Figure S3

- Mapped
- Unmapped Whole Scaffold
- Unmapped Scaffold End
- Crossover/Misassembly Gap

Region Type vs. Region length (basepairs)
**Figure S3** Ranges of mapped and unmapped region lengths across all Hmel1.1 scaffolds. Crossover/Misassembly Gaps occur within scaffolds between markers, either consecutive on one chromosome (Crossover) or distant on one chromosome or on different chromosomes (Misassembly).