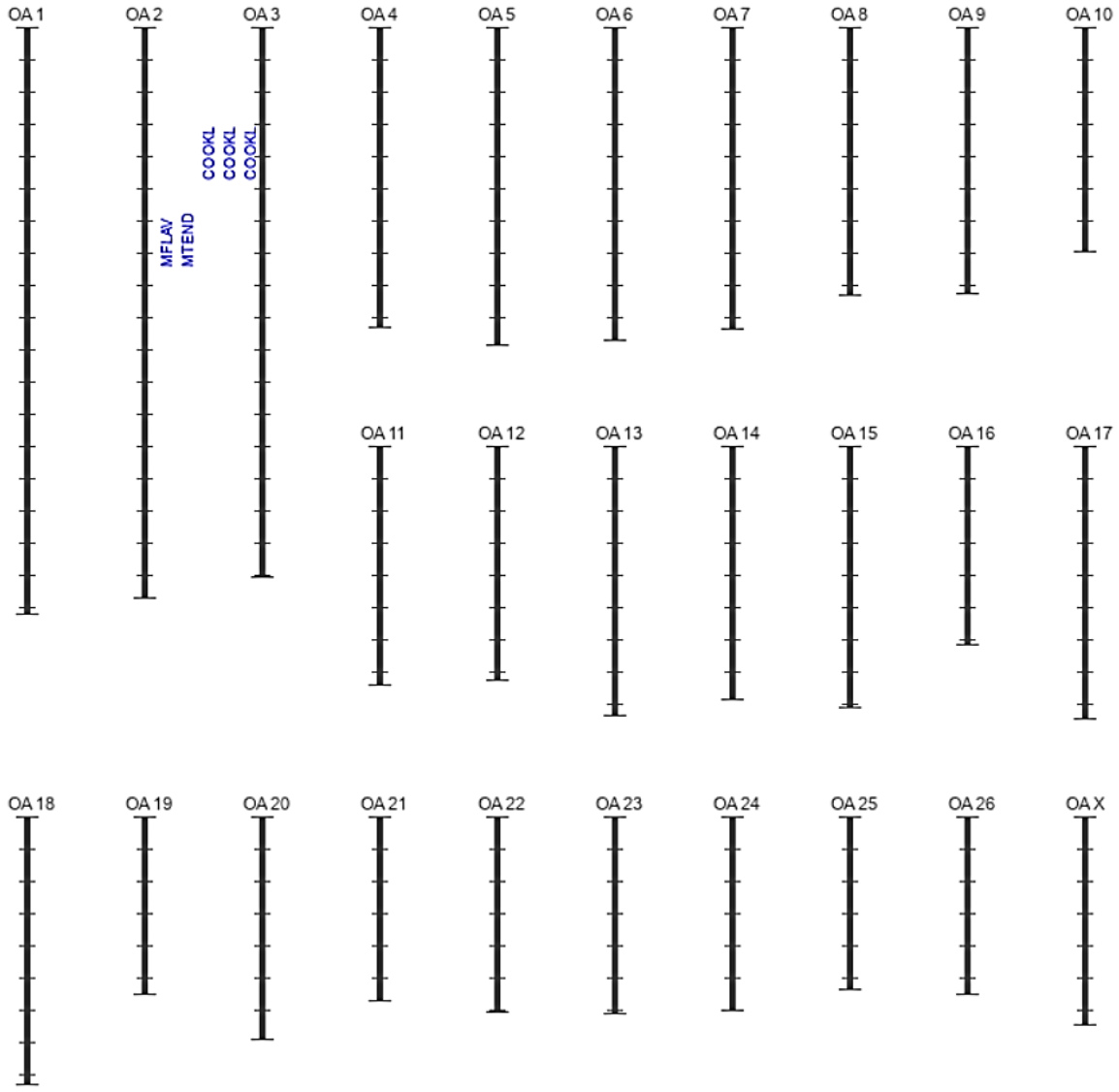


Figure S1. QTL/associations for meat sensory characteristics in the Sheep Genome (It shows genome locations where sensory characteristics traits are mapped by QTL or SNP associations).

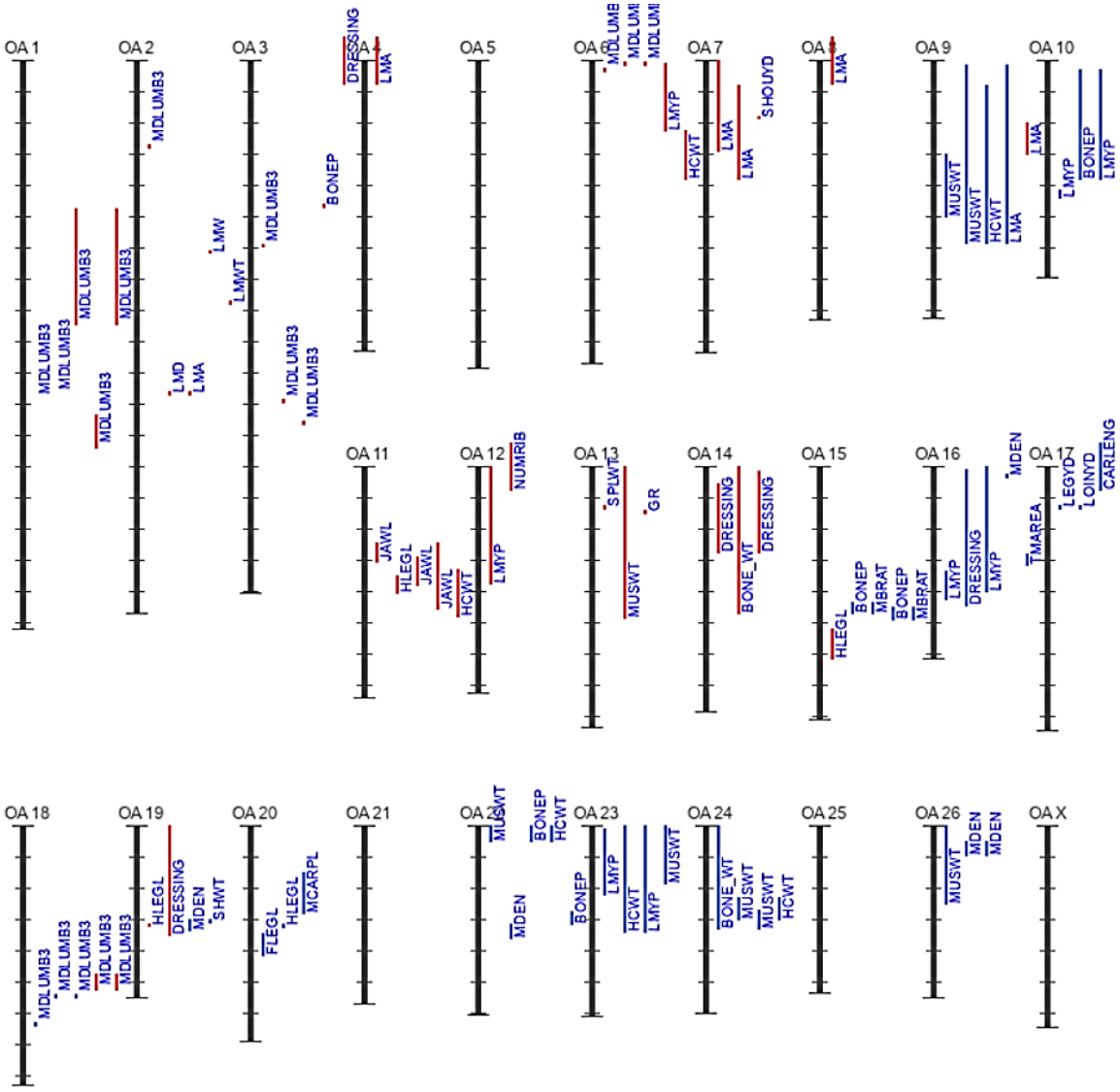


Figure S2. QTL/associations for meat carcass traits (Anatomy) in the Sheep Genome (It shows genome locations where anatomy traits are mapped by QTL or SNP associations).

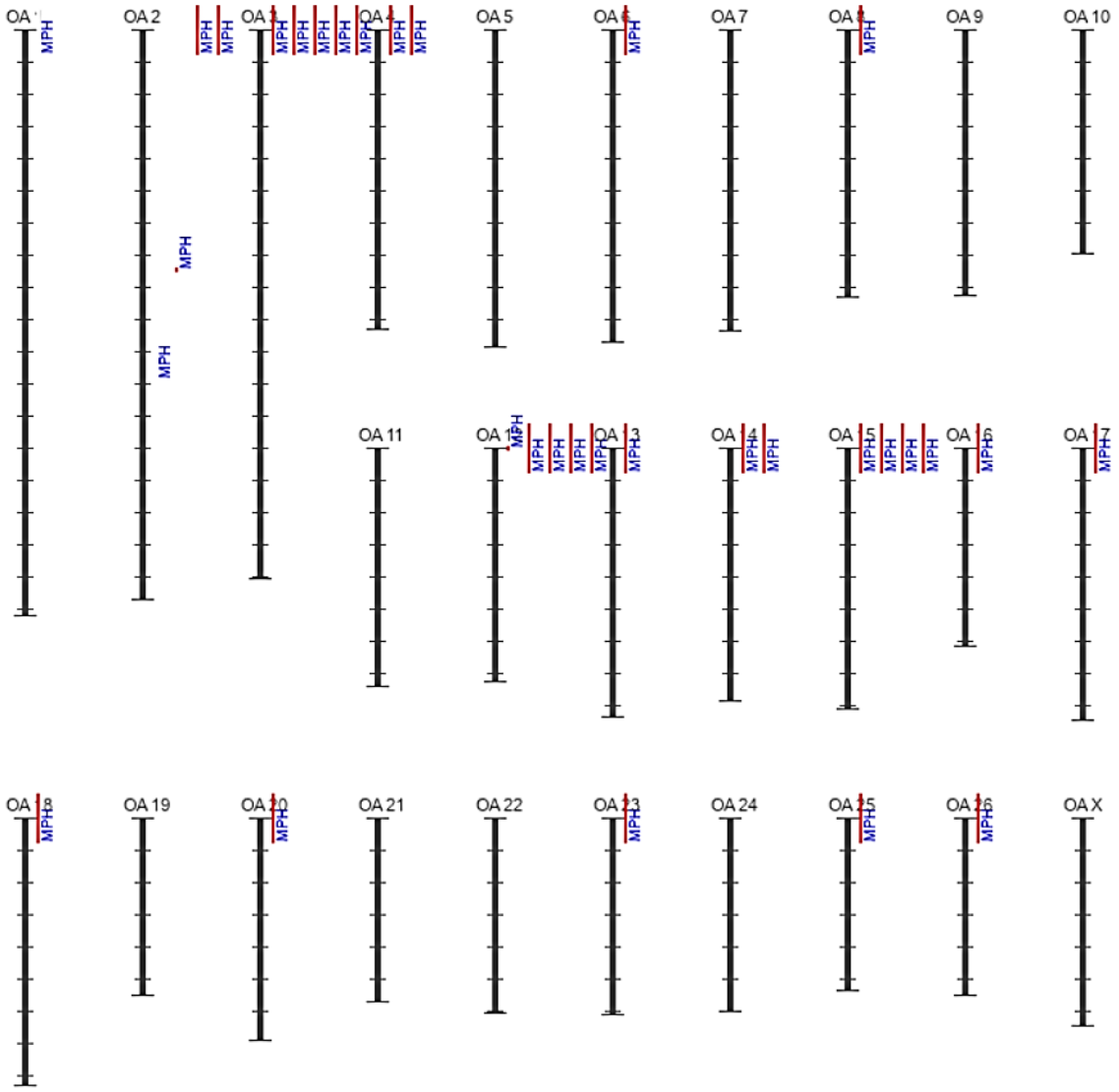


Figure S3. QTL/associations for meat carcass traits (Chemistry/ pH) in the Sheep Genome (It shows genome locations where chemistry traits are mapped by QTL or SNP associations).

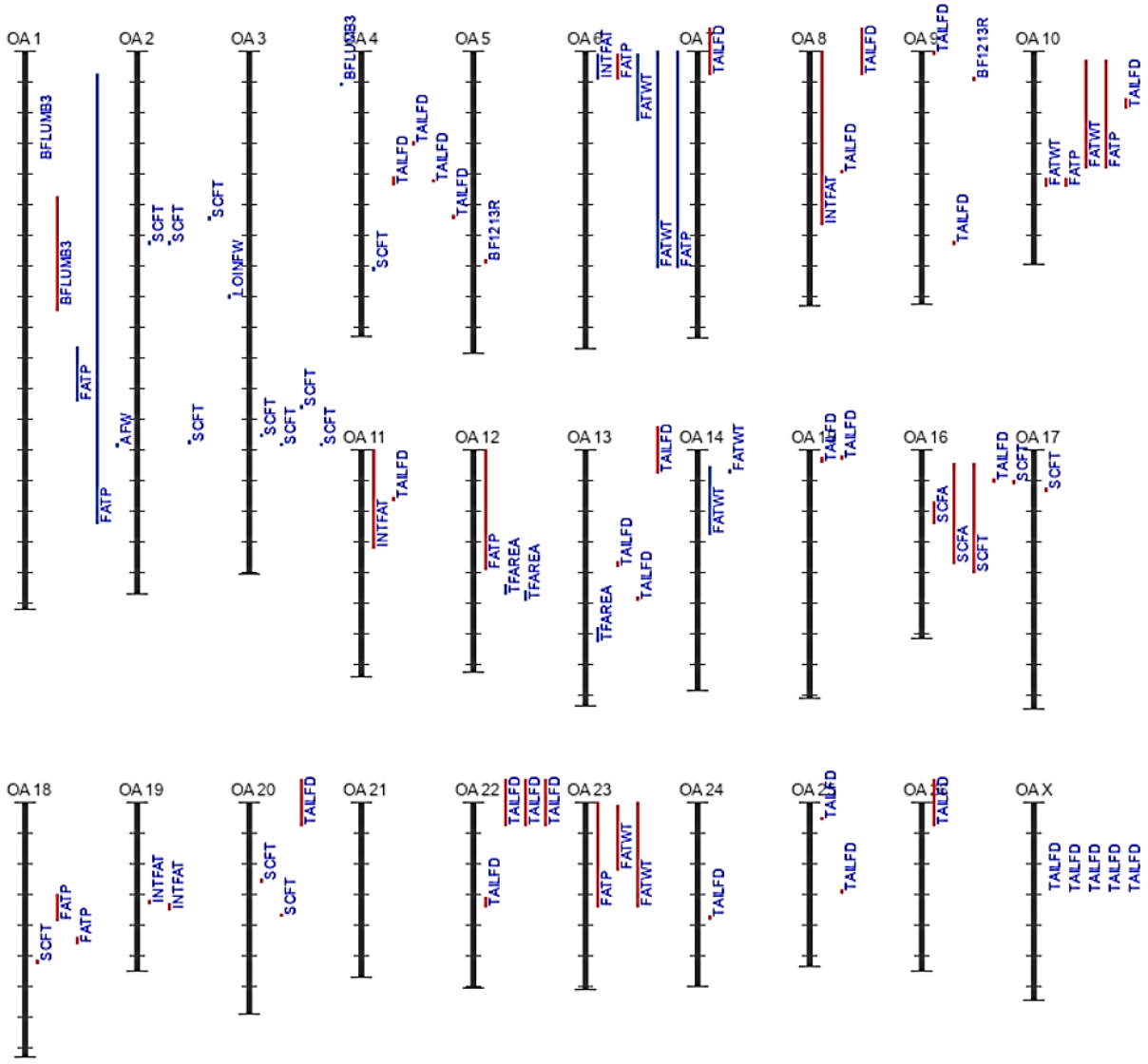


Figure S4. QTL/associations for meat carcass traits (Fatness) in the Sheep Genome (It shows genome locations where fatness traits are mapped by QTL or SNP associations).

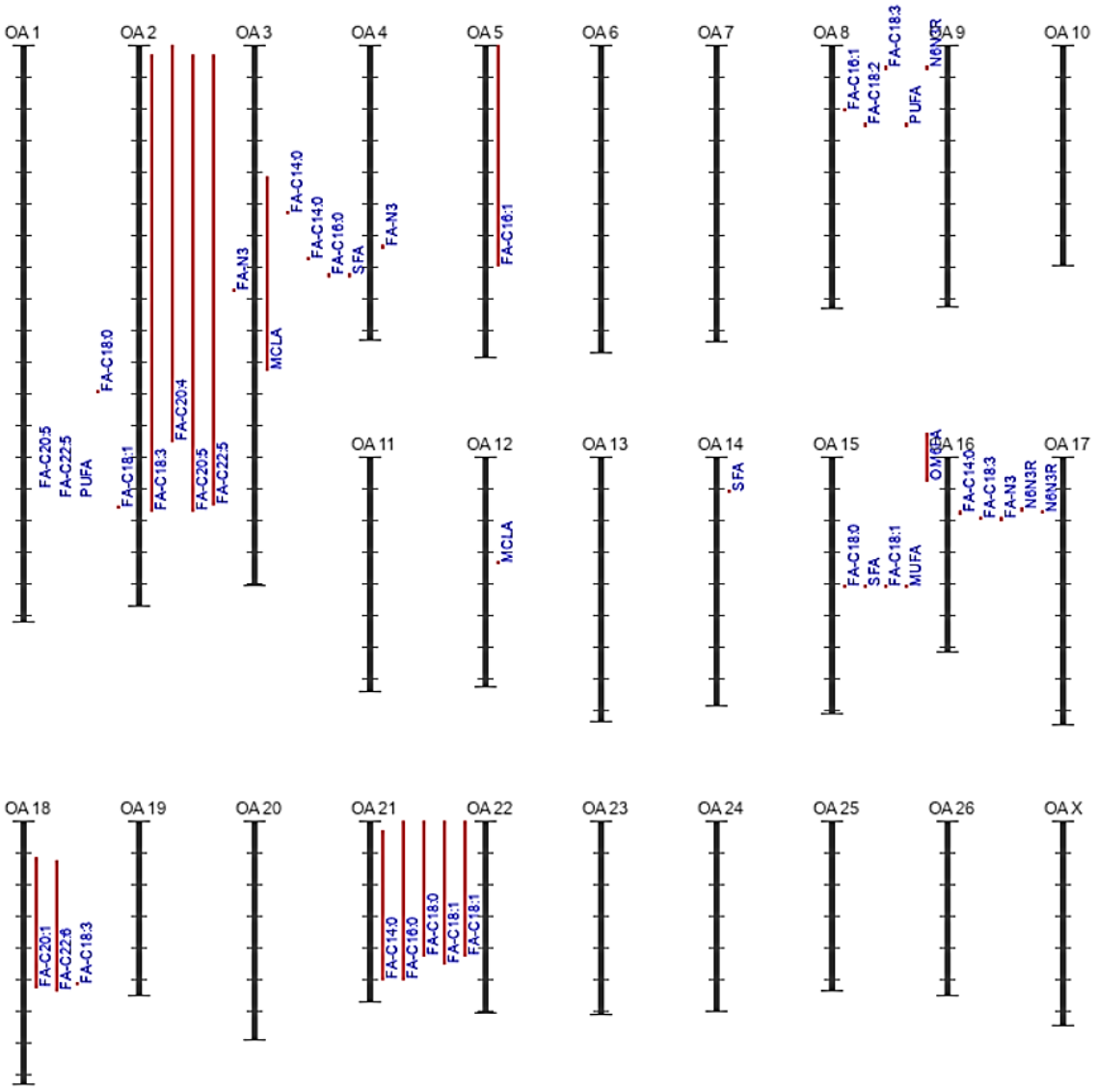


Figure S5. QTL/associations for meat carcass traits (Fat acid content) in the Sheep Genome (It shows genome locations where fat acid traits are mapped by QTL or SNP associations).

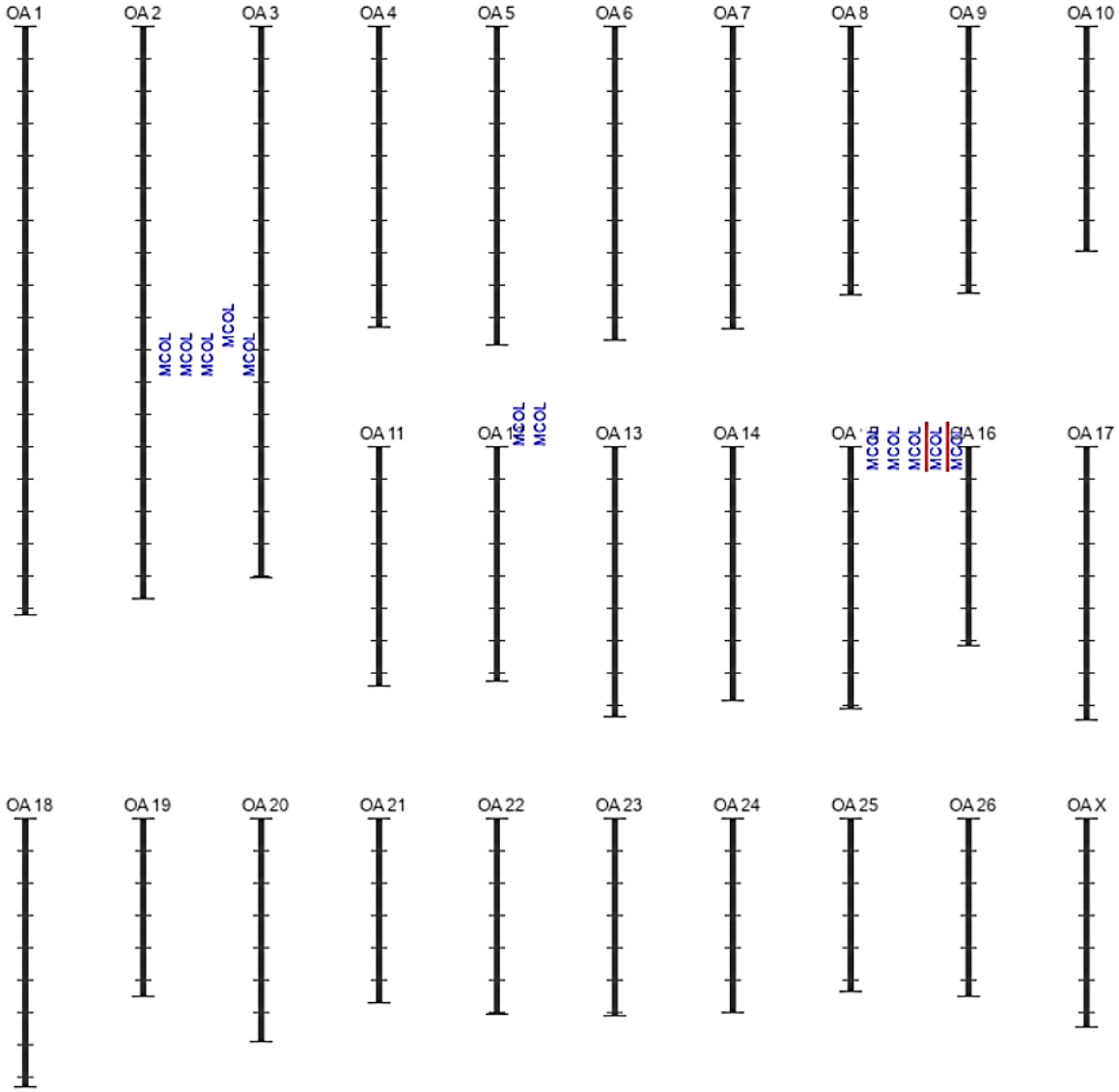


Figure S6. QTL/associations for meat carcass traits (Meat Colour) in the Sheep Genome (It shows genome locations where meat colour are mapped by QTL or SNP associations).

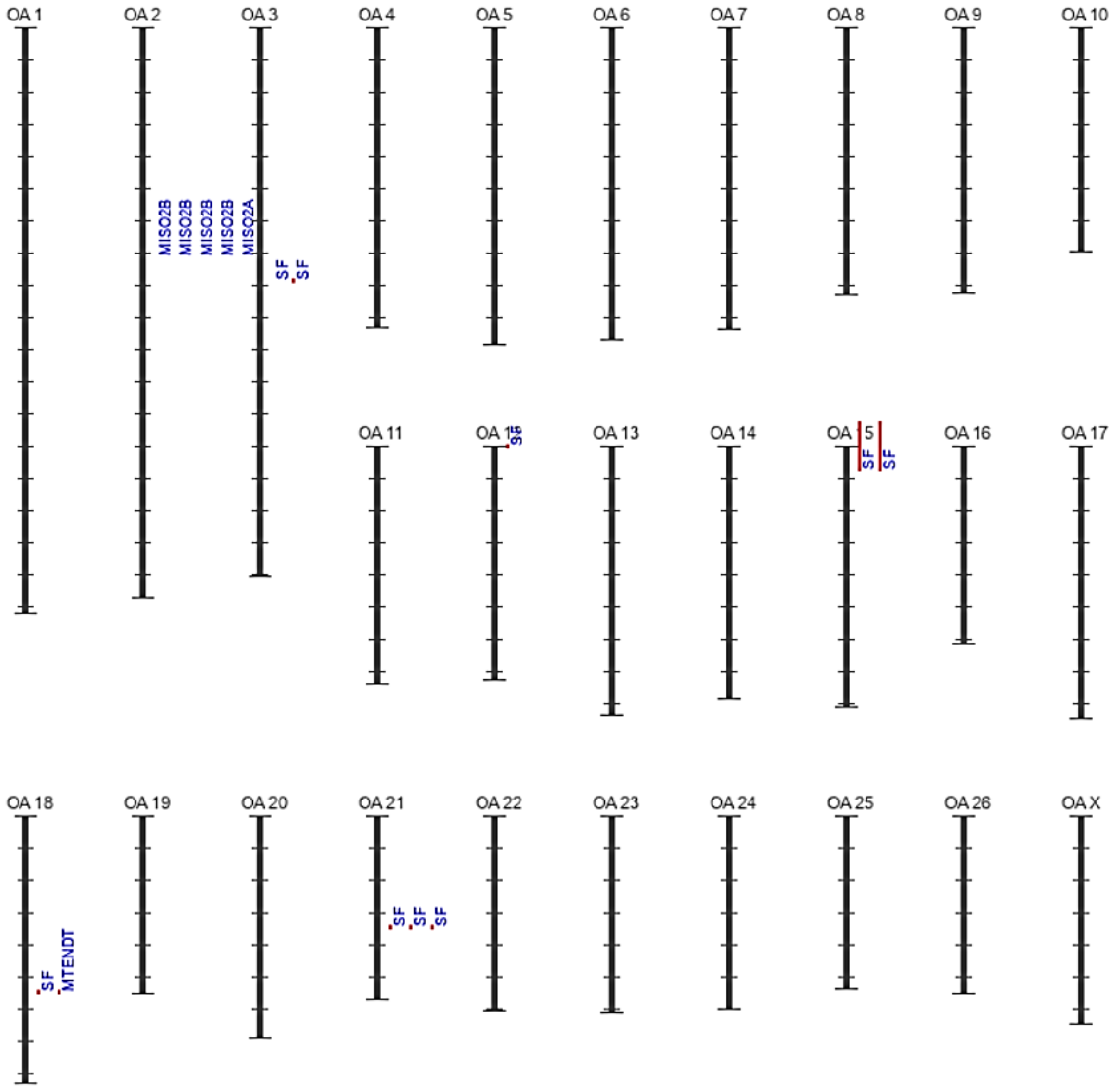


Figure S7. QTL/associations for meat carcass traits (Meat Texture) in the Sheep Genome (It shows genome locations where meat texture traits are mapped by QTL or SNP associations).

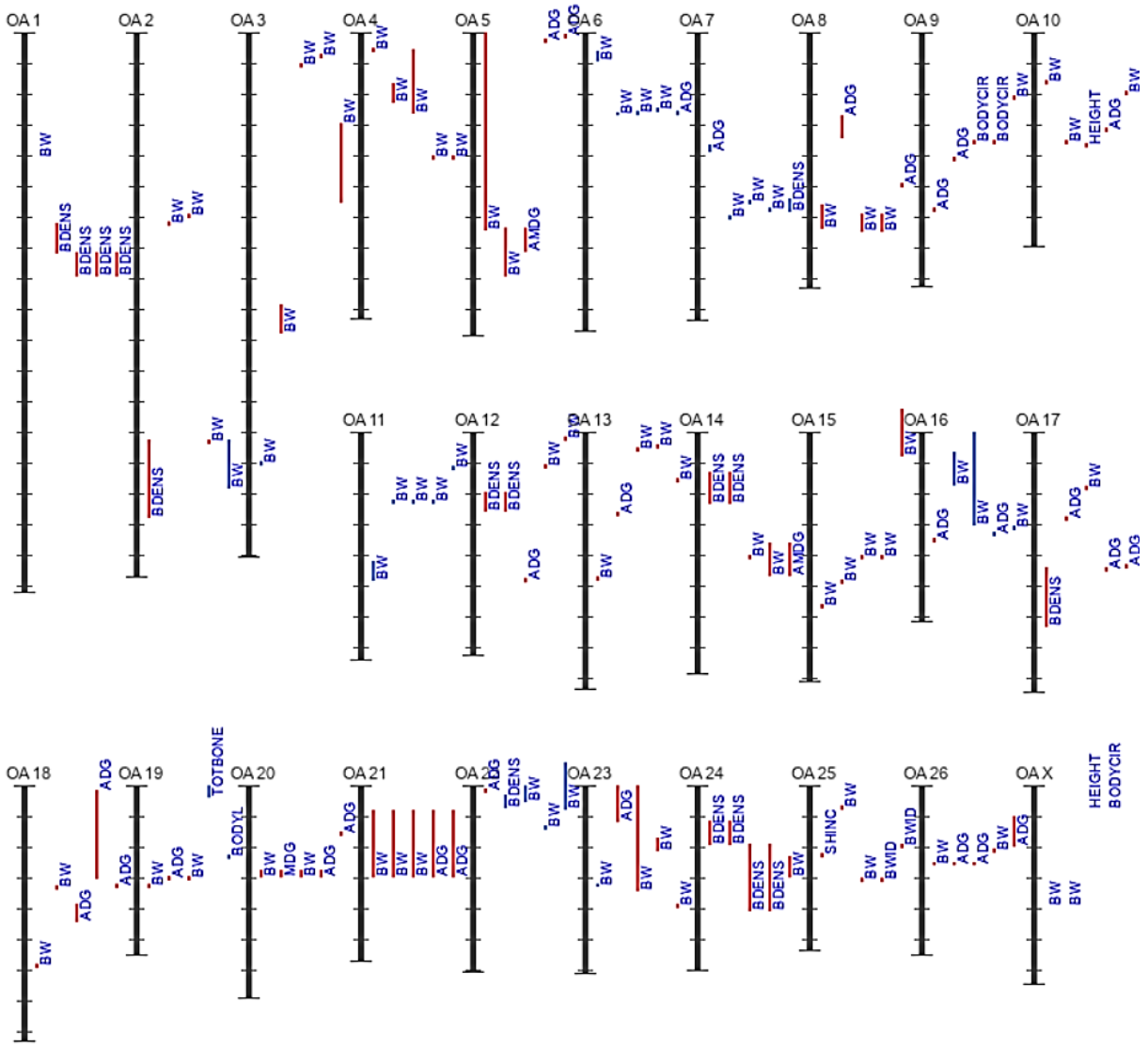


Figure S8. QTL/associations for Growth and carcass characteristics in the Sheep Genome (It shows genome locations where growth traits are mapped by QTL or SNP associations).