



Supplementary Materials

Structure-Based Virtual Screening and De Novo Design to Identify Submicromolar Inhibitors of G2019S mutant of Leucine-Rich Repeat Kinase 2

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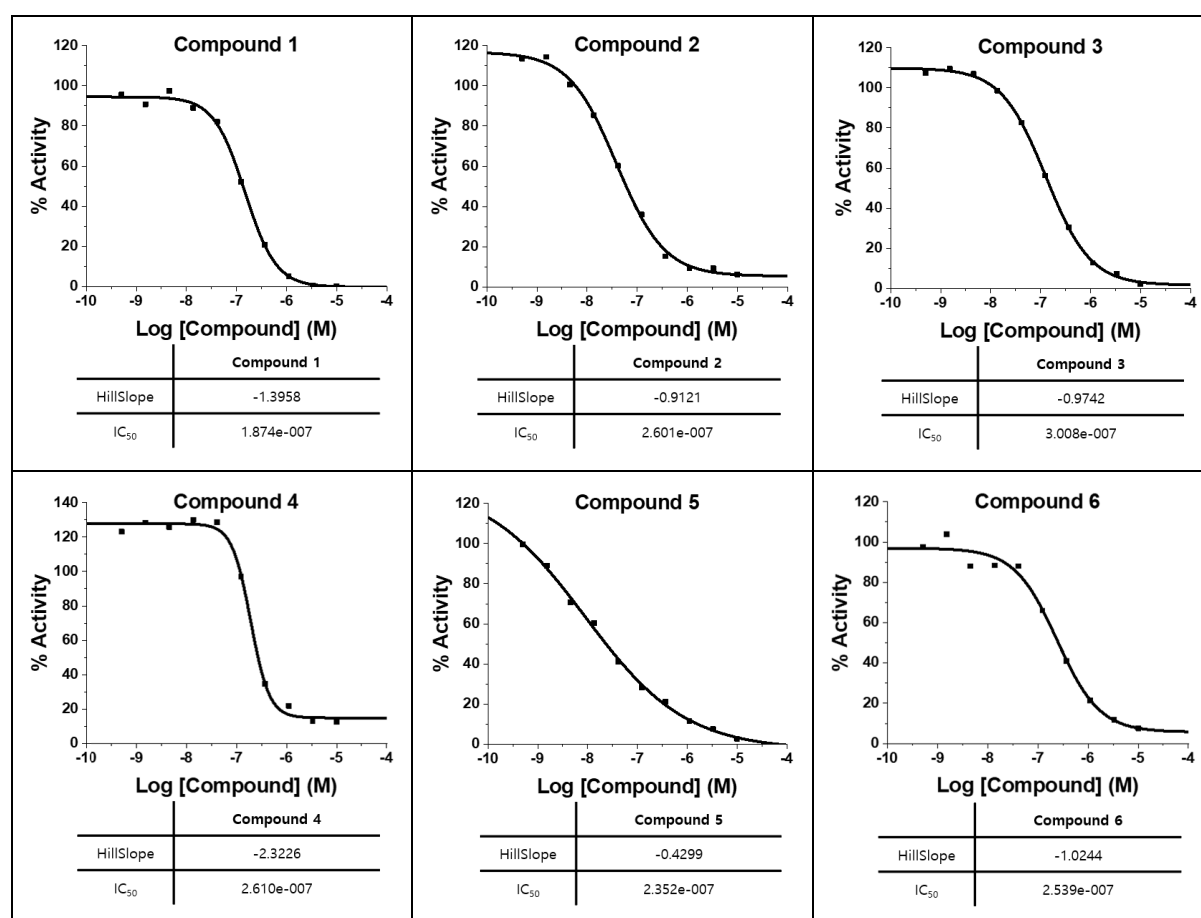
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Dose-response curve fits of **1-18** and **2a-2p**

Figure S1

Figure S1. Dose-response curve fits of **1-18** and **2a-2p**, which were used to obtain their IC₅₀ values with respect to (a) wild type and (b) G2019S mutant of LRRK2.

(a)



(b)

