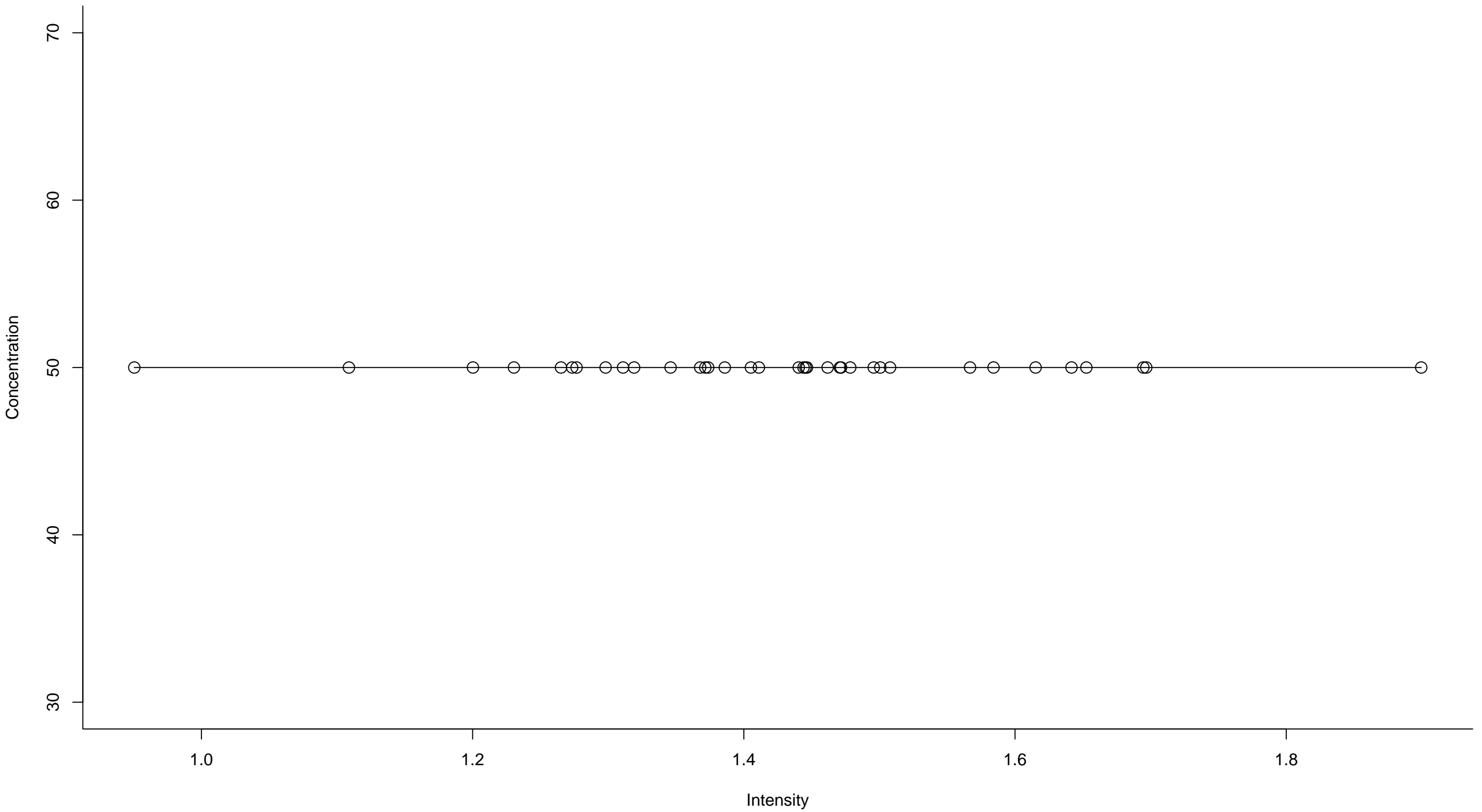


compound: b-MCA 13C6 batch: 1 raw_points: 36 conc: 50



compound: b-MCA 13C6 batch: 1 is_correction: none used_points: 36 fit_points: 36 conc: 50
R-square: 0.985 $y=34.5x$

