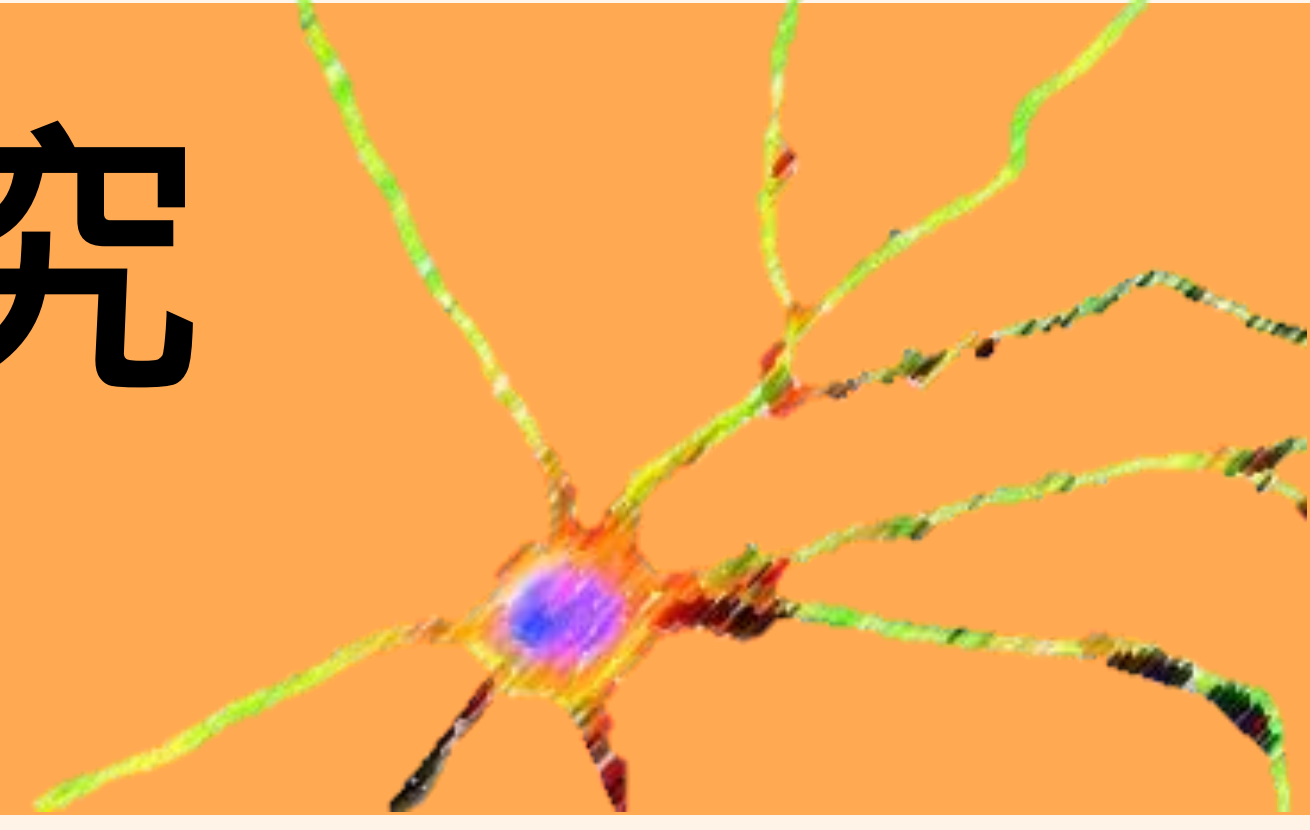


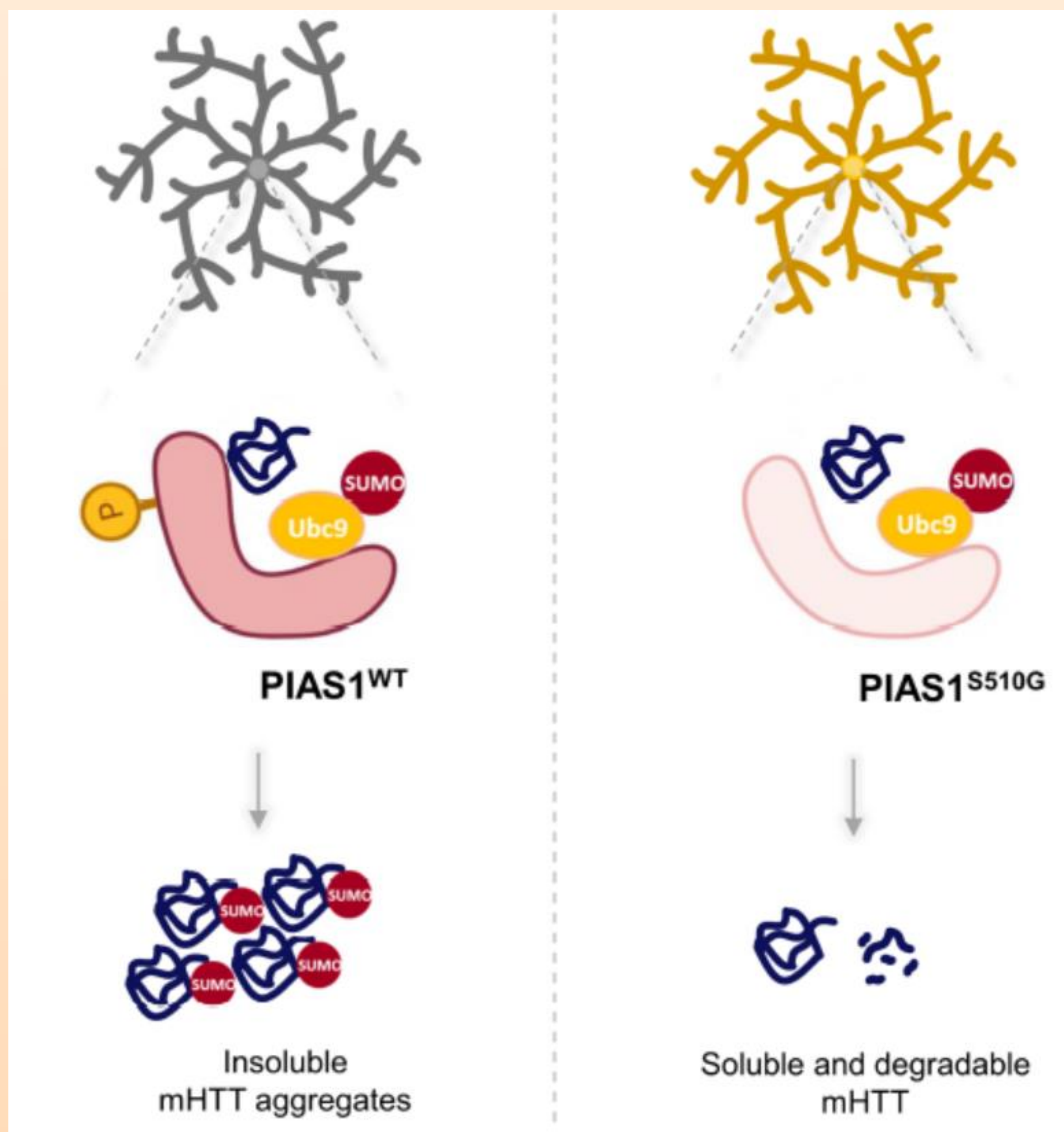
神經退化性疾病之轉譯醫學研究

(鄭子豪團隊)



遺傳性腦神經退化性疾病-亨式舞蹈症

- Huntington's Disease (HD) is an inherited neurodegenerative disorder, caused by a monogenic mutation in the coding sequence of the huntingtin (HTT) gene.
- **Genetic modifiers** are genetic variants that can modify the phenotypic outcome of the primary disease-causing gene.



We identified PIAS1 as a genetic modifier associated with late onset of polyQ diseases. The naturally-occurring PIAS1^{S510G} variant interact with mutant huntingtin (mHTT) poorly, resulting in a lower SUMOylation and accumulation of mHTT. Knock-in of PIAS1^{S510G} ameliorates major symptoms of HD in mice, supporting its protective role.

A PIAS1 protective variant S510G delays polyQ disease onset by modifying protein homeostasis. (Movement Disorders, IF=10.3, in press)

遺傳性腦神經退化性疾病-小腦萎縮症第三型

- Spinocerebellar Ataxia Type 3 (SCA3) is caused by expression of mutant *ATXN3*, encoding a translated product with a long stretch of polyQ that is aggregation-prone and detrimental to neurons.
- Using cellular and biochemical assays, we confirmed that PIAS1 (SUMO E3) enables to enhance *ATXN3* protein stability via the mode of sumoylation.

A *PIAS1* gene variant 3, associated with late onset of SCA3, exhibits a decreased interaction with its E2 (UBC9) in the presence of mutant *ATXN3*, which lead to less amount of SUMO conjugation, protein aggregation and cell death. Our findings provide an insight for the causal role of *PIAS1* gene variant 3 in late disease onset of patients with SCA3.

