PPMI Status Update

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Disclosure

- Co-founder on Molecular Neuroimaging LLC PET and SPECT imaging services
- Consultant –BMS, GEHC, Lilly, Merck, Navidea, Piramal
- Pfizer, Sanofi,





Parkinson Progression Marker Initiative

- Disease modifying PD therapeutics remain a major unmet need
- A major obstacle to current phase 2/3 neuroprotection studies is the lack of biomarkers for
 - Disease mechanism
 - Drug mechanism
 - Dosage determination
 - Study eligibility
 - Stratification into PD sub-types
 - Correlation with clinical signals

Requirements for Biomarker Infrastructure

Specific Data Set

- Appropriate population (early stage PD and controls)
- Clinical (motor/non-motor) and imaging data
- Corresponding biologic samples (DNA, blood, CSF)

Standardization

- Uniform collection of data and samples
- Uniform storage of data and samples
- Strict quality control/quality assurance

Access/Sharing

- Data available to research community → data mining, hypothesis generation & testing
- Samples available for studies



PPMI Study Synopsis

Study population	 400 de novo PD subjects (newly diagnosed and unmedicated) 200 age- and gender-matched healthy controls 70 SWEDD 100 Prodromal - Olfactory/RBD/LRRK2 500 LRRK2 - PD manifest and non-manifesting family members 100 Synuclein - PD manifest and non-manifesting family members Subjects will be followed for 3 to 5 years
Assessments/ Clinical data collection	 Motor assessments Neurobehavioral/cognitive testing Autonomic, Olfaction, Sleep DaTSCAN, VMAT, Amyloid imaging, DTI/RS MRI
Biologic collection/	 DNA collected at screening Serum and plasma collected at each visit; urine collected annually CSF collected at baseline, 6mo 12 mo and then annually Samples aliquotted and stored in central biorepository
Initial Verification studies	 Lead biologic candidates to be tested: Alpha-synuclein (CSF) DJ-1 (CSF and blood) Urate (blood) Abeta 1-42 (CSF) Total tau, Phospho-tau (p-181) (CSF)

PPMI Sites

PPMI SITES IN THE UNITED STATES:

- Arizona PD Consortium (Sun City, AZ)
- Baylor College of Medicine (Houston, TX)
- Boston University (Boston, MA)
- Cleveland Clinic (Cleveland, OH)
- Emory University (Atlanta, GA)
- Institute of Neurodegenerative Disorders (New Haven, CT)
- Johns Hopkins University (Baltimore, MD)
- Northwestern University (Chicago, IL)
- Oregon Health and Science University (Portland, OR)
- The Parkinson's Institute (Sunnyvale, CA)
- PD & Movement Disorders Center at Boca Raton (Boca Raton, FL)
- University of Alabama at Birmingham (Birmingham, AL)
- University of California at San Diego (San Diego, CA)
- University of Cincinnati (Cincinnati, OH)
- University of Pennsylvania (Philadelphia, PA)
- University of Rochester (Rochester, NY)
- University of South Florida (Tampa, FL)
- University of Washington (Seattle, WA)

PPMI SITES IN EUROPE:

- Imperial College (London, UK)
- Innsbruck University (Innsbruck, Austria)
- Paracelsus-Elena Clinic Kassel/University of Marburg (Kassel and Marburg, Germany)
- University of Napoli (Naples, Italy)
- University of Tübingen (Tübingen, Germany)
 PPMI SITES IN AUSTRALIA:
- Macquarie University (Sydney, Australia)

Sites to enroll LRRK2 and synuclein subjects will be added.





PPMI SC and Study Cores

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Steering Committee	PI-K Marek, C Tanner, T Foroud, D Jennings, K Kieburtz, W Poewe, B Mollenhauer, T Simuni, (core leaders, MJFF, ISAB), S Lasch
Clinical Coordination Core	 University of Rochester's Clinical Trials Coordination Center PI: Karl Kieburtz, irina Lazurenko, Alice Rudolph, Cindy Casaceli
Imaging Core	Institute for Neurodegenerative Disorders;PI: John Seibyl, Norbert Schuff,
Statistics Core	University of IowaPI: Chris Coffey
Bioinformatics Core	Laboratory of Neuroimaging (LONI) at UCLAPI: Arthur Toga, Karen Crawford
BioRepository	Coriell/BioRepPI: Alison Ansbach, Paola Casalin,
Bioanalytics Core	University of PennsylvaniaPI: John Trojanowski, Les Shaw
Genetics Core	National Institute on Aging/NIHPI: Andy Singleton
RBD Core	Hephata Hessisches Diakoniezentrum e. V.PI: Geert Mayer
Olfactory Core	Institute for Neurodegenerative DisordersPI: Danna Jennings
Genetics Coordinating Core	Indiana UniversityPI: Tatiana Foroud



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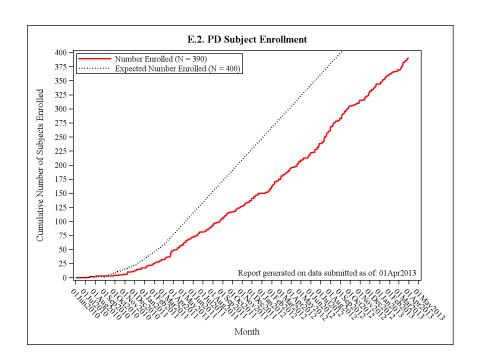


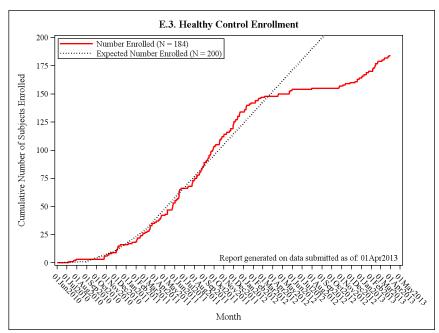






ENROLLMENT





- •Enrollment 419 PD 191 HS 59 SWEDD 669 subjects
- Retention 413 PD 183 HS 58 SWEDD 654 subjects



Baseline Demographics and Motor Characteristics					
Baseline Assessment	PD Subjects (N = 423)	Healthy Controls (N = 196)	SWEDD Subjects (N = 64)	PD p-value relative to HC	PD p-value relative to SWEDD
Mean Age (Range)	61.7 (33 - 85)	60.8 (31 - 84)	60.9 (38 - 79)	0.33	0.58
Gender (M %/F %)	277 (65%) / 146 (35%)	126 (64%) / 70 (36%)	40 (63%) / 24 (37%)	0.79	0.67
MDS-UPDRS Mean Score & Sub Scores					
MDS-UPDRS Total Score	32.3	4.7	29	< 0.01	0.08
MDS-UPDRS Part I	5.5	3	8.7	< 0.01	< 0.01
MDS-UPDRS Part II	5.9	0.4	5.9	< 0.01	0.98
MDS-UPDRS Part III (Motor Exam)	20.9	1.2	14.3	<0.01	<0.01
Hoehn & Yahr N(%)					
Stage 0	0 (0%)	184 (97%)	0 (0%)		
Stage 1	179 (43%)	2 (1%)	35 (59%)	<0.01	0.7
Stage 2	229 (56%)	0 (0%)	24 (41%)	<0.01	0.7
Stage 3-5	2 (1%)	0 (0%)	0 (0%)		
Modified Schwab & England (mean)	93.1	NA	94.7	NA	0.05
First degree family Member with PD (%)	54 (13%)	0 (0%)	14 (24%)	<0.01	0.22
Mean Duration of Disease (months)	6.6 (0.4 - 35.8)	NA	7.9 (0.5 - 37)	NA	0.16
Initial Symptoms*					
Resting Tremor	321 (78%)	NA	50 (85%)	NA	0.23
Rigidity	314 (76%)	NA	33 (56%)	NA	<0.01
Bradykinesia	339 (82%)	NA	46 (78%)	NA	0.42
Postural Instability	29 (7%)	NA	7 (12%)	NA	0.19
Other	72 (17%)	NA	8 (14%)	NA	0.45



Baseline Non-motor Characteristics					
Baseline Assessment	PD Subjects (N = 423)	Healthy Controls (N = 196)	SWEDD Subjects (N = 64)	PD p-value relative to HC	PD p- value relative to SWEDD
MOCA Total Score	27.1	28.2	27.1	< 0.01	0.94
SCOPA AUT Total Score	9.5	5.9	113.8	<0.01	<0.01
GDS	2.3	1.3	3.3	<0.01	<0.01
State Trait Anxiety Score	65.2	57	70.3	<0.01	0.04
QUIP	0.3	0.3	0.6	0.92	<0.01
Benton Judgment of Line Orientation Score	12.7	13.1	12.8	0.05	0.84
HVLT Immediate Recall	9.7	10.2	9.7	< 0.01	0.84
HVLT Delayed Recognition	11.2	11.5	10.8	< 0.01	0.07
HVLT Delayed False Alarms	1.2	1.1	1.7	0.2	0.02
Letter Number Sequencing Raw Score	10.5	11	9.8	0.07	0.05
Semantic Fluency Total Score	48.6	51.9	45	<0.01	0.03
Symbol Digit Modalities (SDM)	41.3	46.8	41	<0.01	0.83
UPSIT Raw Score	22.3	34	31.3	< 0.01	<0.01
Epworth Sleepiness Scale (ESS)					
Not Sleepy (9 or below)	345 (84%)	163 (88%)	40 (68%)	<0.01	<0.01
Sleepy (10 or above)	65 (16%)	23 (12%)	19 (32%)	<0.01	<0.01
REM Sleep Disorder					
Negative (< 5)	257 (62%)	152 (80%)	34 (58%)	<0.01	0.57
Positive (5 or greater)	157 (38%)	37 (20%)	25 (42%)	<0.01	0.37



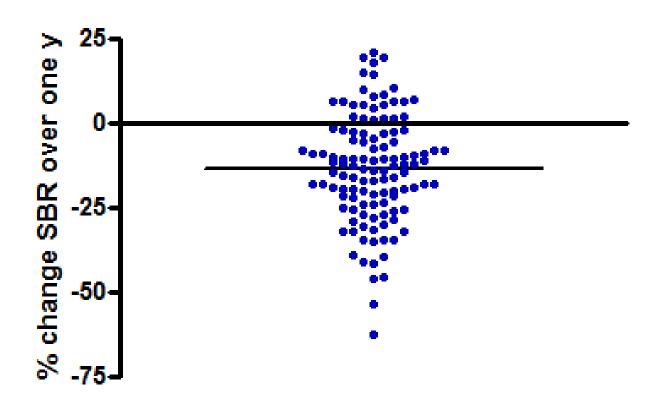


MoCA Cut-off Scores

MoCA	Frequency	Percentage	Cumulative Frequency	Cumulative Percent
17	1	0.26	1	0.26
19	1	0.26	2	0.52
20	2	0.52	4	1.04
21	5	1.30	9	2.34
22	8	2.08	17	4.43
23	13	3.39	30	7.81
24	13	3.39	43	11.20
25	36	9.38	79	20.57
26	49	12.76	128	33.33
27	64	16.67	192	50.00
28	68	17.71	260	67.71
29	70	18.23	330	85.94
30	54	14.06	384	100.00

Consistent with research reporting 15-20% of de novo PD patients have MCI.

Longitudinal DAT





N= 117 Mean 13.3% ± 16.0% 78.6% going down at yr 1

CSF Acquisition

_	Visit (months)					
Group	0 Baseline	6	12	24		
PD	401 (98%)	275 (91%)	171 (87%)	29 (83%)		
Healthy controls	184 (97%)	146 (87%)	140 (84%)	25 (80%)		
SWEDD	59 (92%)	36 (89%)	25 (84%)	N/A		

LP well tolerated – HA – 4-7%
CSF Volume collected 15.25 (mean)
Sprotte needle used in 82%
Syringe suction 63%
Sitting position in 63%
Flouroscopy in 5%

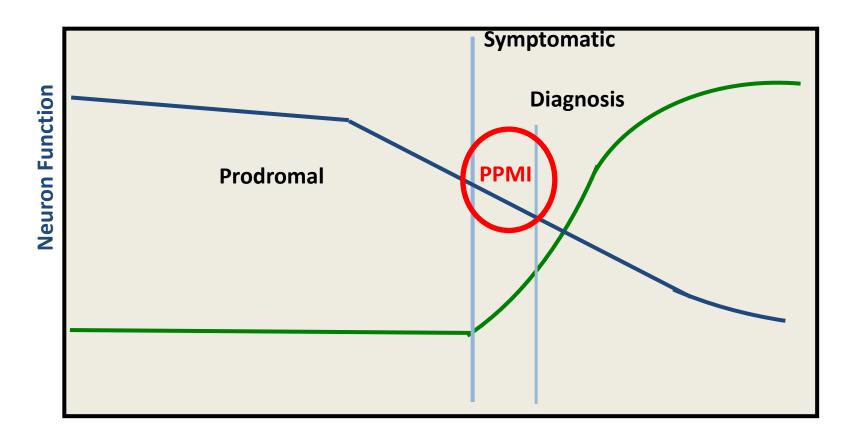
Play a Part in Parkinson's Research

CSF Pilot Baseline Data

	HC (N = 39)	PD (N = 63)	<i>P</i> value [#]	
Aβ ₁₋₄₂ (pg/mL)	242.8 ± 49.95	228.7 ± 45.63	0.0466	
	(226.7 – 259.0)*	(217.2 – 240.2)	0.0466	
t tou (ng/ml)	53.9 ± 19.33	46.1 ± 24.71	0.0276	
t-tau (pg/mL)	(47.6 – 60.1)	(39.8 - 52.3)	0.0276	
n-tau (ng/ml.)	24.9 ± 8.45	21.0 ± 7.83	0.0093	
p-tau ₁₈₁ (pg/mL)	(22.2 – 27.6)	(19.0 - 23.0)	0.0093	
t tou/AD ratio	0.240 ± 0.141	0.215 ± 0.157	0.0451	
t-tau/Aβ ₁₋₄₂ ratio	(0.195 – 0.286)	(0.176 – 0.255)	0.0451	
n tau IAO ratio	0.113 ± 0.075	0.099 ± 0.063	0.1482	
p-tau ₁₈₁ /Aβ ₁₋₄₂ ratio	(0.089 – 0.138)	(0.084 – 0.115)	0.1402	
p-tau ₁₈₁ /t-tau ratio	0.491 ± 0.160	0.543 ± 0.263	0.6820	
	(0.439 – 0.543)	(0.477 – 0.609)	0.0020	
α-syn (pg/mL)	1264 ± 425.7	1082 ± 611.1	0.0120	
	(1126 – 1403)	(928 – 1235)	0.0120	

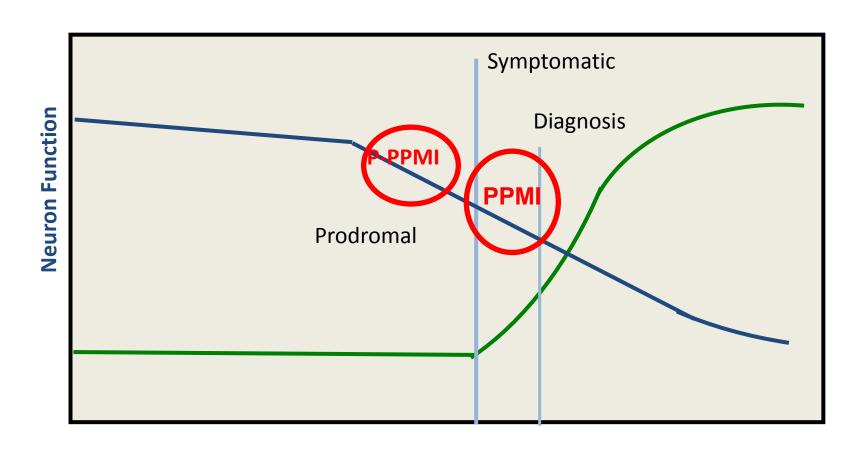
Clinical Ratings

Natural history of Parkinson's disease

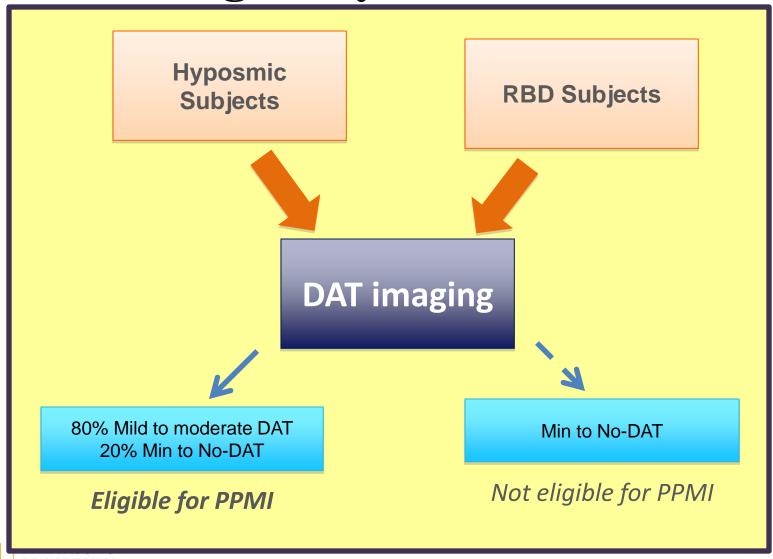




Natural History of Parkinson disease

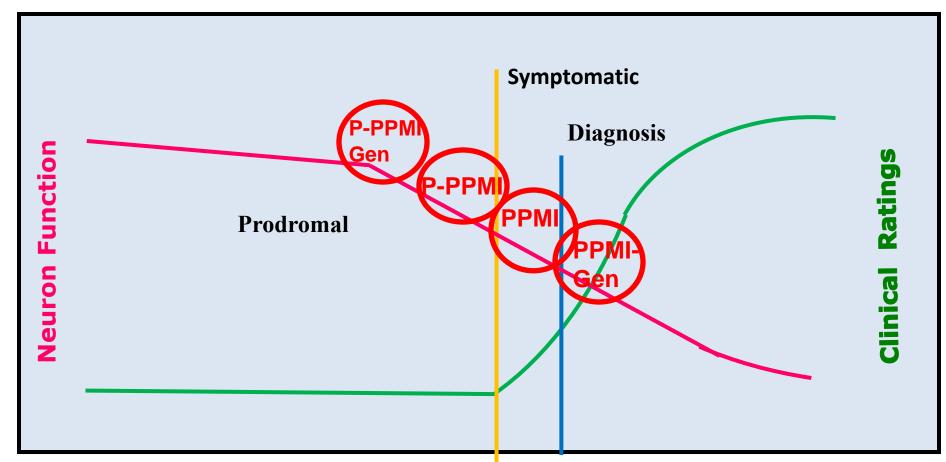


Eligibility for P-PPMI





PPMI - Cohorts Stage of PD



PPMI-LRRK2/SNCA

- Leverage existing PPMI infrastructure and add sites with existing expertise and experience with LRRK2 patients and families.
- Enroll 250-300 LRRK2/SNCA + PD and 250-300 LRKK2/SNCA + unaffected family members with and intensive longitudinal clinical assessment protocol.
- Follow PD and unaffected family members for for 3-5 years
 - Establish pre-motor biomarker signature
 - Define phenoconversion
- Maintain PPMI database structure and commitment to rapid access to data



Current Status

- PD, healthy and SWEDD cohorts and has established standardized procedures for acquisition and analysis of all study data
- PPMI strategy for comprehensive biomarker acquisition including CSF has been successful.
- PPMI longitudinal follow-up underway-subject retention 16/662 subjects withdrawn from the study
- Robust web-based access(<u>www.ppmi-info,org</u>) for data and biospecimen - >70000 data downloads >20 biologic specimen requested.
- PPMI Prodromal and Genetic cohorts incorporated to assess prodromal PD biomarkers

$PPMI \longleftrightarrow ADNI$

- Data Mining
- Imaging core Amyloid imaging, DAT imaging, Tau, Inflammatory
- CSF

 Tau, pTau, Amyloid, alphasynuclein
- Cognitive outcomes
- Genetics full sequence data
- Prodromal cohorts ethical issues

