

**Supplementary Figure 1. Characterisation of the rabbit monoclonal Rab8A-total (MJF-22-74-3) antibody.** (A & B) Wild-type MEFs treated  $\pm$  100 nM MLI-2 inhibitor for 60 min prior to cell lysis. For immunoprecipitation analysis 100  $\mu$ g of whole cell extract was subjected to immunoprecipitation employing either the Rab8A-total (MJF-22-74-3) antibody or Pre-immune IgG (used as a control). The immunoprecipitates were subjected to Rab8A-total (10% of sample) and Rab8-pT72 immunoblot (90% of sample) (all at 1  $\mu$ g/ml primary antibody concentration). 10  $\mu$ g of whole cell extract or supernatants after immunoprecipitation were subjected to Rab8A-total immunoblot. (C) The indicated lysates of A549 wild-type and A549 Rab8A knock-out cells (10  $\mu$ g), HEK293 cells (10  $\mu$ g), human PBMCs (5  $\mu$ g), wild-type MEFs (10  $\mu$ g), as well as mouse lung, spleen, kidney and brain lysates (20  $\mu$ g) were subjected to Rab8A-total (MJF-22-74-3) immunoblot analysis.

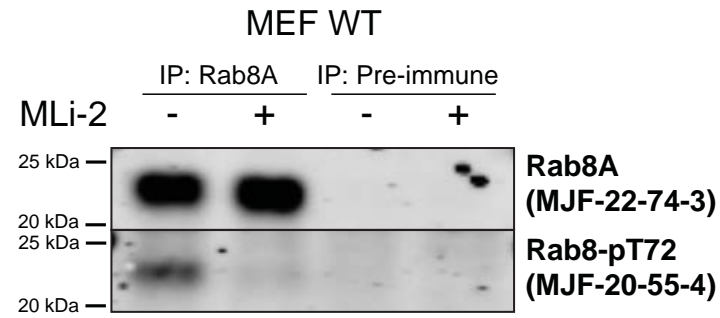
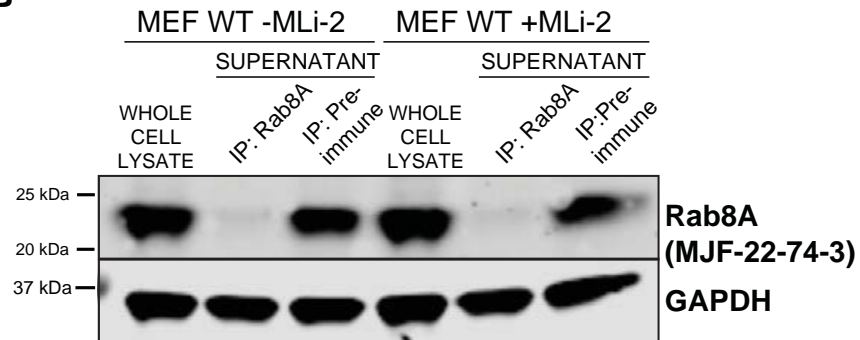
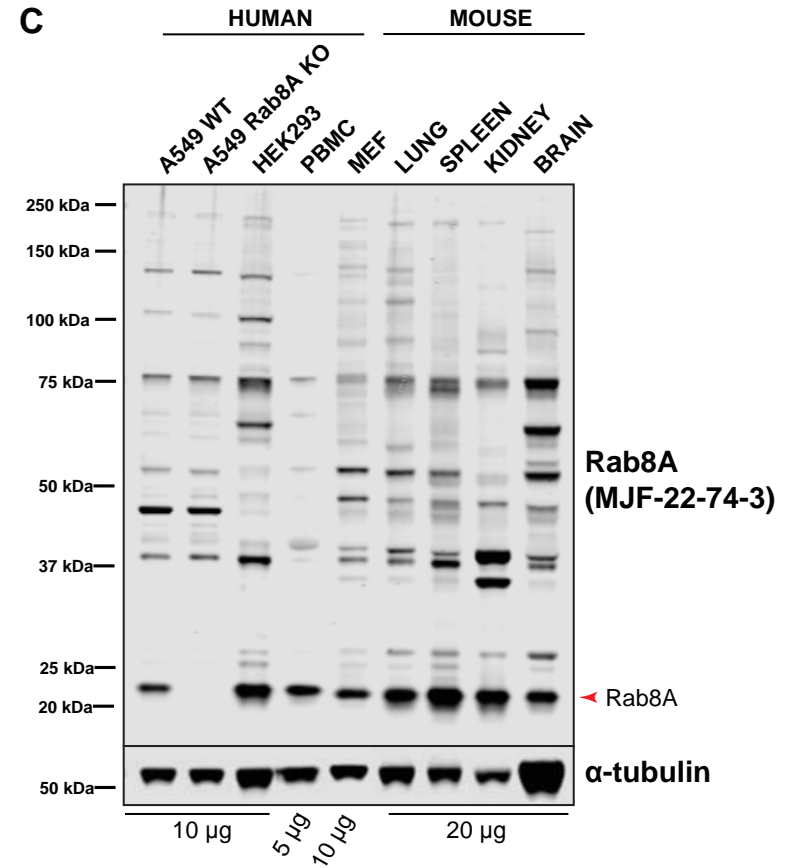
**Supplementary Figure 2. Characterisation of the rabbit monoclonal Rab12-pS106 (MJF-25-9-1) antibody.** (A) HEK293 cells were co-transfected with plasmids encoding the LRRK2[Y1699C] pathogenic mutation and the indicated Rab protein that all possessed an N-terminal HA tag. 36 hours post transfection, cells were treated  $\pm$  150 nM MLI-2 inhibitor for 90 min prior to cell lysis. 0.5  $\mu$ g of whole cell extract was subjected to quantitative immunoblot analysis with the indicated antibodies (all at 1  $\mu$ g/ml primary antibody concentration), and the membranes developed using the Odyssey CLx scan Western Blot imaging system. (B) The indicated wild type and Rab12 knock-out A549 cells were treated  $\pm$  100 nM MLI-2 for 90 min prior to cell lysis. Generation of the Rab12 knock-out cells is described in the materials and methods. 30  $\mu$ g of whole cell extract was subjected to quantitative immunoblot analysis with the indicated antibodies (all at 1  $\mu$ g/ml primary antibody concentration). The membranes developed using the Odyssey CLx scan Western Blot imaging system. A duplicate Rab12-pS106 immunoblot analysis was developed using enhanced chemiluminescence (ECL) for comparison.

**Supplementary Figure 3: Three structurally unrelated LRRK2 inhibitors reduce elevated Rab10 phosphorylation in dose dependent manner in homozygous VPS35[D260N] knock-in cells.** Homozygous VPS35[D620N] knock-in and wild-type MEFs were treated with the indicated doses of MLI-2 [34, 35], GSK2578215A [36] and PF-06447475 [37] LRRK2 inhibitor for 60 min prior to cell lysis. 7.5  $\mu$ g of whole cell extracts were subjected to quantitative immunoblot analysis with the indicated antibodies (all at 1  $\mu$ g/ml primary antibody concentration), and the membranes developed using the Odyssey CLx scan Western Blot imaging system. Immunoblots were quantified by LiCor and presented as average  $\pm$  SEM $\pi$ . Each lane represents cell extract obtained from a different dish of cells.

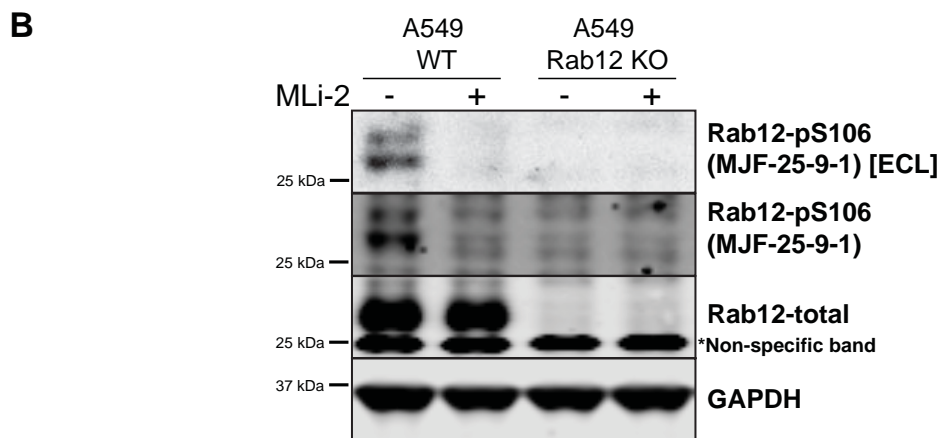
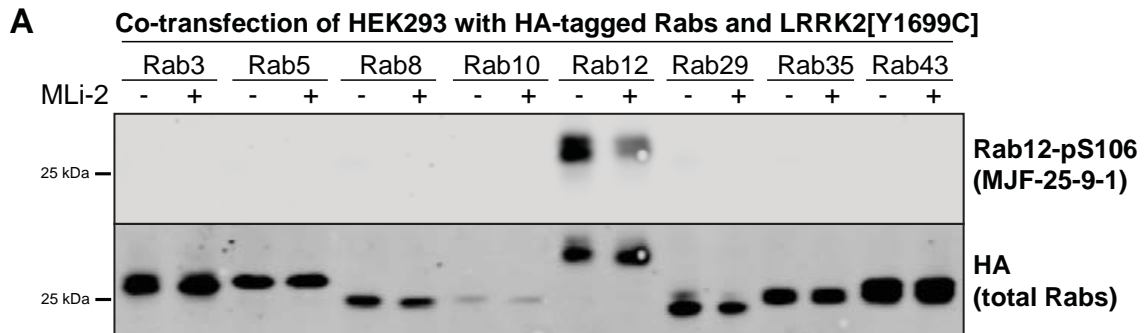
**Supplementary Figure 4: Flow cytometry analysis to determine the purity and viability of isolated monocytes and neutrophils, and the composition of PBMC.**

Purified monocytes were stained with the monocyte specific marker CD14 employing anti-human CD14 V500 antibody while purified neutrophils were stained with the granulocyte specific marker CD66b using anti-human CD66b FITC antibody. Cells were also stained with the cell viability dye DAPI. Cells were analysed by flow cytometry revealing that isolated monocytes had a viability >99% and a purity 87-90%, while isolated neutrophils had a viability >99% and a purity 94-98%. PBMC were stained with the B cell marker CD19 APC, the T cell marker CD3 PE, the monocyte marker CD14 V500, the granulocyte marker CD66b FITC and

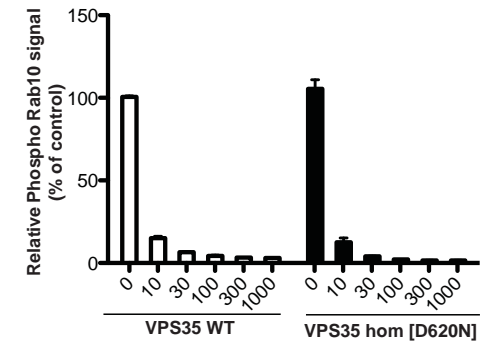
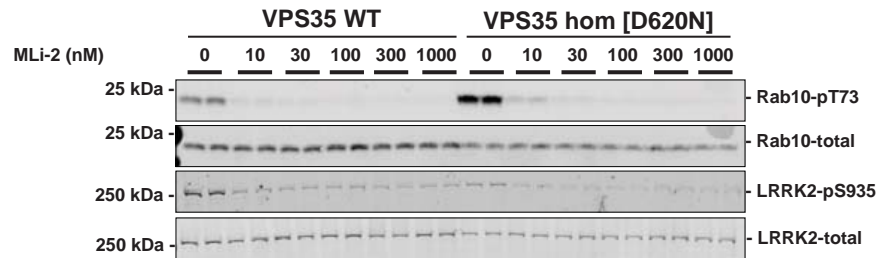
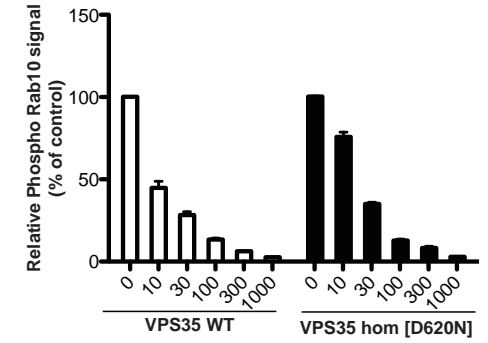
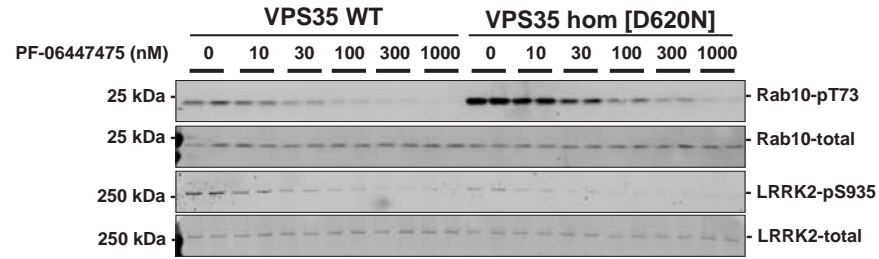
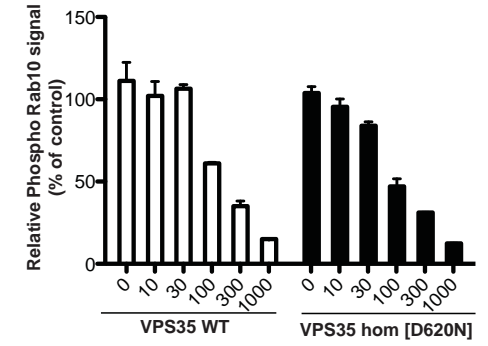
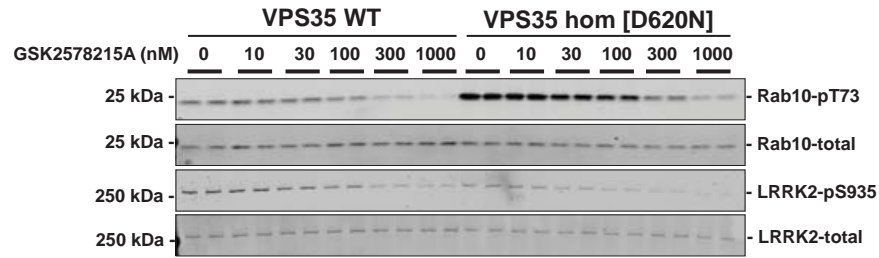
DAPI. \* corresponds to the monocyte and granulocyte parent gate. The proportion of each cell type within the isolated PBMC is provided. The data shown is for Donor A. A summary of the viability and purity data for all three donors is provided in Figure 6A.

**A****B****C**

**Supplementary Figure 1**

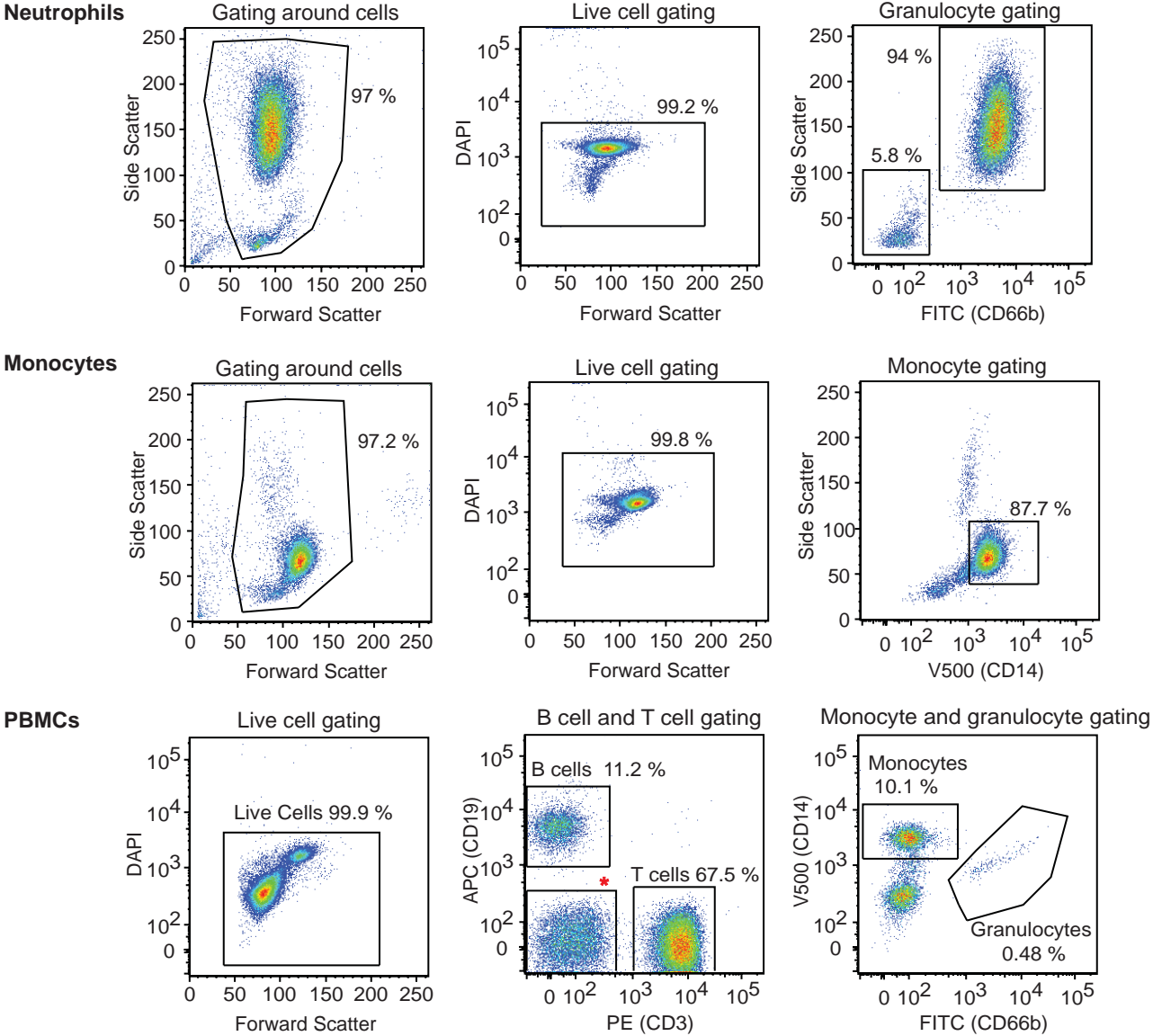


**Supplementary Figure 2**



Supplementary Figure 3

**S Figure 4**



**Supplementary Table S1: Demographic and clinical data for subjects analysed in Figures 4 and 6. Abbreviations iPD idiopathic Parkinson's disease; N/A not applicable; Age at Onset of Parkinson's disease AAO**

Donor number	Group	Gender	Age	AAO
1	iPD	m	57	50
2	iPD	f	66	52
3	iPD	m	71	69
5	iPD	f	53	50
6	iPD	m	75	68
7	iPD	m	82	72
8	iPD	m	58	55
9	Control non-PD	f	42	N/A
12	iPD	m	80	75
13	iPD	m	65	60
14	Control non-PD	f	27	N/A
15	VPS35het D620N	f	64	51
16	Control non-PD	m	65	N/A
17	VPS35het D620N	f	75	47
18	Control non-PD	m	58	N/A
19	Control non-PD	m	44	N/A
22	Control non-PD	m	52	N/A
24	Control non-PD	m	60	N/A
25	VPS35het D620N	f	54	47
26	Control non-PD	f	56	N/A
27	Control non-PD	f	70	N/A