Whole genome saddle strength profile at 500kb Control B2 4.5 Brd2_dep_rep1 4.0 Brd2 dep rep2 (YB + BB) / (BB + BY) / (BB + BY) / (BB + BY) 1.5 1.0 0.0 0.2 0.4 0.6 8.0 1.0 extent