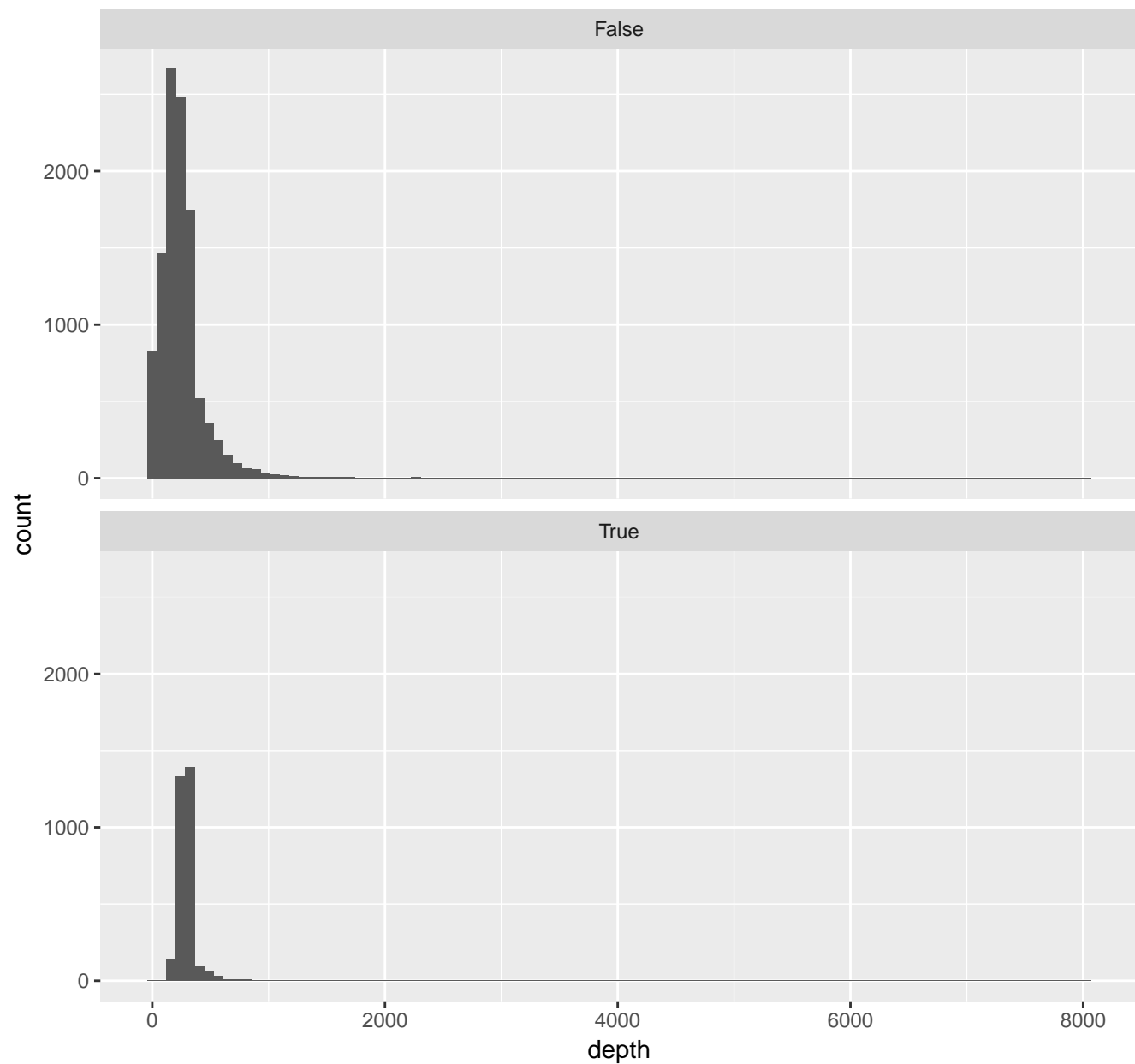


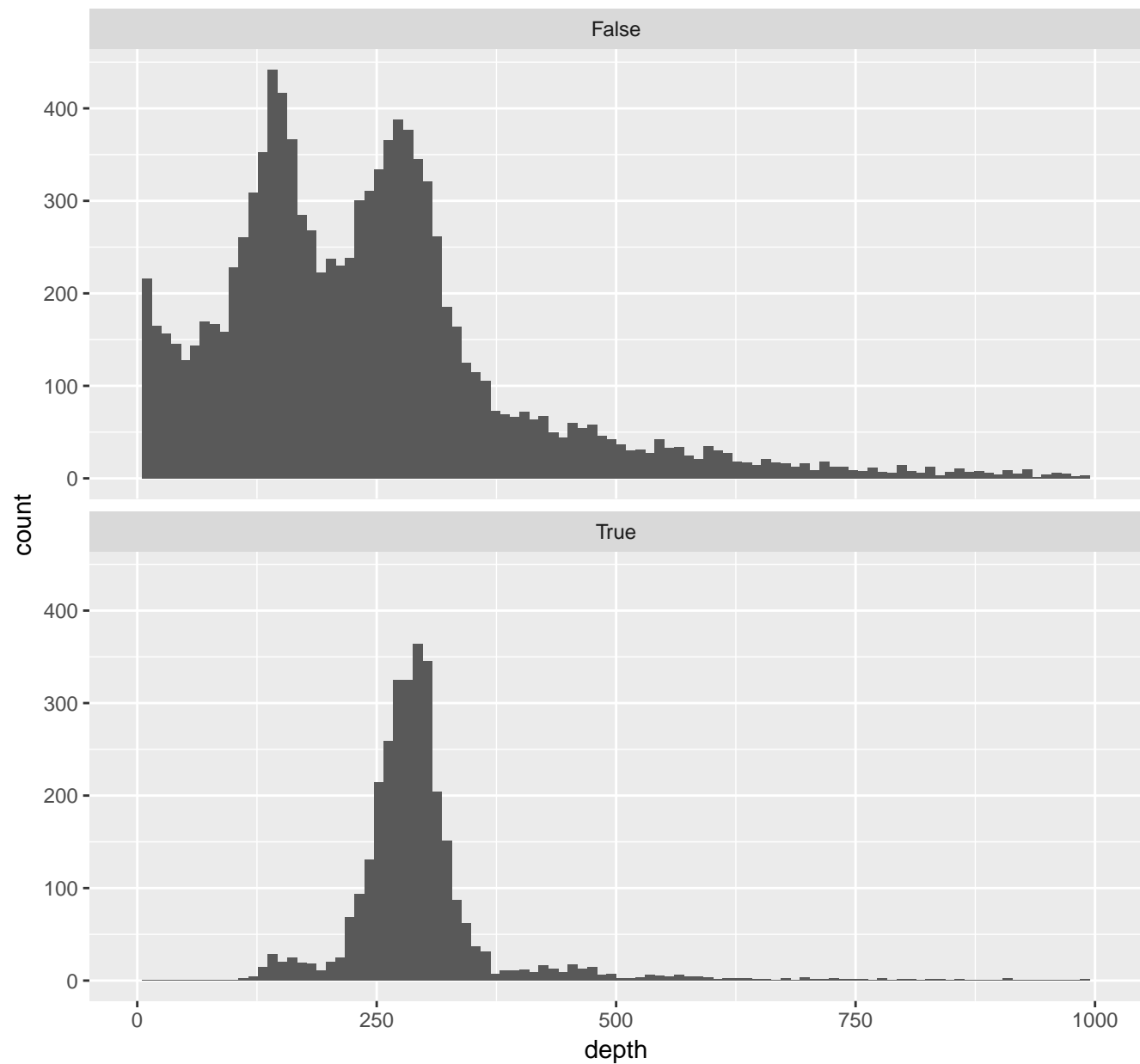
Detection rate

Coverage per site, with facets for whether or not an in-silico mutation was recovered



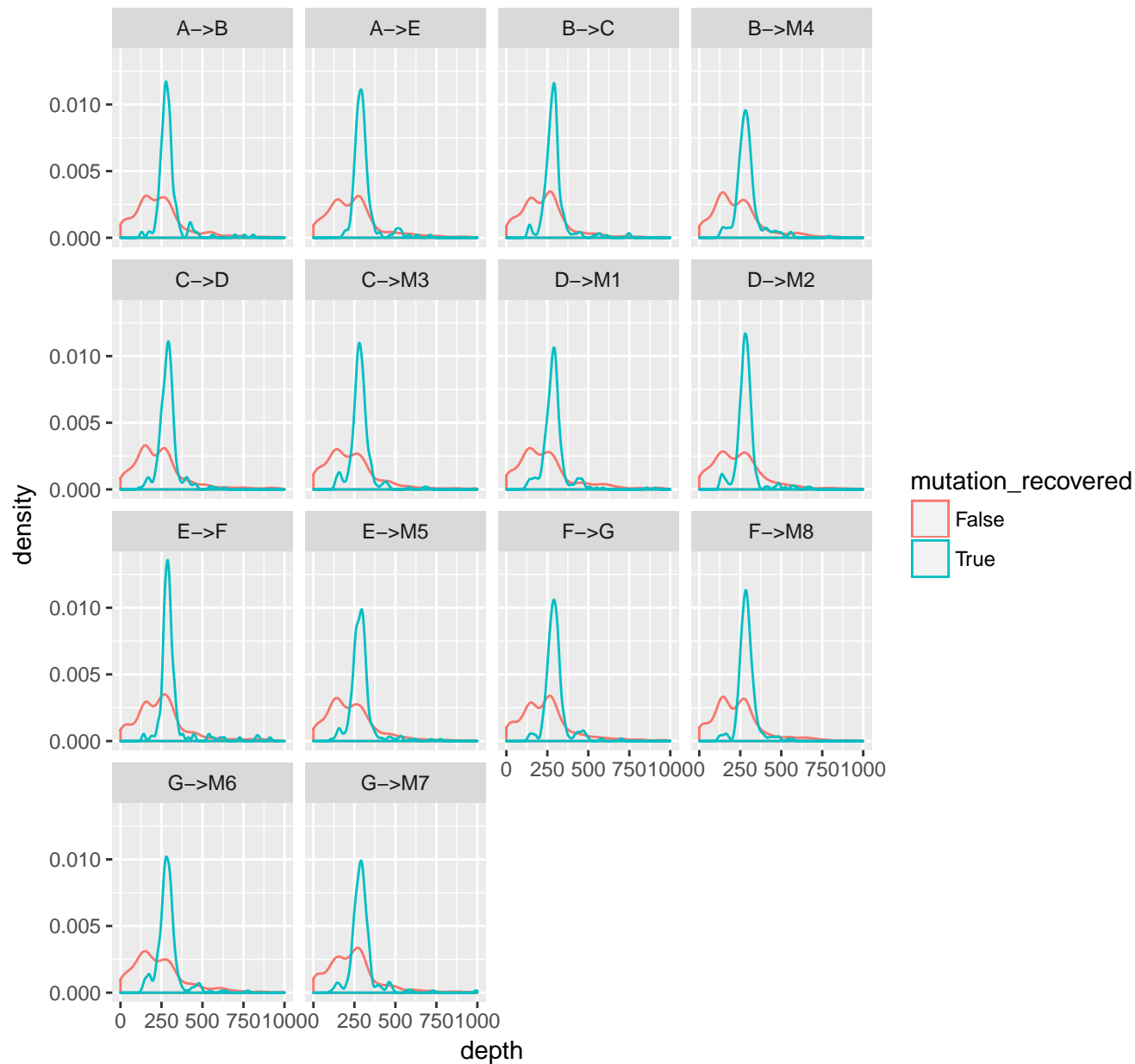
Detection rate, zoomed to x[0, 1000]

Coverage per site, with facets for whether or not an in-silico mutation was recovered



Detection rate per branch x[0, 1000]

Coverage per site, with facets for whether or not an in-silico mutation was recovered



Detection rate per scaffold x[0, 1000]

Coverage per site, with facets for whether or not an in-silico mutation was recovered

