QTL mapping of traits across CO2 levels

Legend:
- biomass
- biomass_log
- leaf_mass
- leaf_mass_log
- seed_mass
- seed_mass_log
- stem_mass
- stem_mass_log
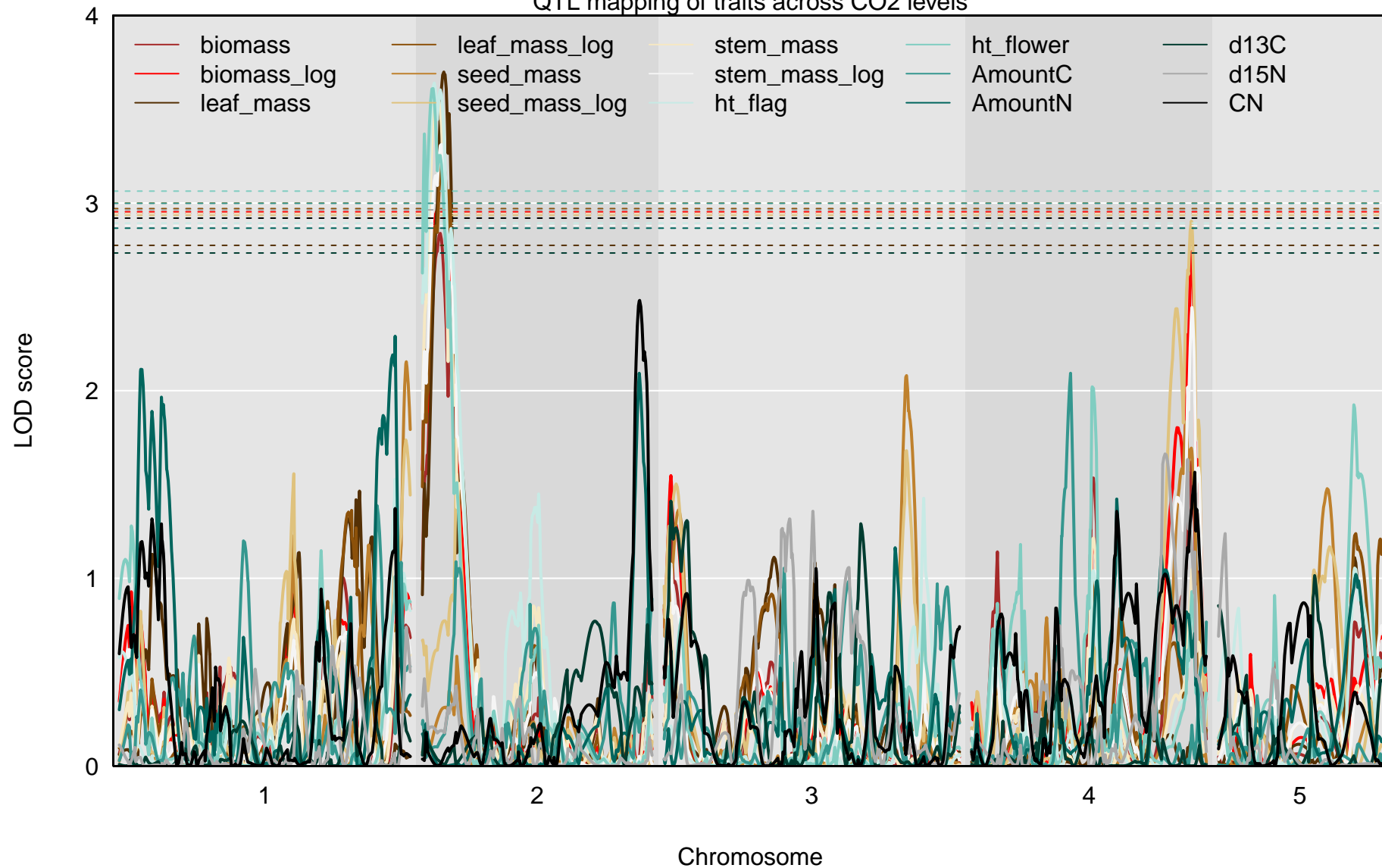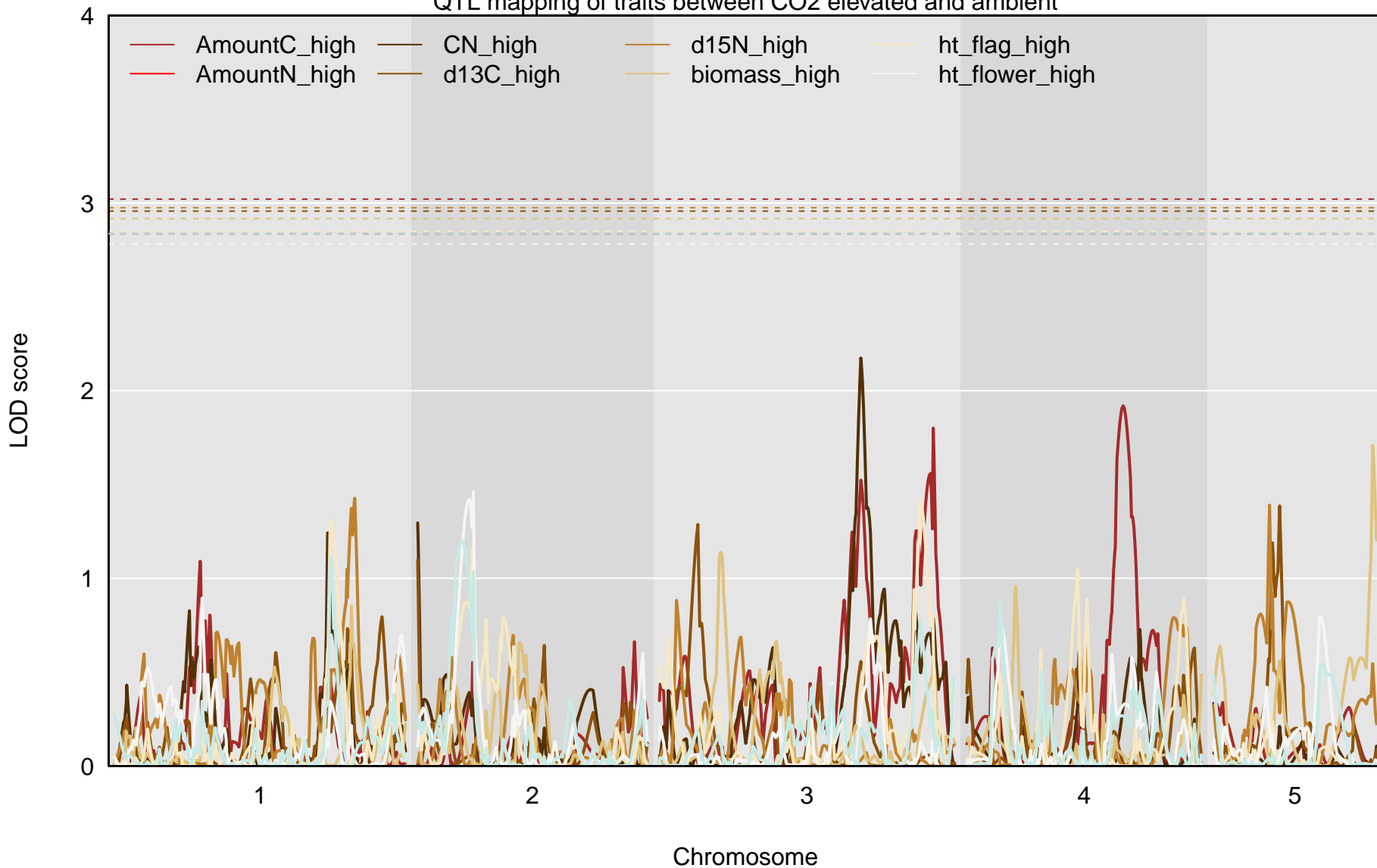- ht_flag
- ht_flower
- AmountC
- AmountN
- d13C
- d15N
- CN

Y-axis: LOD score
X-axis: Chromosome

QTL mapping of traits between CO2 elevated and ambient

QTL mapping of traits between CO2 subambient and ambient