Sample: DO – each point represents one individual

total number of crossovers on autosomes – qtl2

total number of crossovers on autosomes – HaploQA
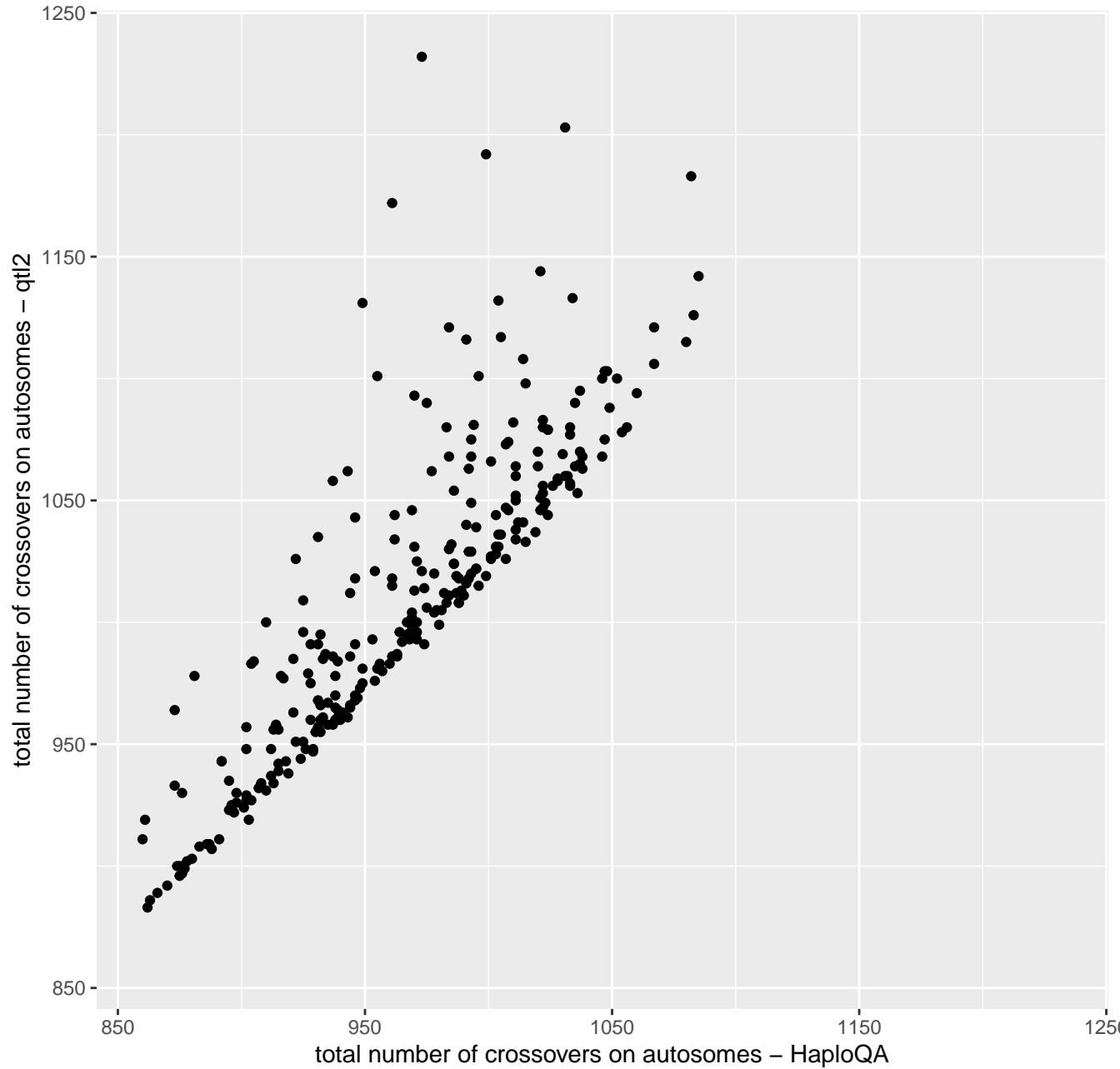
Sample: CC – each point represents one individual
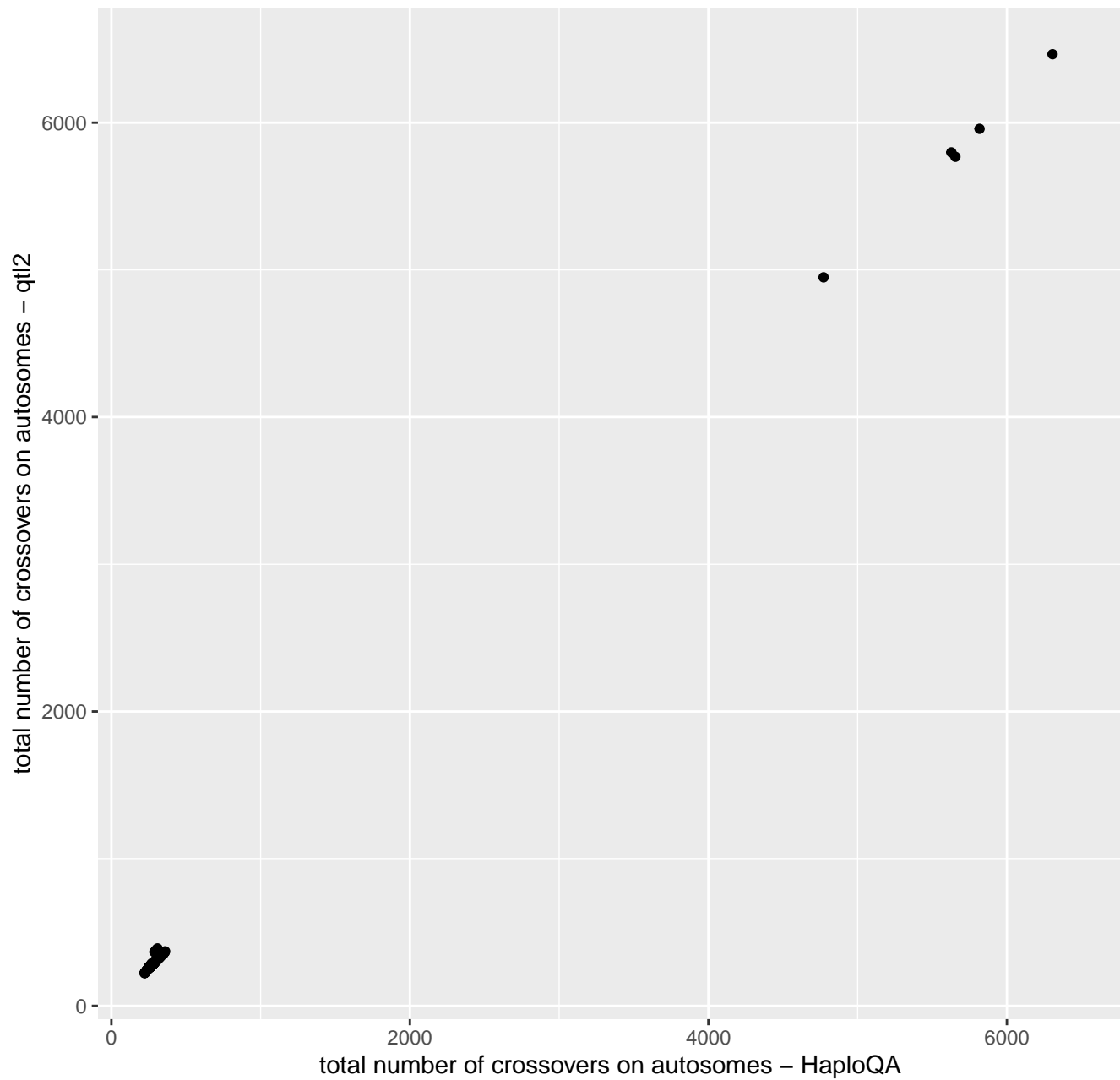
Sample: BXD – each point represents one individual

total number of crossovers on autosomes – qtl2 (y-axis)

total number of crossovers on autosomes – HaploQA (x-axis)