

Total Structural Variations

500

400

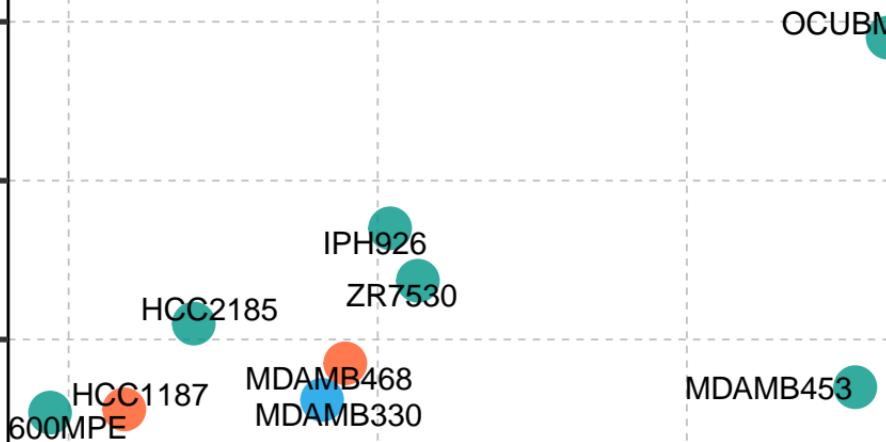
300

200

100

200

Total Mutations



$R = 0.43, p = 0.0831$