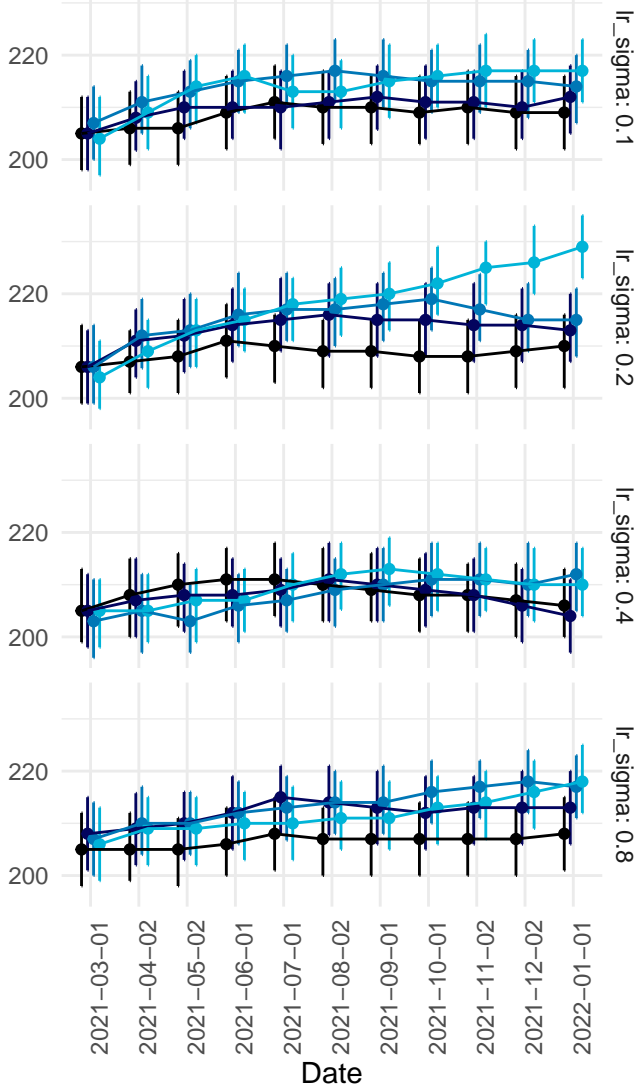


Number of selected features



P(mutation categorical genes)

- 0.25
- 0.1
- 0.04
- 0.016