Whole genome saddle strength profile at 200kb Control B2 4.0 Brd2_dep_rep1 Brd2_dep_rep2 3.5 (AA + BB) / (BB + AA) / (BB + AB) / (BB + AB) 1.5 1.0 0.0 0.2 0.4 0.6 1.0 0.8 extent