

Total Structural Variations

500

400

300

200

100

200

400

600

Total Mutations

SKBR3

OCUBM

IPH926

ZR7530

HCC2185

HCC1187

600MPE

MDAMB468

MDAMB330

MDAMB453

SUM44PE

UACC3133

MDAMB134VI

HCC2218

WCRC25

CAMA1

BCK4

$R = 0.43, p = 0.0831$