

Expanded View Figures

Q17

Figure EV1. (A) interaction fingerprint clustering of LRRK2's modulators (MLi-2 and RAB29) interactomes. Cells contain the minimum expected distance obtained from the distograms of any residue of each interactor (row) to every residue in LRRK2 (column). Rows are color annotated based on confidence score, Jaccard, RMSD from reference LRRK2 structure (PDB: 7LHW), and experiment type (either MLI-2 or RAB29 overexpression); (B) interaction fingerprint of LRRK2 and interacting RABs (from IntAct).

