
- 2) Kick-off meeting, Paris, 18. Mai 2010
- 3) Board meetings, (i) Tübingen, 19. November 2010, (ii) Amsterdam, 9. April 2011, Web conference, 18. November 2011.
- 4) Annual meeting and board meeting, Amsterdam, 8./9. April 2011

Results

Homepage

www.mefopa.eu

The screenshot shows the MEFOPA homepage with a navigation menu on the left and a central flowchart. The flowchart is titled 'European Project on Mendelian Forms of Parkinson's Disease' and is divided into two main sections: 'Sp1: Dominant PD' and 'Sp2: Recessive PD'. The 'Sp1: Dominant PD' section includes four steps: (i) α-SYN toxicity & aggregation, (ii) α-SYN oligomer formation, (iii) LRRK2 biology & pathology, and (iv) LRRK2 mouse & rat models. The 'Sp2: Recessive PD' section includes five steps: (i) parkin ubiquitin ligase activity, (ii) mito targets of parkin, (iii) modulators of PINK1/parkin, (iv) neuroprotective drugs, and (v) viral models of ARPD. A central box labeled 'Sp3: Biobank and Biomarkers' is connected to the flowchart. The flowchart also includes two boxes for 'Identification and validation of disease-related molecular pathways, drug-targets and biomarkers for disease susceptibility and progression'. The homepage also features a 'HOME' section with a welcome message and a 'News' section with a link to the 'MEFOPA Study Proposal Study Proposal Template'.

Publications out already:

- Winslow AR, Chen CW, Corrochano S, Acevedo-Arozena A, Gordon DE, Peden AA, Lichtenberg M, Menzies FM, Ravikumar B, Imarisio S, Brown S, O'Kane CJ, Rubinsztein DC. (2010) α-Synuclein impairs macroautophagy: implications for Parkinson's disease. **Journal of Cell Biology**. 190(6):1023-37. 20/09/2010
- Ashley R Winslow and David C Rubinsztein. (2011) The Parkinson disease protein α-synuclein inhibits autophagy. **Autophagy** 7(4): 429–431.
- Lichtenberg M, Mansilla A, Zecchini VR, Fleming A, Rubinsztein DC. (2011) The Parkinson's disease protein LRRK2 impairs proteasome substrate clearance without affecting proteasome catalytic activity. **Cell Death and Disease**. Aug 25;2:e196. doi: 10.1038/cddis.2011.81.

- Katerina Vamvaca et al. (2011) N-terminal deletion does not affect alpha-synuclein membrane binding, self-association and toxicity in human neuroblastoma cells, unlike yeast. **Journal of Neurochemistry**. 10.1111/j.1471-4159.2011.07431.
- Evangelia Emmanouilidou et al. (2011) Assessment of α -synuclein secretion in mouse and human brain parenchyma. **PLoS ONE** 6(7):e22225.
- Marina Moraitou et al. (2011) β -Glucocerebrosidase gene mutations in two cohorts of Greek patients with sporadic Parkinson's disease. **Molecular Genetics and Metabolism** 2011 Sep-Oct;104(1-2):149-52.
- Kathrin Brockmann et al. (2011) GBA-associated PD presents with nonmotor characteristics. **Neurology** 77(3):276-801.
- Van der Perren A., Toelen J., Coun F., Van den Haute C., Heeman B., Carlon M., Reumers V., Vandenberghe L.H., Wilson J.M., Debyser Z. and Baekelandt V. (2011) Efficient transduction of dopaminergic neurons in rat substantia nigra by different adeno-associated viral vector serotypes. **Gene Therapy** 18(5):517-27.
- Heeman B., Van den Haute C., Aelvoet S.-A., Valsecchi F., Rodenburg R., Reumers V., Debyser Z., Callewaert G., Koopman W.J.H., Willems P.H.G.M. and Baekelandt V. (2011) Depletion of PINK1 affects mitochondrial metabolism, calcium homeostasis and energy maintenance. **J of Cell Science** 124 (7):1115-25.
- Deleersnijder A., Desender L., Munck S., Pottel H., Buée L., Debyser Z., Baekelandt V. and Gerard M. (2011) Comparative analysis of different peptidyl-prolyl isomerases reveals FK506-binding protein 12 as the most potent enhancer of α -synuclein aggregation. **J. Biol. Chem** 286:26687-26701.
- Claudia Schulte et al. (2011) Genetic basis of Parkinson's disease: inheritance, penetrance, and expression. **The Application of Clinical Genetics** 2011;4; 67-80
- Thomas Gasser et al. (2011) Milestones in PD Genetics. **Movement Disorders** 6 26(6):1042-1048.
- Sorin Breit et al. (2010) Effective long-term subthalamic stimulation in PARK8 positive Parkinson's disease. **Journal of Neurology** 7: 1205-7.
- Thomas Gasser et al. (2010) Identifying PD-causing genes and genetic susceptibility factors: current approaches and future prospects. **Progress in brain research** 2010;183:3-20.
- Susanne Gräber et al. (2011) Self estimated quality of life in monogenetic Parkinson's disease. **Movement Disorders** 1: 182-190.
- Imam SZ et al. (2011) Novel regulation of parkin function through c-Abl-mediated tyrosine phosphorylation: implications for Parkinson's disease. **J Neurosci**. 31(1): 157-163.
- Springer W et al. (2011) Regulation of PINK1-Parkin-mediated mitophagy. **Autophagy** 7(3): 266 – 278.
- Geisler S et al. (2011) The PINK1/Parkin-mediated mitophagy is compromised by PD-associated mutations. **Autophagy** 6(7): 871 – 878.
- Kieper N et al. (2010) Modulation of mitochondrial function and morphology by interaction of Omi/HtrA2 with the mitochondrial fusion factor OPA1. **Experimental Cell Research** 316(7): 1213

Recruitment of patients after half of the project (18 months):

Entity	Number
All	268 (319)
SNCA	29
symptomatic	24
asymptomatic	5
LRRK2	128
symptomatic	87
asymptomatic	41
Parkin	63
homozygous	37
heterozygous	26
PINK1	48
homozygous	16
heterozygous	32
Other mutations and controls	51