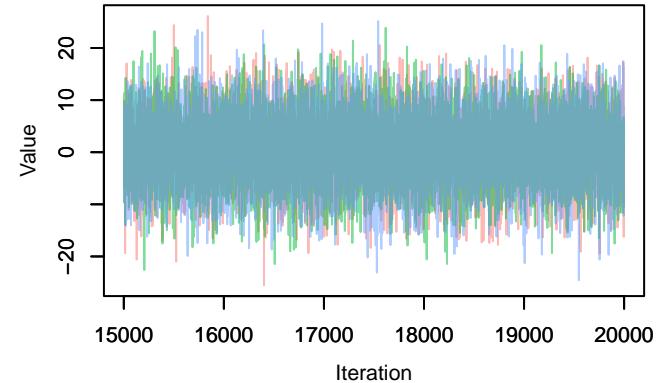
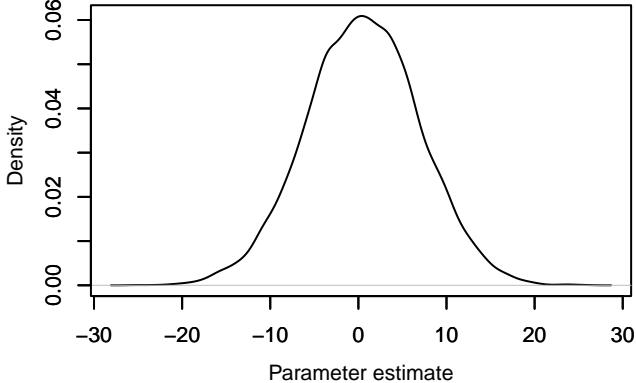


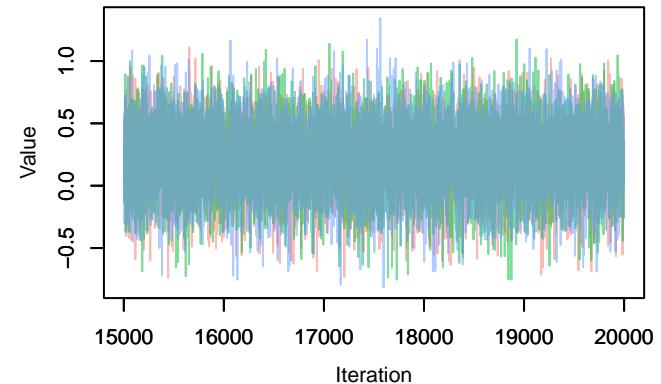
Trace – $B[(\text{Intercept}) (\text{C1}), \text{Angiostrongylus} (\text{S1})]$



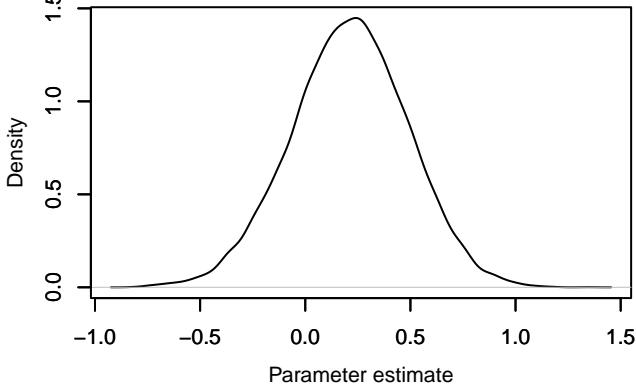
Density – $B[(\text{Intercept}) (\text{C1}), \text{Angiostrongylus} (\text{S1})]$



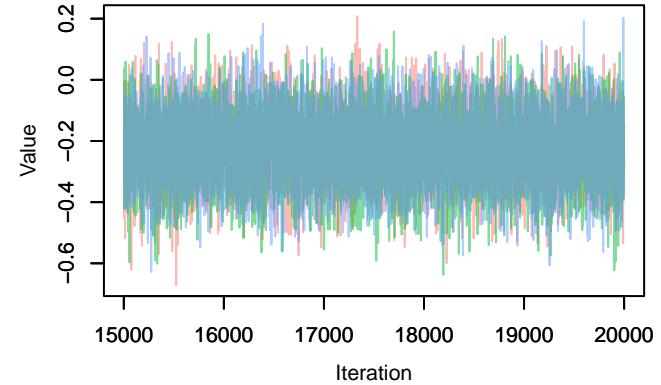
Trace – $B[\text{sexmale} (\text{C2}), \text{Angiostrongylus} (\text{S1})]$



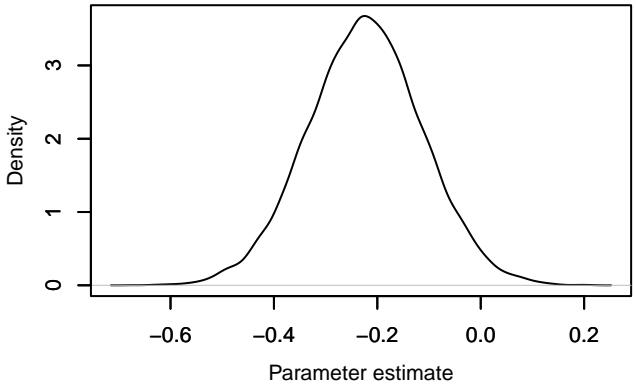
Density – $B[\text{sexmale} (\text{C2}), \text{Angiostrongylus} (\text{S1})]$



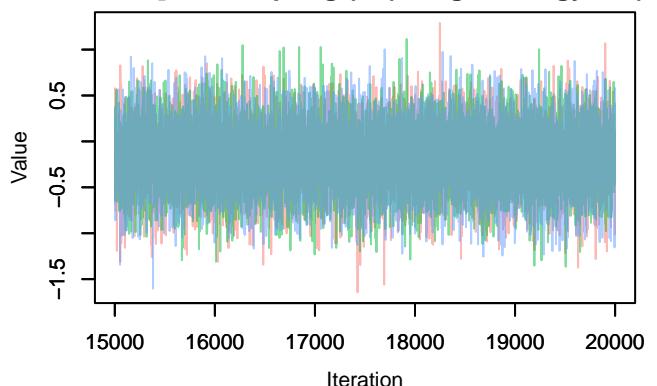
Trace – $B[\text{weight_kg} (\text{C3}), \text{Angiostrongylus} (\text{S1})]$



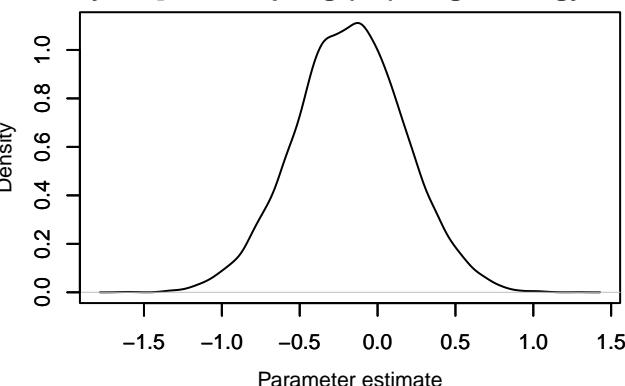
Density – $B[\text{weight_kg} (\text{C3}), \text{Angiostrongylus} (\text{S1})]$



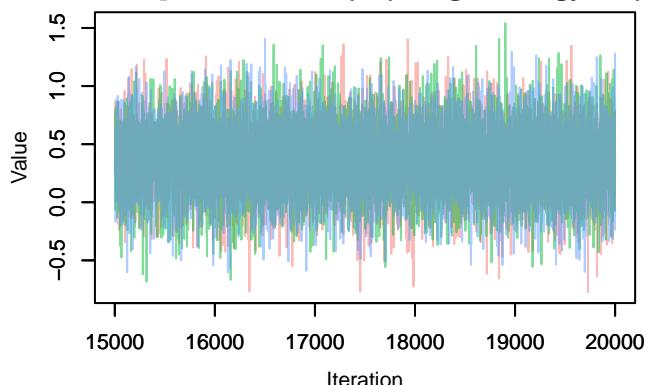
Trace – $B[\text{seasonspring (C4)}, \text{Angiostrongylus (S)}]$



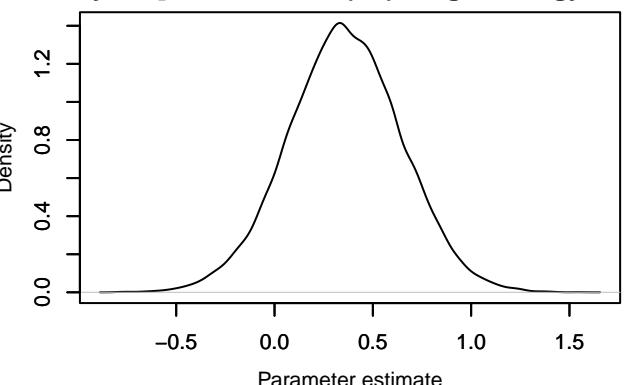
Density – $B[\text{seasonspring (C4)}, \text{Angiostrongylus (S)}]$



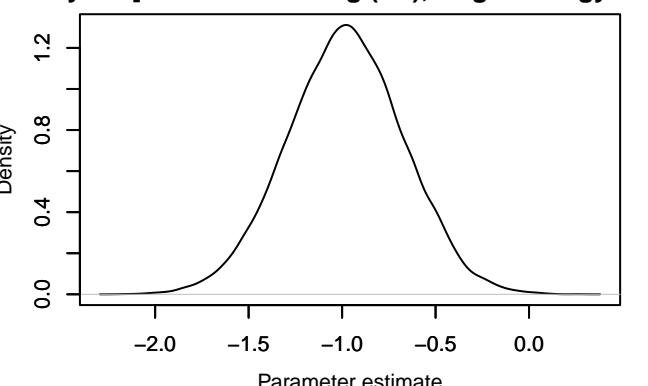
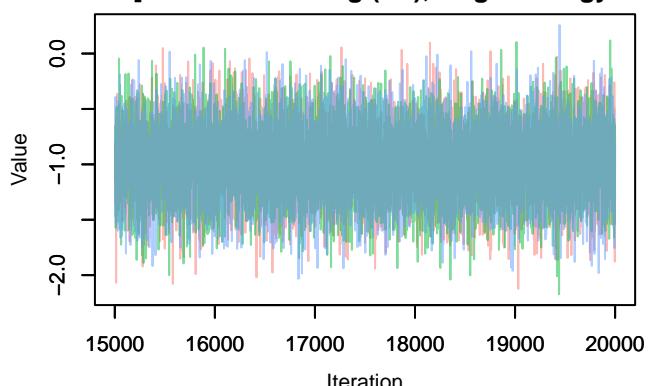
Trace – $B[\text{seasonwinter (C5)}, \text{Angiostrongylus (S)}]$



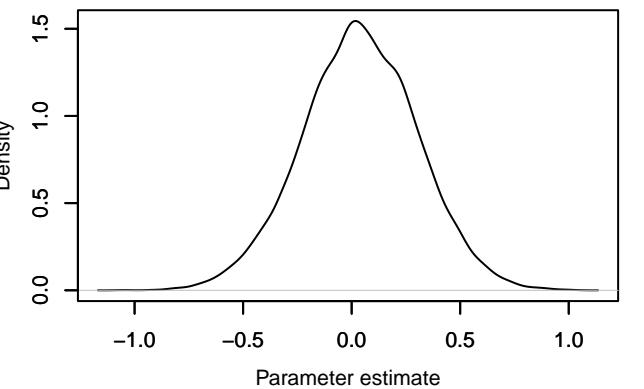
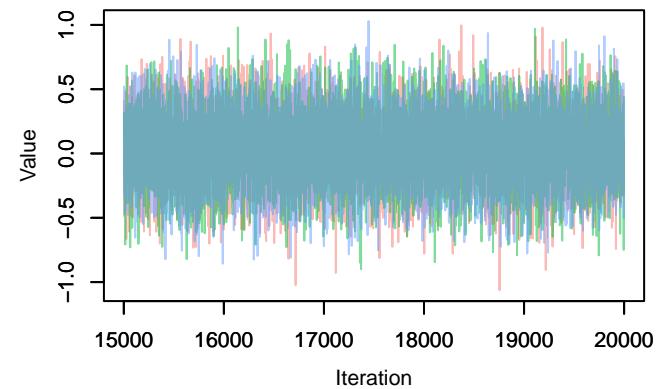
Density – $B[\text{seasonwinter (C5)}, \text{Angiostrongylus (S)}]$



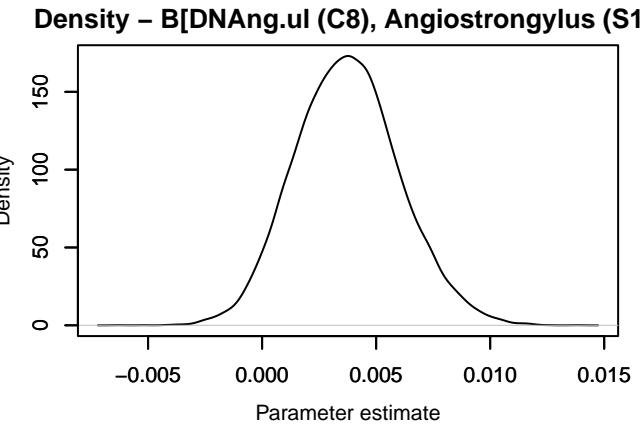
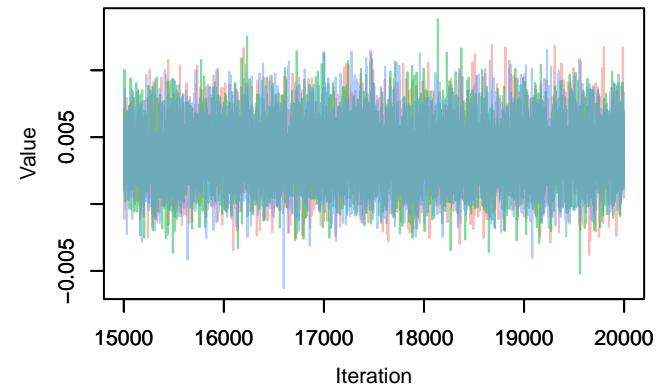
Trace – $B[\text{areaBrandenburg (C6)}, \text{Angiostrongylus (S)}]$



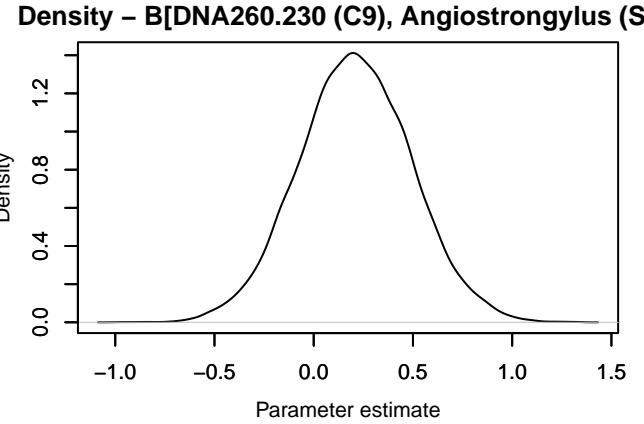
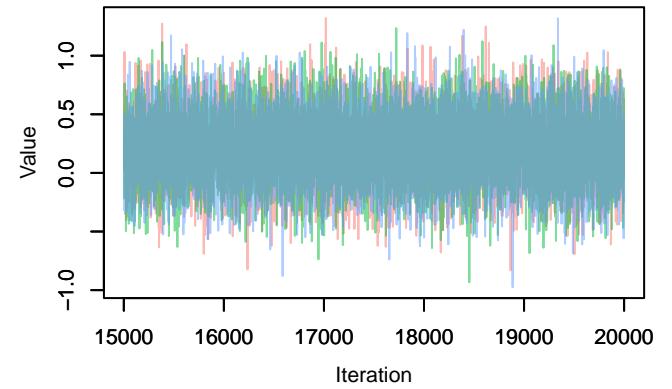
Trace – $B[\text{conditionexcellent (C7), Angiostrongylus}]$ Density – $B[\text{conditionexcellent (C7), Angiostrongylus}]$



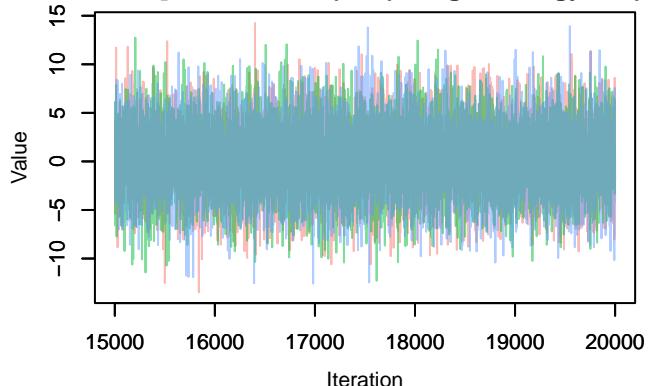
Trace – $B[\text{DNAng.ul (C8), Angiostrongylus (S1)}]$ Density – $B[\text{DNAng.ul (C8), Angiostrongylus (S1)}]$



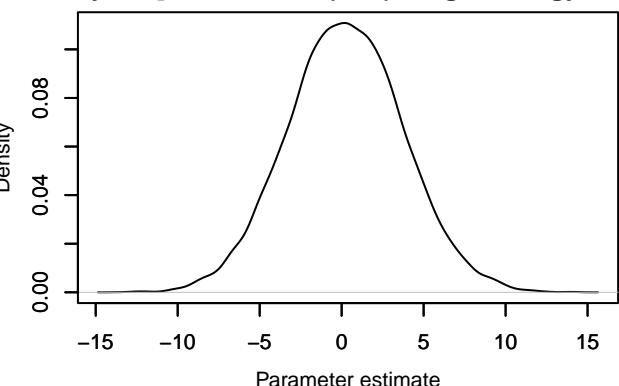
Trace – $B[\text{DNA260.230 (C9), Angiostrongylus (S1)}]$ Density – $B[\text{DNA260.230 (C9), Angiostrongylus (S1)}]$



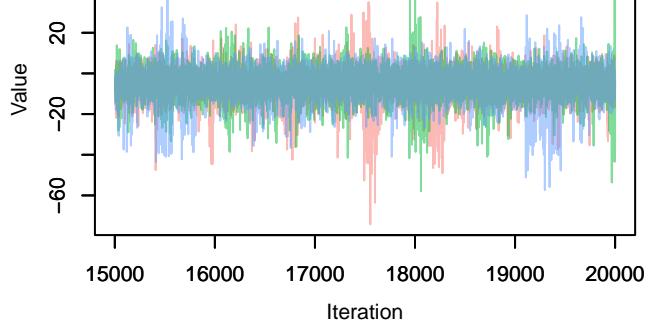
Trace – $B[\text{DNA260.280 (C10), Angiostrongylus (S1)}]$



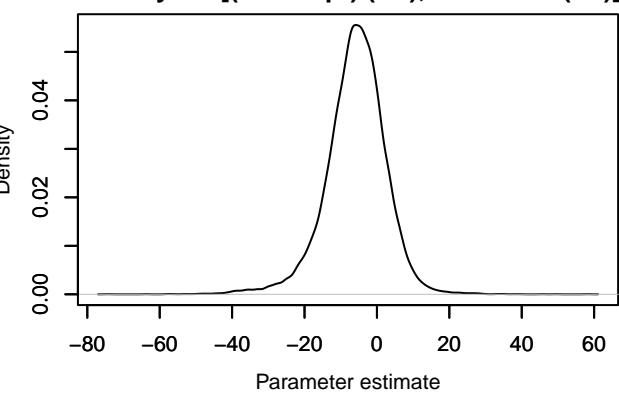
Density – $B[\text{DNA260.280 (C10), Angiostrongylus (S1)}]$



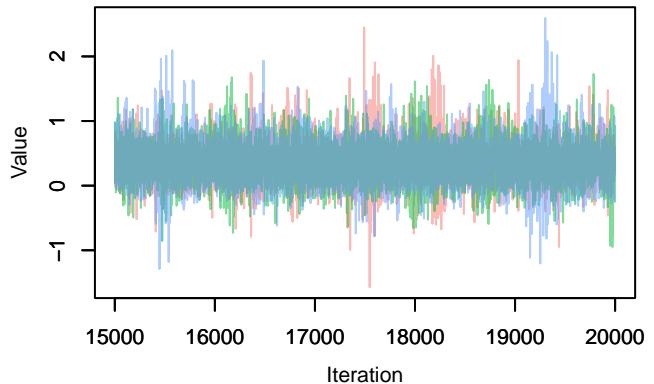
Trace – $B[(\text{Intercept}) (\text{C1}), \text{Uncinaria (S2)}]$



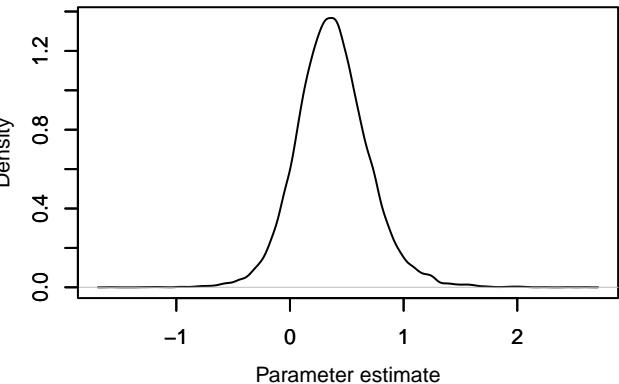
Density – $B[(\text{Intercept}) (\text{C1}), \text{Uncinaria (S2)}]$



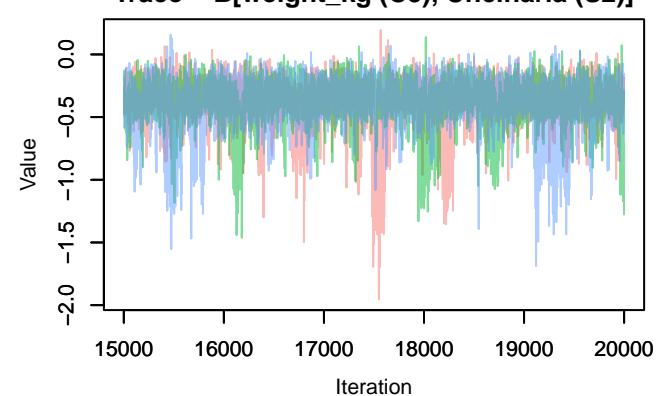
Trace – $B[\text{sexmale (C2)}, \text{Uncinaria (S2)}]$



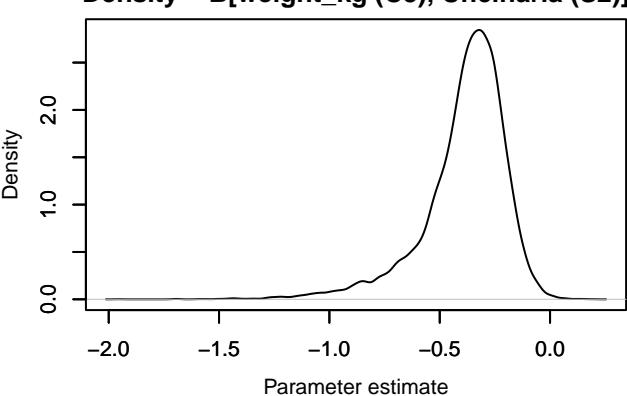
Density – $B[\text{sexmale (C2)}, \text{Uncinaria (S2)}]$



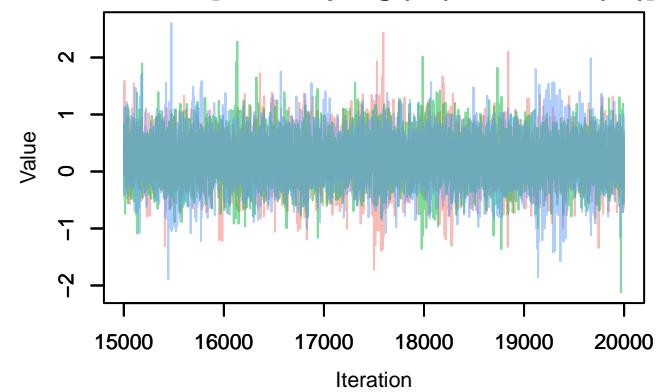
Trace – B[weight_kg (C3), Uncinaria (S2)]



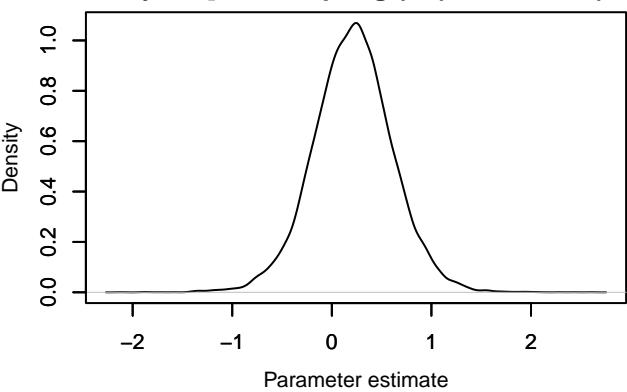
Density – B[weight_kg (C3), Uncinaria (S2)]



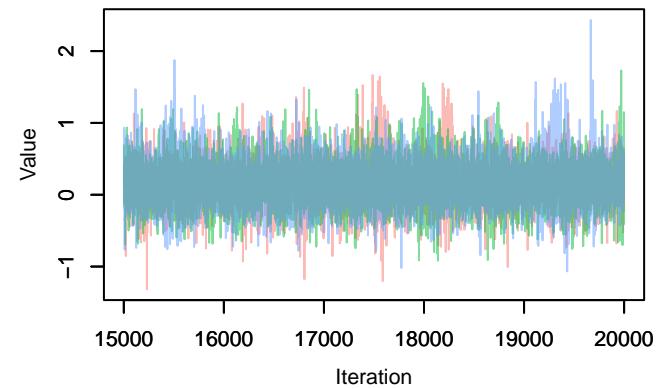
Trace – B[seasonspring (C4), Uncinaria (S2)]



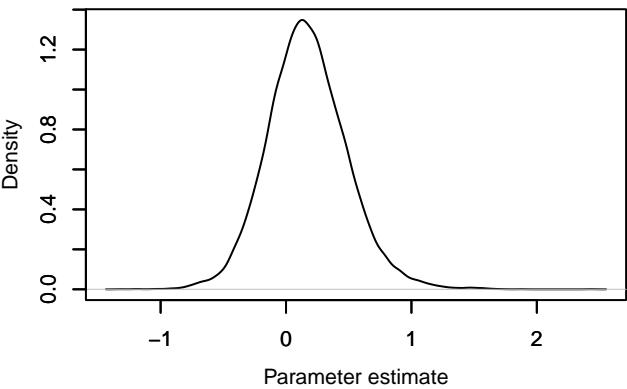
Density – B[seasonspring (C4), Uncinaria (S2)]



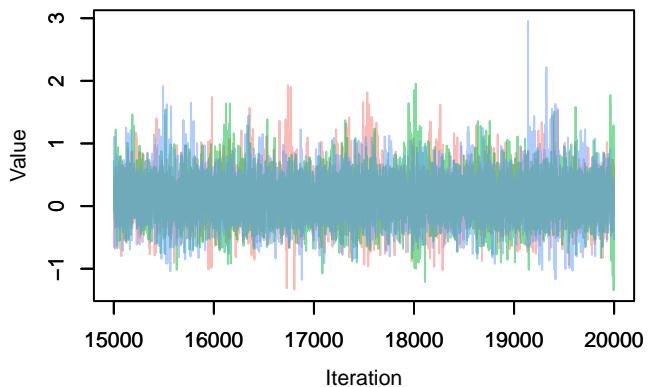
Trace – B[seasonwinter (C5), Uncinaria (S2)]



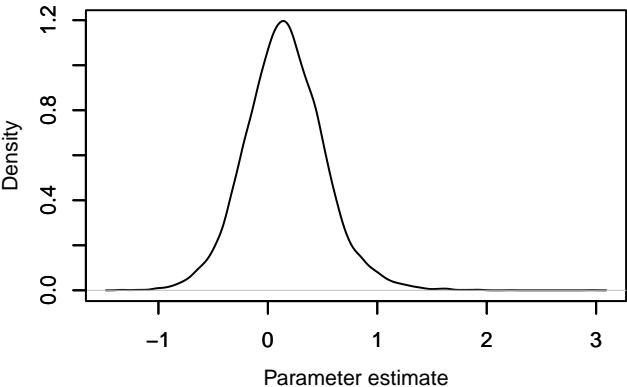
Density – B[seasonwinter (C5), Uncinaria (S2)]



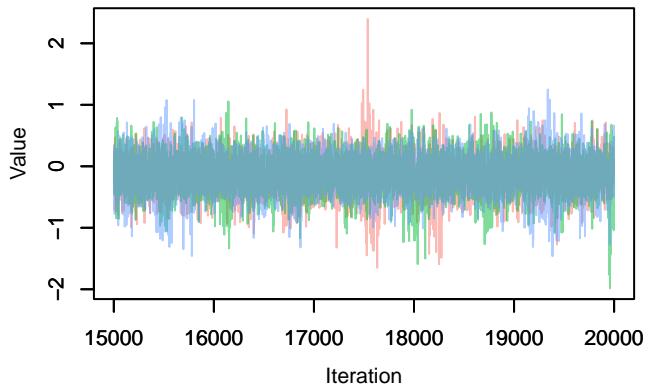
Trace – $B[\text{areaBrandenburg (C6), Uncinaria (S2)}]$



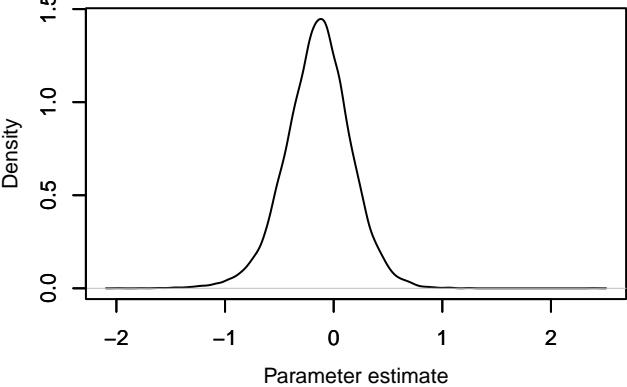
Density – $B[\text{areaBrandenburg (C6), Uncinaria (S2)}]$



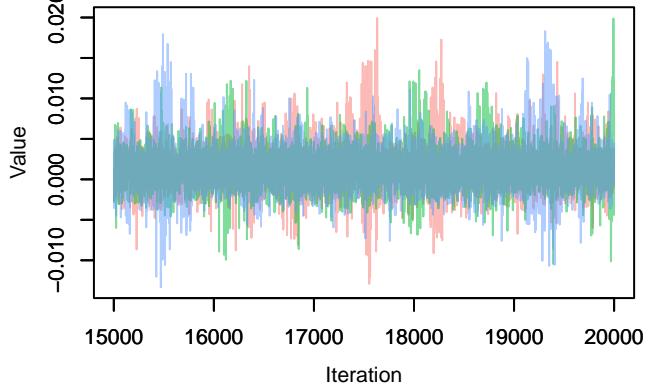
Trace – $B[\text{conditionexcellent (C7), Uncinaria (S2)}]$



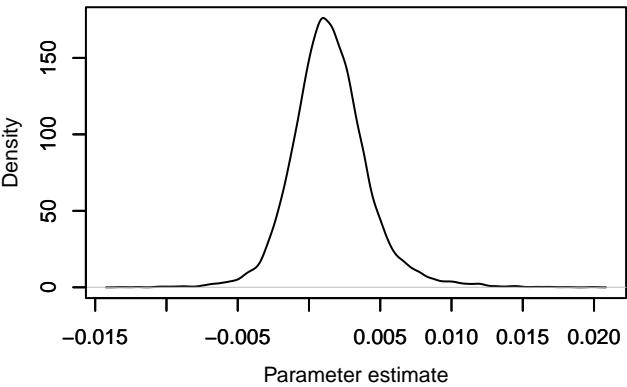
Density – $B[\text{conditionexcellent (C7), Uncinaria (S2)}]$



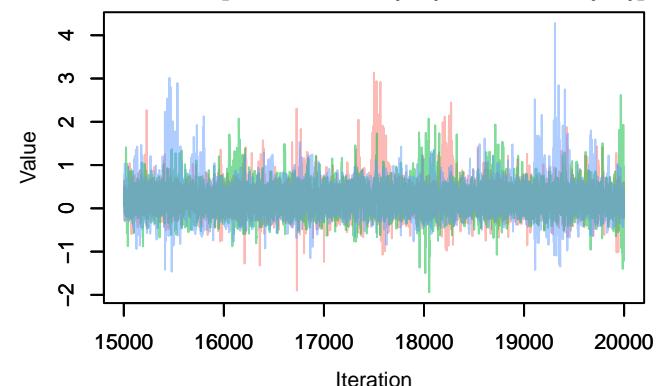
Trace – $B[\text{DNAng.ul (C8), Uncinaria (S2)}]$



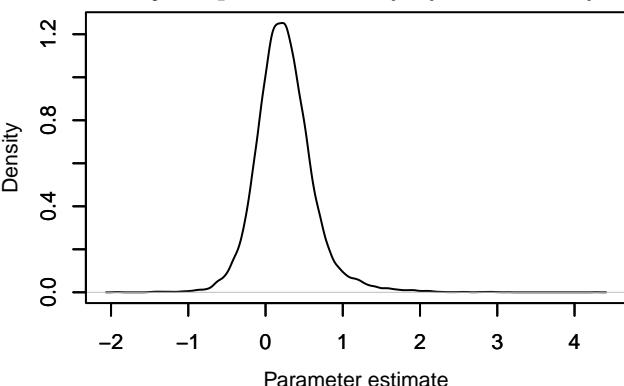
Density – $B[\text{DNAng.ul (C8), Uncinaria (S2)}]$



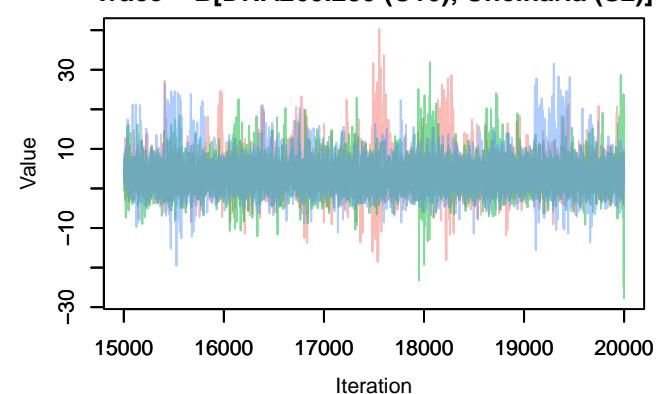
Trace – $B[\text{DNA260.230 (C9), Uncinaria (S2)}]$



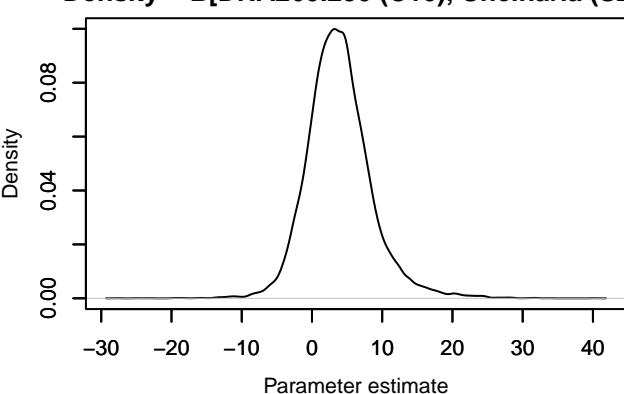
Density – $B[\text{DNA260.230 (C9), Uncinaria (S2)}]$



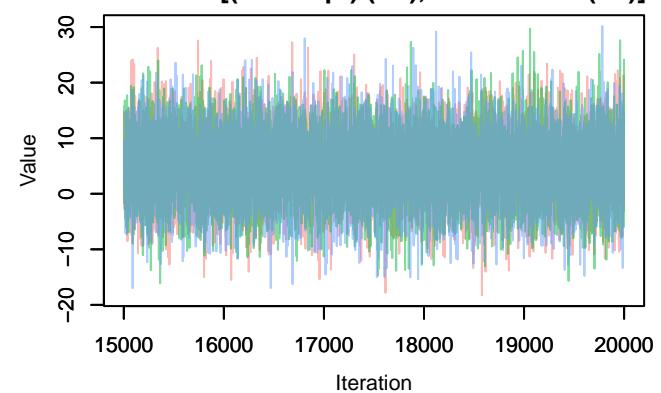
Trace – $B[\text{DNA260.280 (C10), Uncinaria (S2)}]$



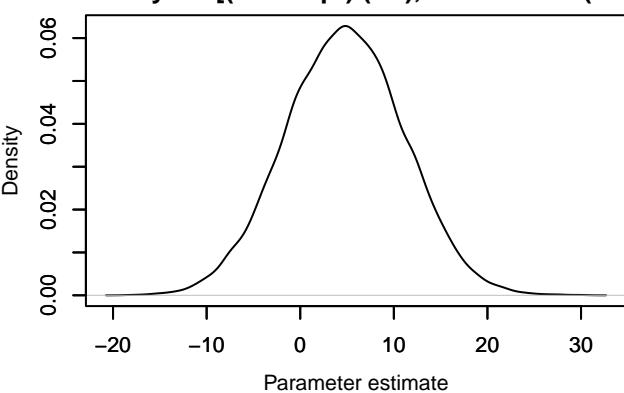
Density – $B[\text{DNA260.280 (C10), Uncinaria (S2)}]$



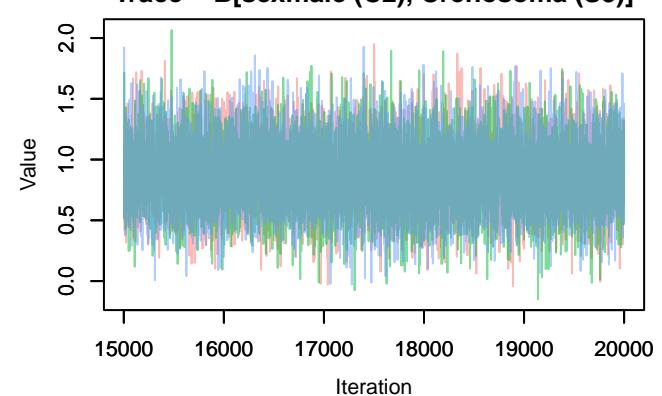
Trace – $B[(\text{Intercept}) (\text{C1}), \text{Crenosoma (S3)}]$



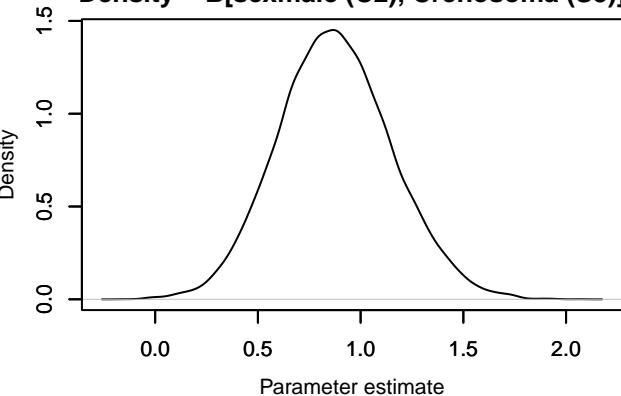
Density – $B[(\text{Intercept}) (\text{C1}), \text{Crenosoma (S3)}]$



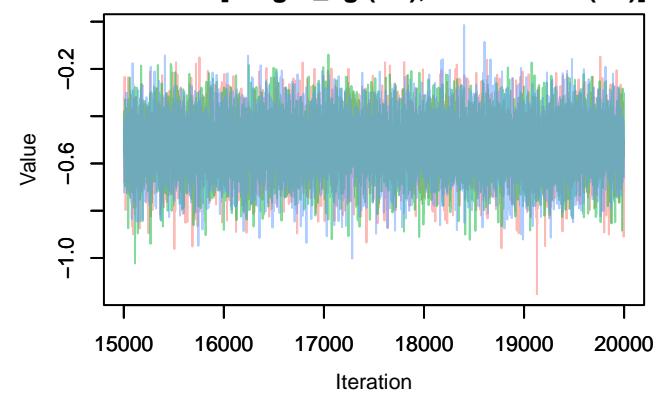
Trace – B[sexmale (C2), Crenosoma (S3)]



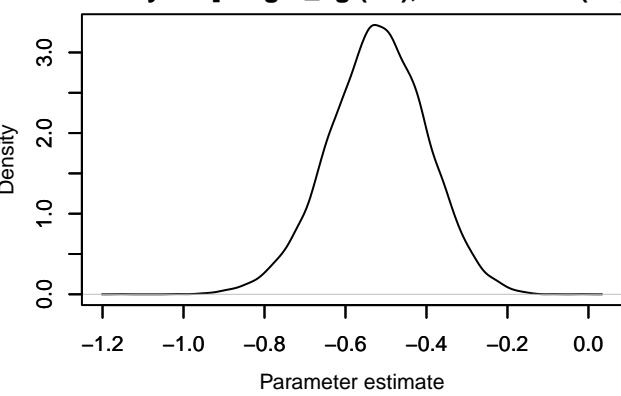
Density – B[sexmale (C2), Crenosoma (S3)]



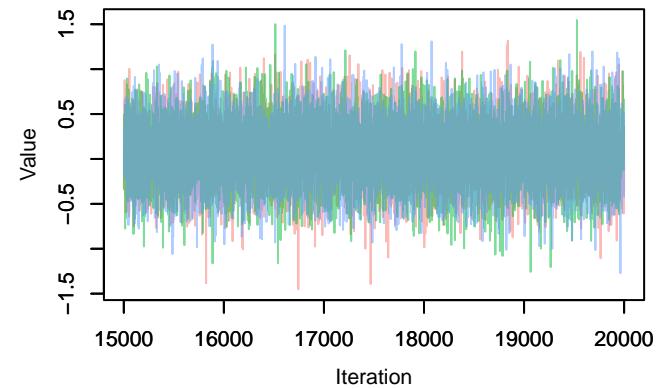
Trace – B[weight_kg (C3), Crenosoma (S3)]



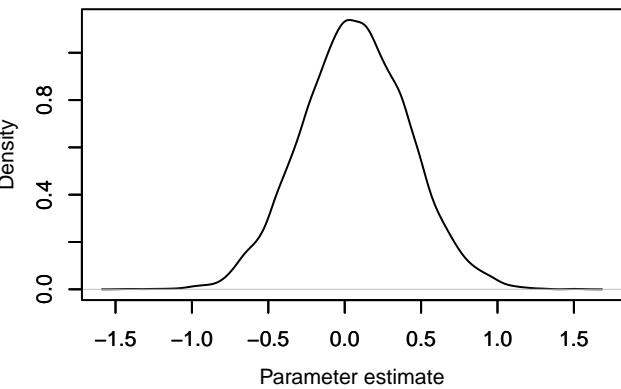
Density – B[weight_kg (C3), Crenosoma (S3)]

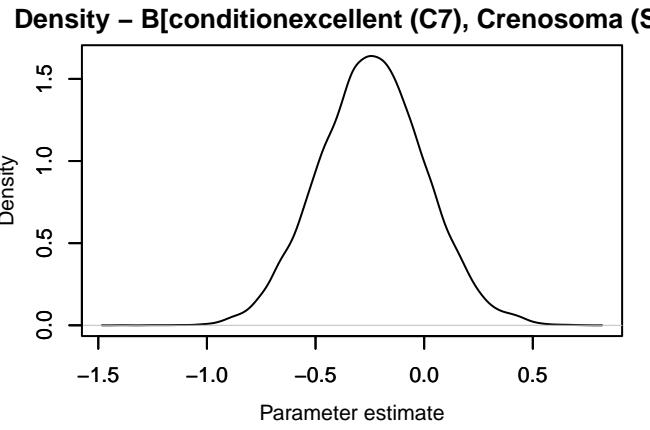
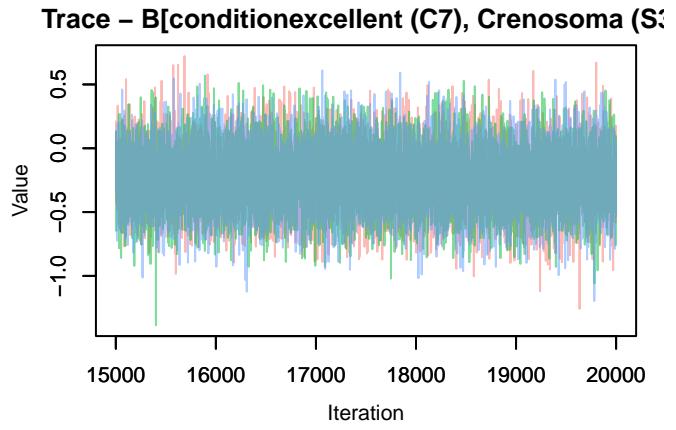
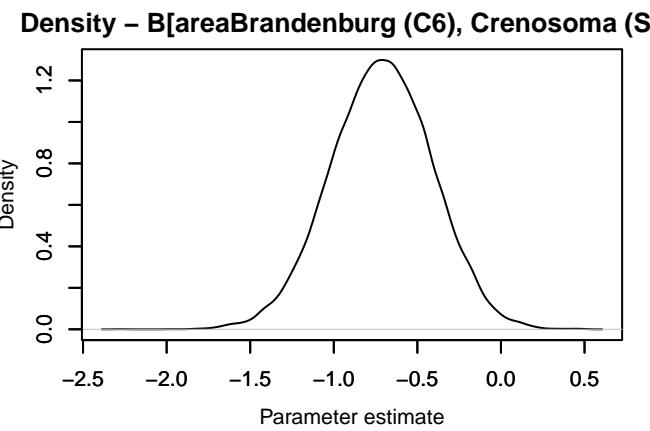
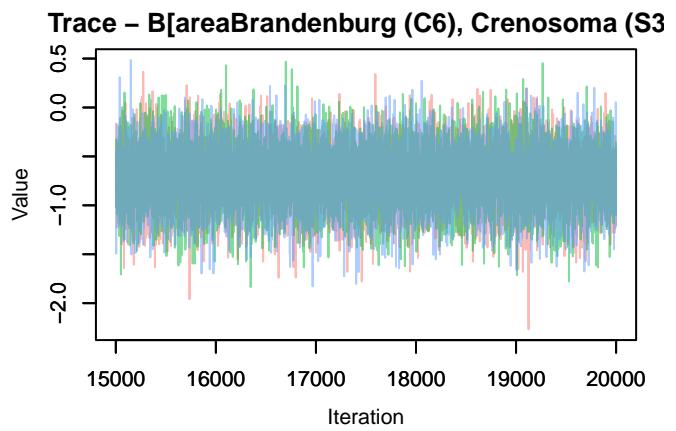
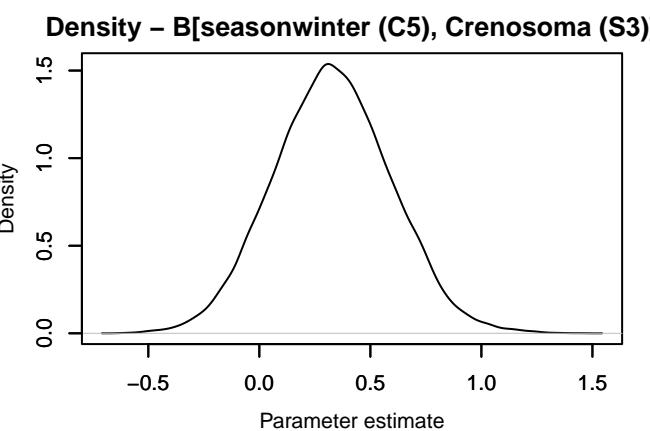
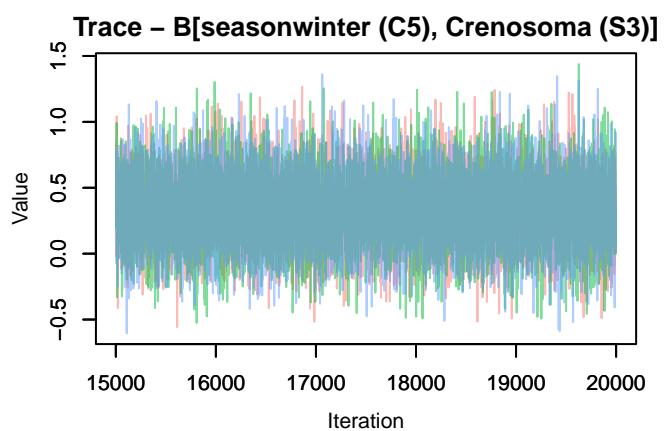


Trace – B[seasonspring (C4), Crenosoma (S3)]

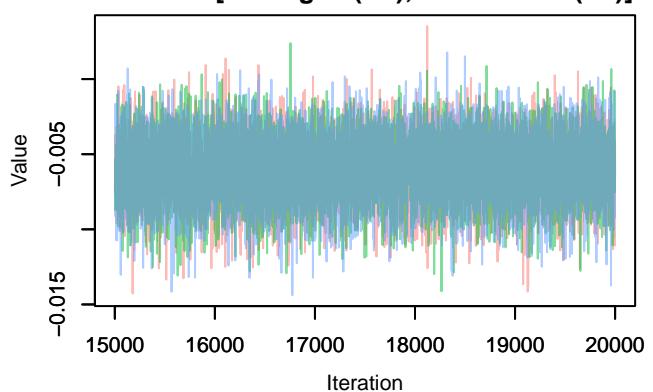


Density – B[seasonspring (C4), Crenosoma (S3)]

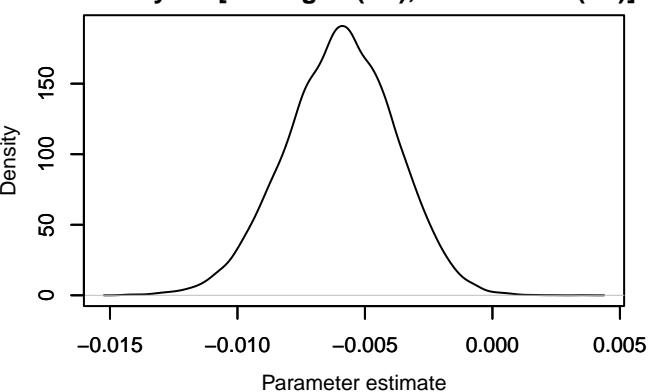




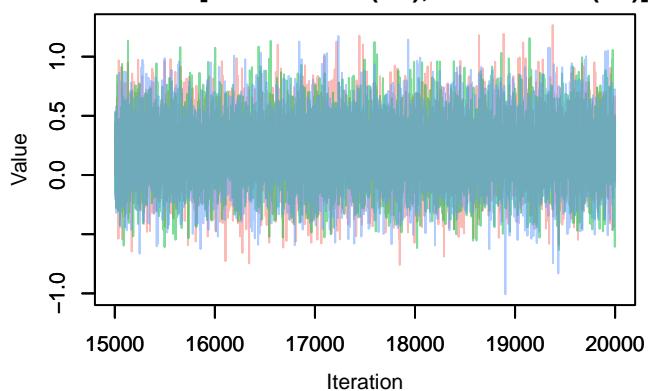
Trace – $B[\text{DNAng.ul (C8), Crenosoma (S3)}]$



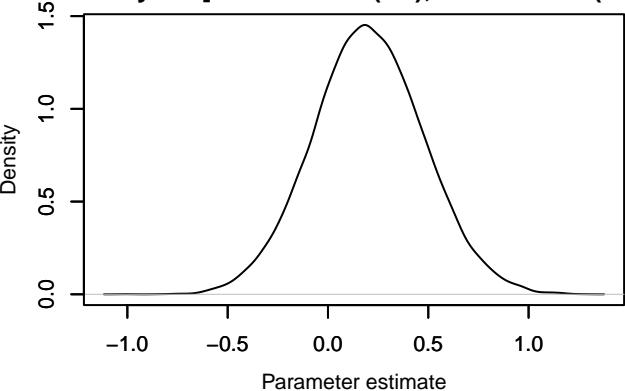
Density – $B[\text{DNAng.ul (C8), Crenosoma (S3)}]$



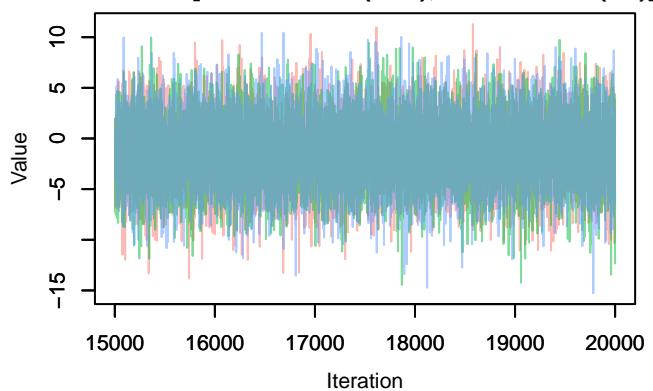
Trace – $B[\text{DNA260.230 (C9), Crenosoma (S3)}]$



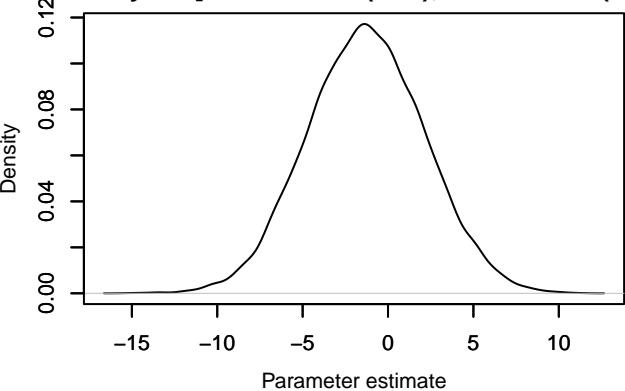
Density – $B[\text{DNA260.230 (C9), Crenosoma (S3)}]$



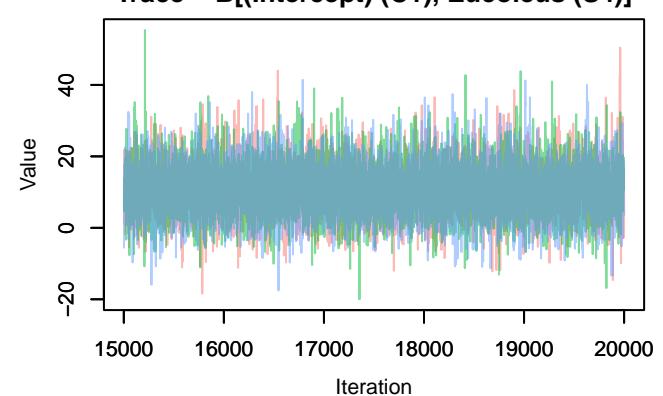
Trace – $B[\text{DNA260.280 (C10), Crenosoma (S3)}]$



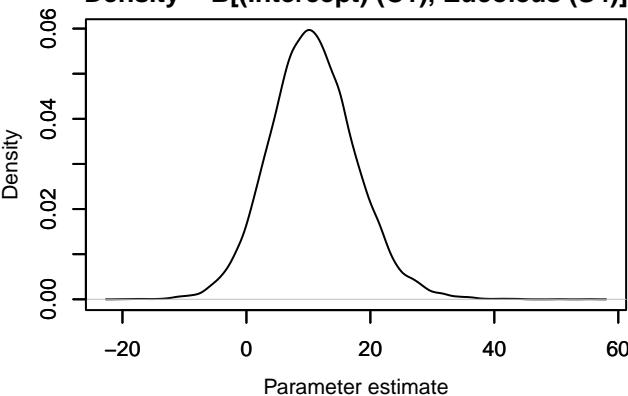
Density – $B[\text{DNA260.280 (C10), Crenosoma (S3)}]$



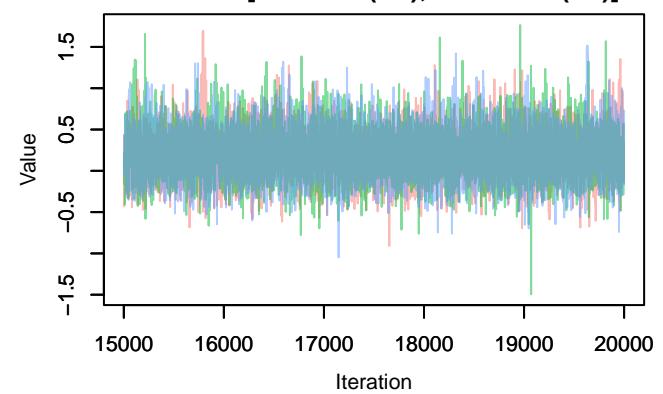
Trace – $B[(\text{Intercept}) (\text{C1}), \text{Eucoleus} (\text{S4})]$



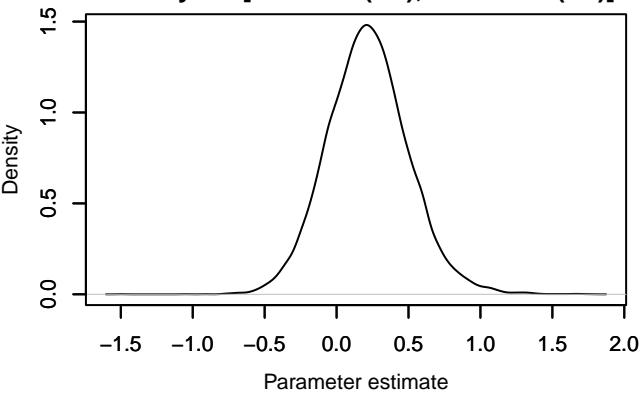
Density – $B[(\text{Intercept}) (\text{C1}), \text{Eucoleus} (\text{S4})]$



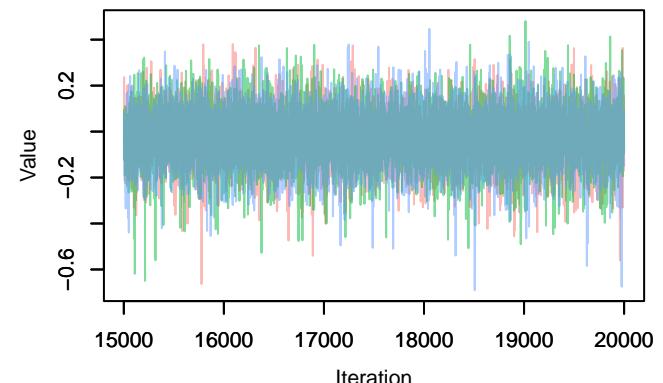
Trace – $B[\text{sexmale} (\text{C2}), \text{Eucoleus} (\text{S4})]$



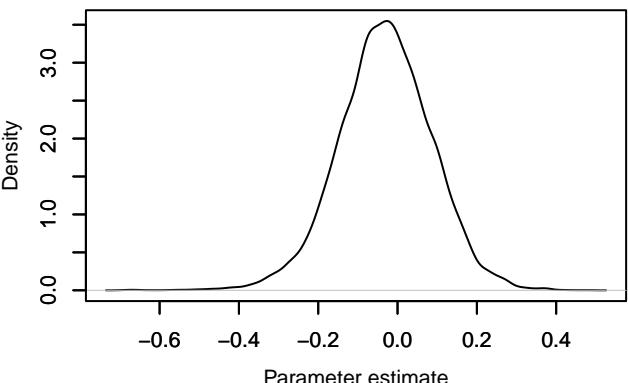
Density – $B[\text{sexmale} (\text{C2}), \text{Eucoleus} (\text{S4})]$

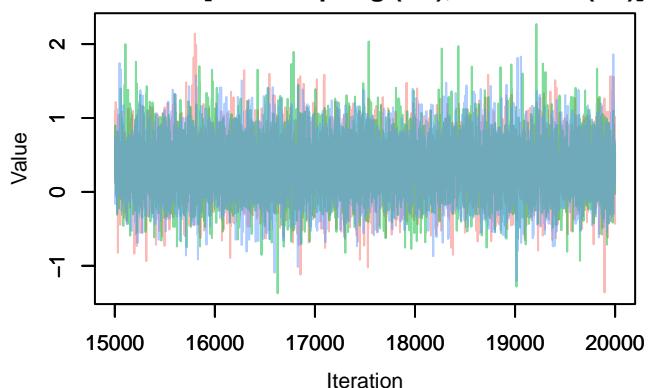
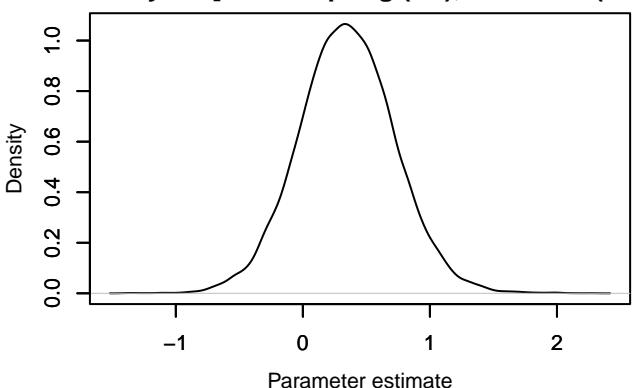
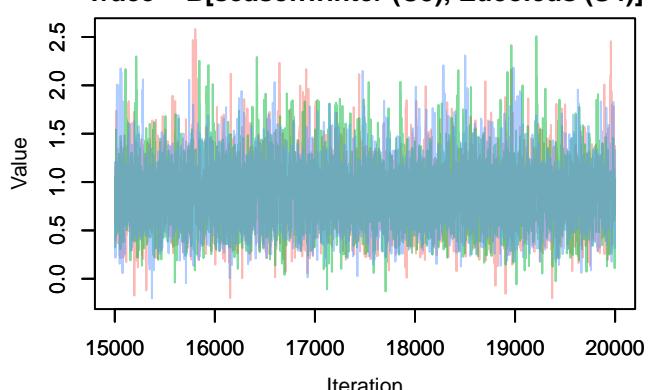
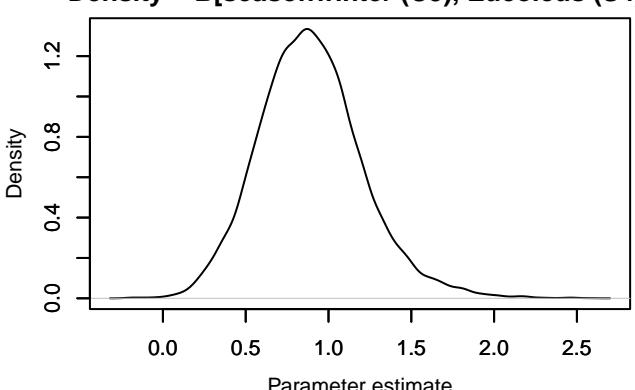
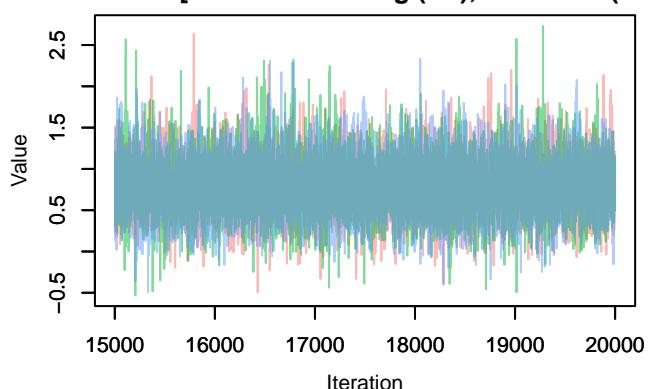
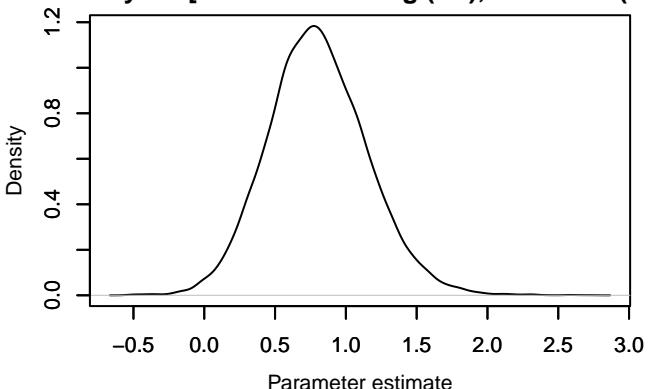


Trace – $B[\text{weight_kg} (\text{C3}), \text{Eucoleus} (\text{S4})]$

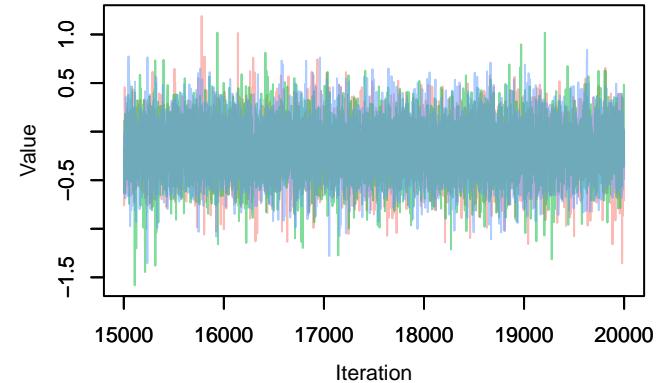


Density – $B[\text{weight_kg} (\text{C3}), \text{Eucoleus} (\text{S4})]$

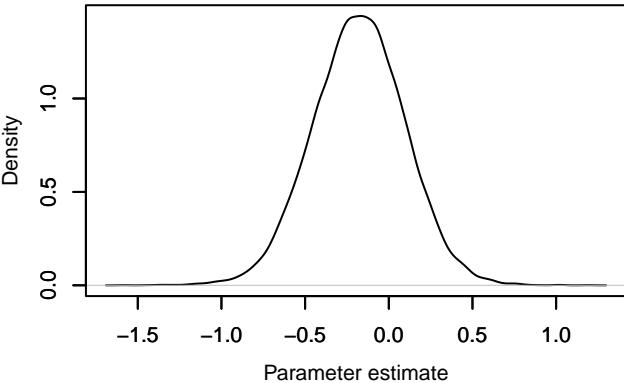


Trace – $B[\text{seasonspring (C4)}, \text{Eucoleus (S4)}]$ Density – $B[\text{seasonspring (C4)}, \text{Eucoleus (S4)}]$ Trace – $B[\text{seasonwinter (C5)}, \text{Eucoleus (S4)}]$ Density – $B[\text{seasonwinter (C5)}, \text{Eucoleus (S4)}]$ Trace – $B[\text{areaBrandenburg (C6)}, \text{Eucoleus (S4)}]$ Density – $B[\text{areaBrandenburg (C6)}, \text{Eucoleus (S4)}]$ 

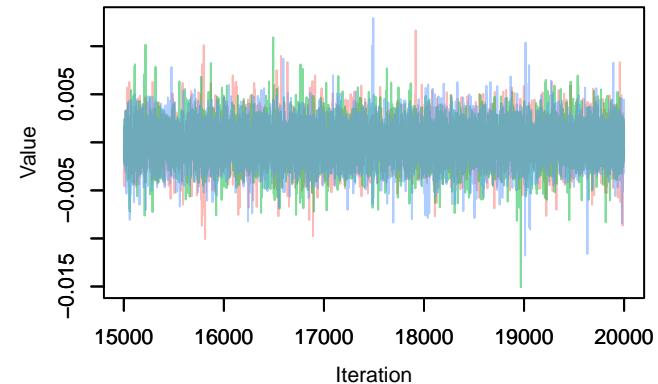
Trace – B[conditionexcellent (C7), Eucoleus (S4)]



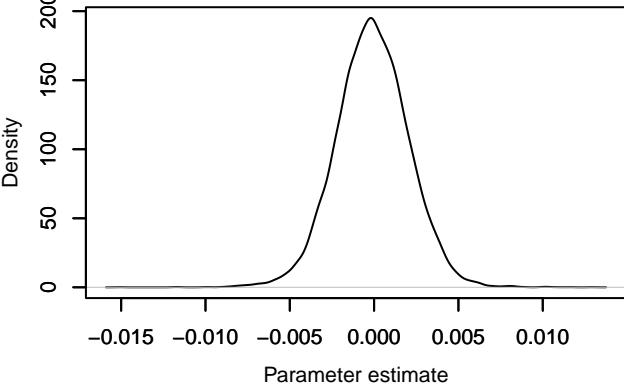
Density – B[conditionexcellent (C7), Eucoleus (S4)]



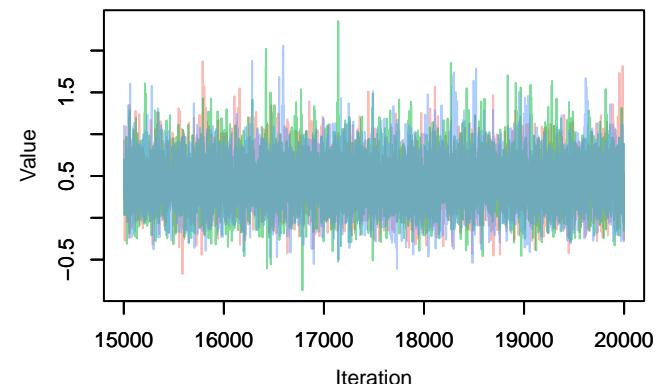
Trace – B[DNAng.ul (C8), Eucoleus (S4)]



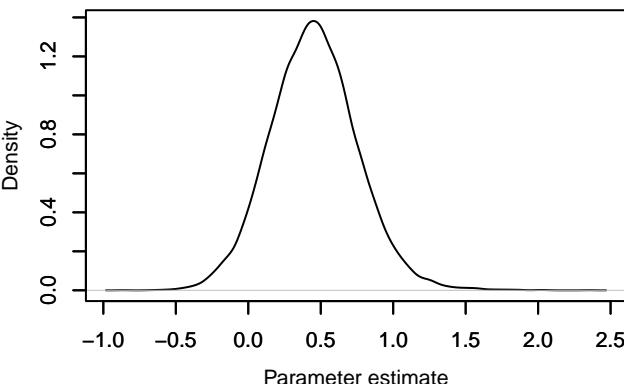
Density – B[DNAng.ul (C8), Eucoleus (S4)]



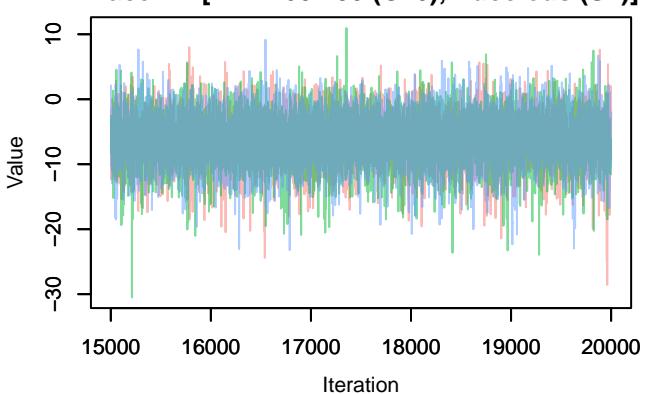
Trace – B[DNA260.230 (C9), Eucoleus (S4)]



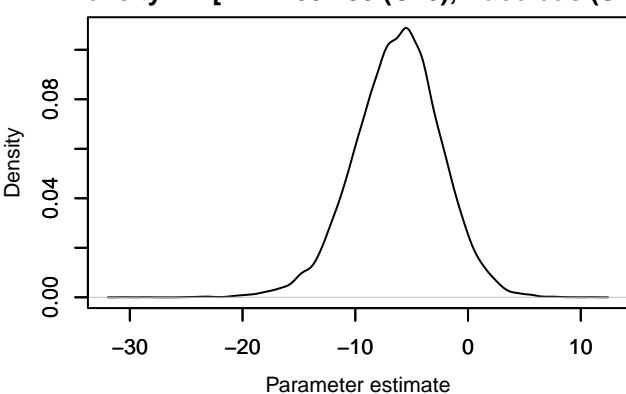
Density – B[DNA260.230 (C9), Eucoleus (S4)]



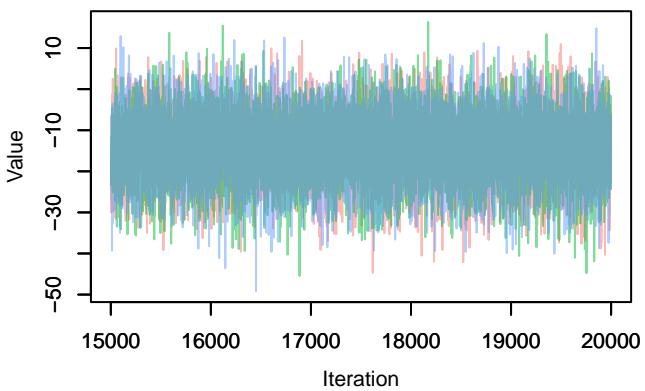
Trace – $B[\text{DNA260.280 (C10)}, \text{Eucoleus (S4)}]$



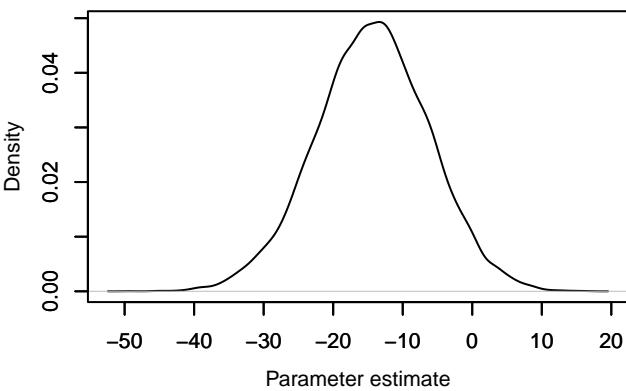
Density – $B[\text{DNA260.280 (C10)}, \text{Eucoleus (S4)}]$



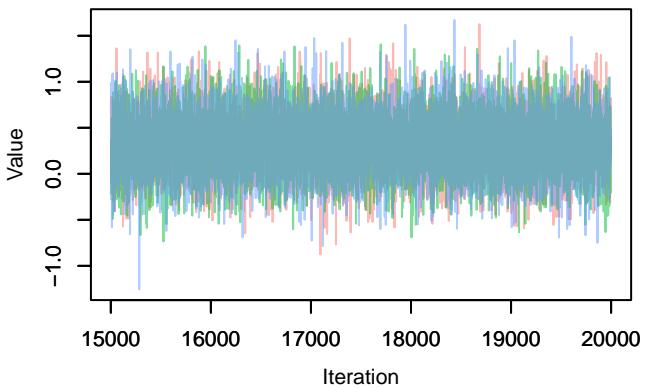
Trace – $B[(\text{Intercept}) (\text{C1})], \text{Alaria (S5)}$



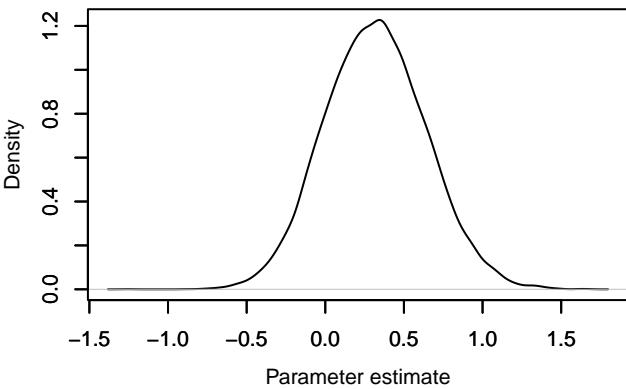
Density – $B[(\text{Intercept}) (\text{C1})], \text{Alaria (S5)}$



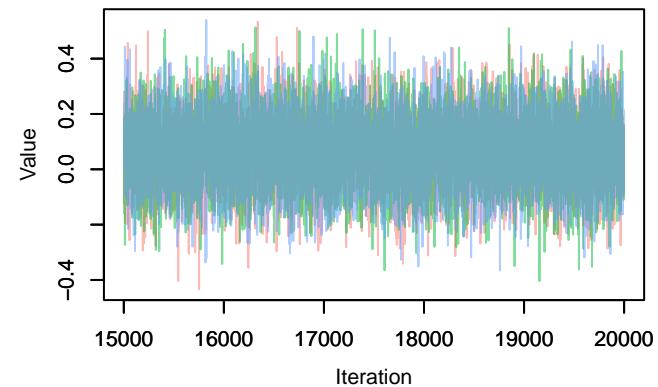
Trace – $B[\text{sexmale (C2)}], \text{Alaria (S5)}$



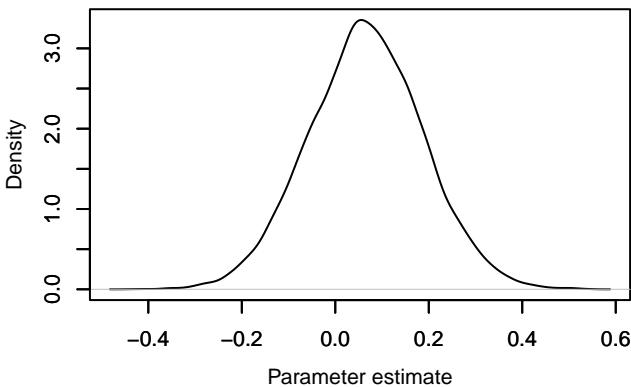
Density – $B[\text{sexmale (C2)}], \text{Alaria (S5)}$



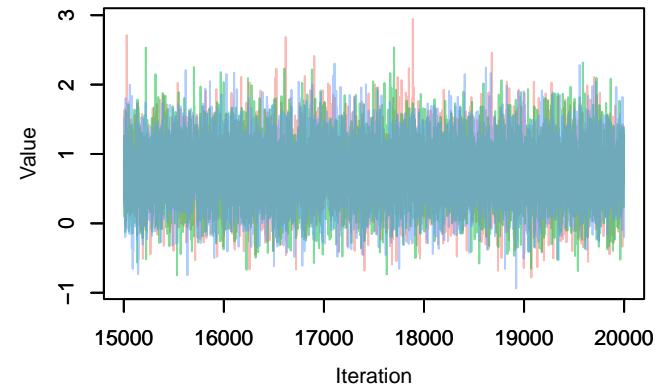
Trace – $B[\text{weight_kg} \text{ (C3), Alaria (S5)}]$



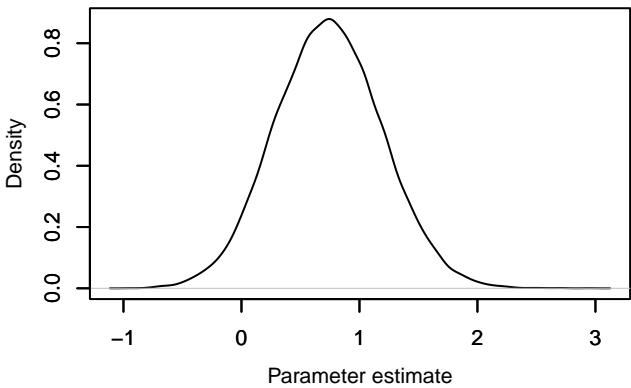
Density – $B[\text{weight_kg} \text{ (C3), Alaria (S5)}]$



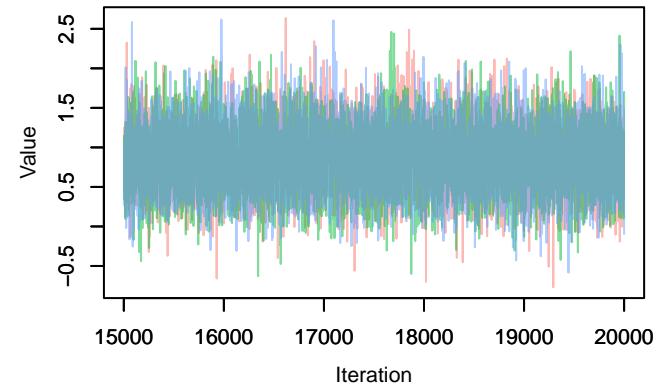
Trace – $B[\text{seasonspring} \text{ (C4), Alaria (S5)}]$



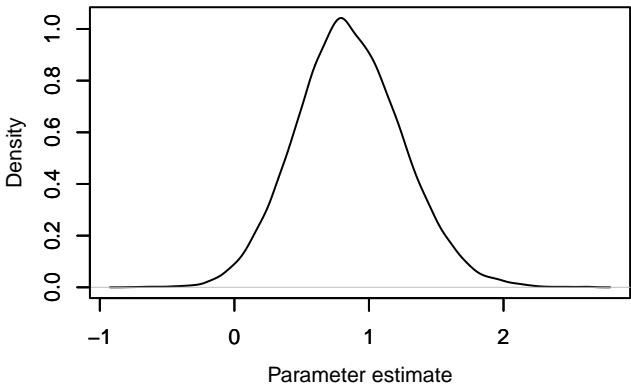
Density – $B[\text{seasonspring} \text{ (C4), Alaria (S5)}]$

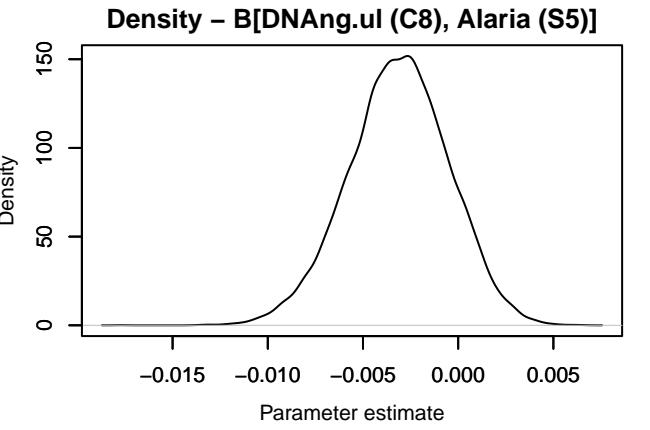
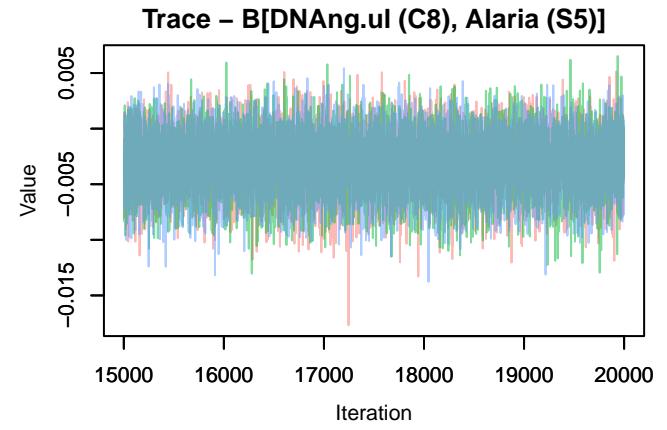
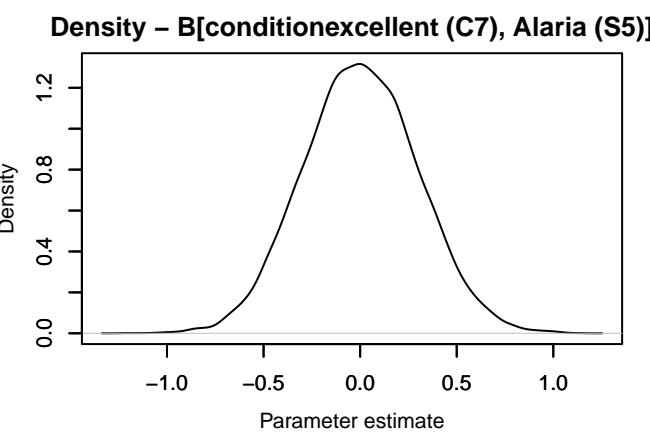
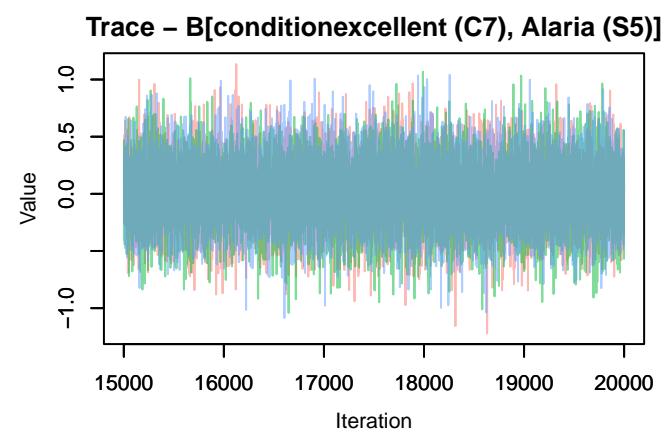
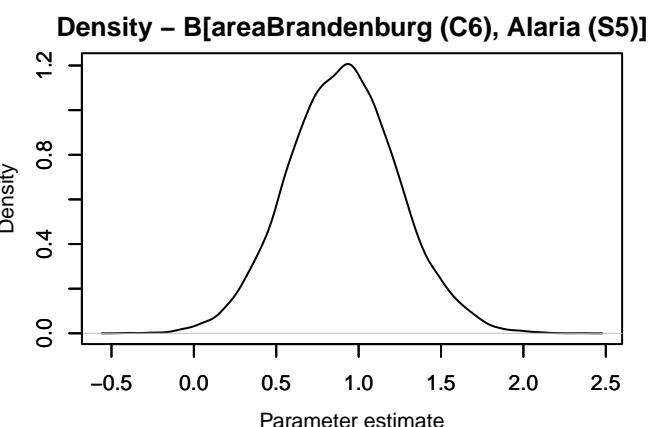
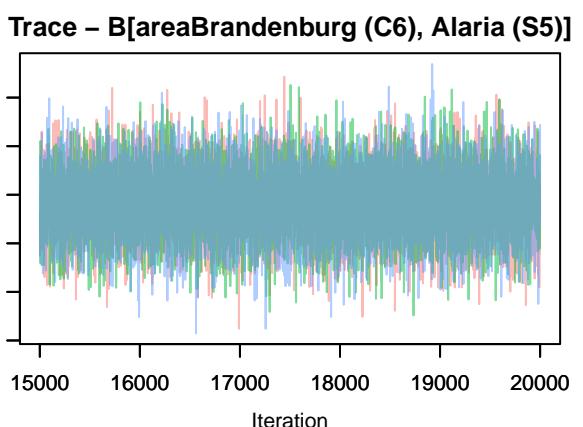


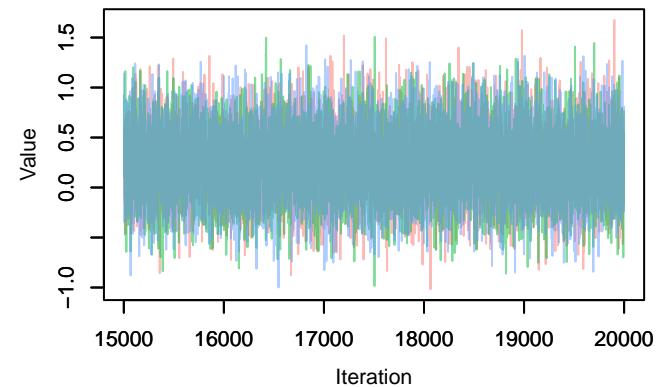
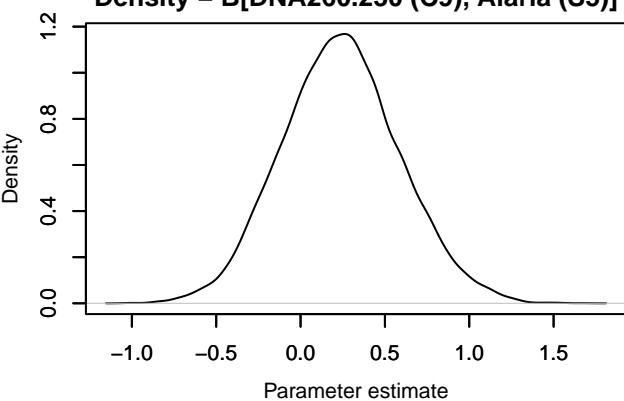
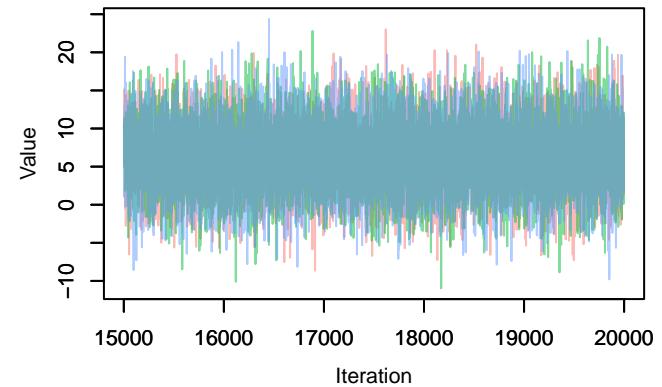
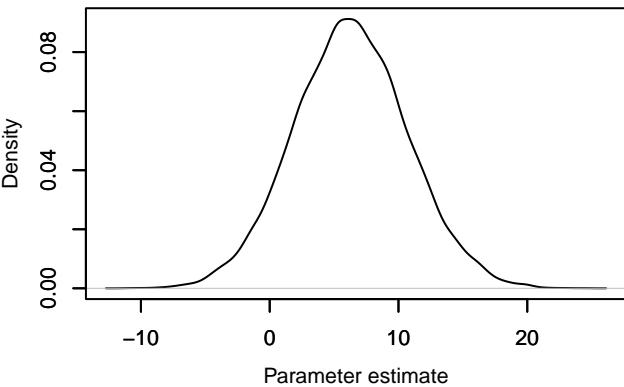
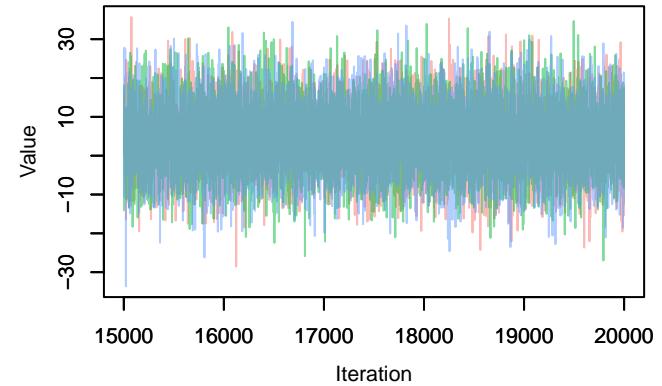
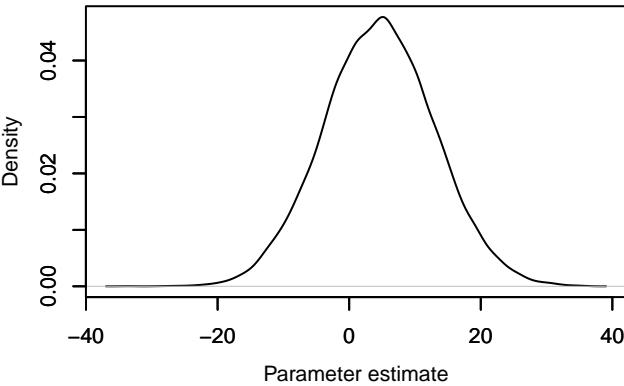
Trace – $B[\text{seasonwinter} \text{ (C5), Alaria (S5)}]$



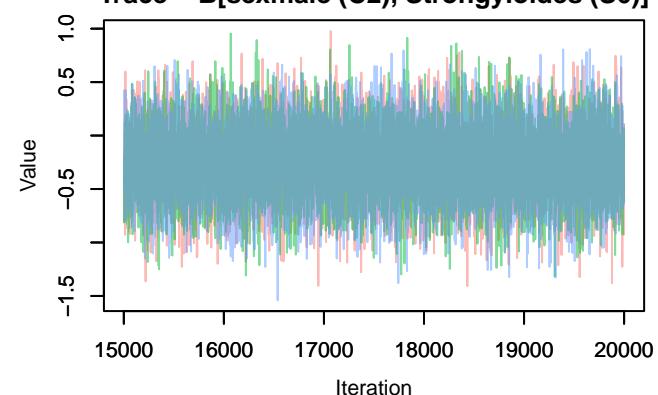
Density – $B[\text{seasonwinter} \text{ (C5), Alaria (S5)}]$



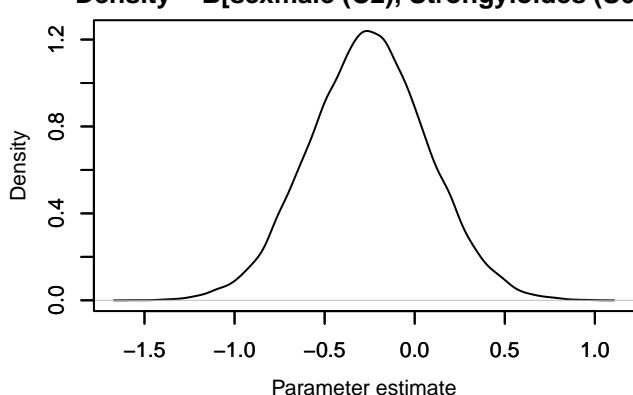


Trace – $B[\text{DNA260.230 (C9), Alaria (S5)}]$ Density – $B[\text{DNA260.230 (C9), Alaria (S5)}]$ Trace – $B[\text{DNA260.280 (C10), Alaria (S5)}]$ Density – $B[\text{DNA260.280 (C10), Alaria (S5)}]$ Trace – $B[(\text{Intercept}) (\text{C1}), \text{Strongyloides (S6)}]$ Density – $B[(\text{Intercept}) (\text{C1}), \text{Strongyloides (S6)}]$ 

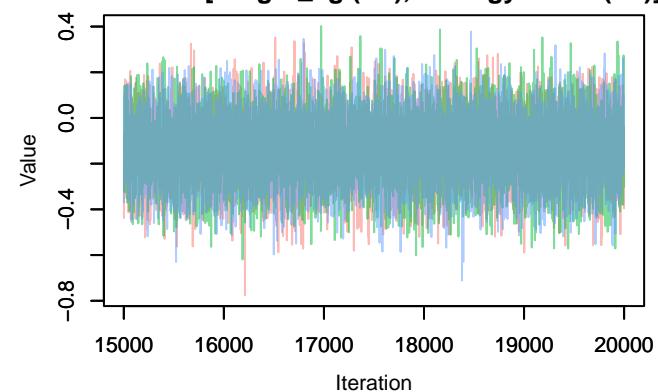
Trace – $B[\text{sexmale (C2)}, \text{Strongyloides (S6)}]$



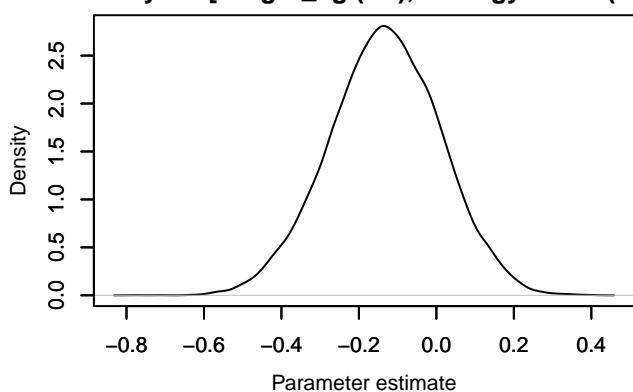
Density – $B[\text{sexmale (C2)}, \text{Strongyloides (S6)}]$



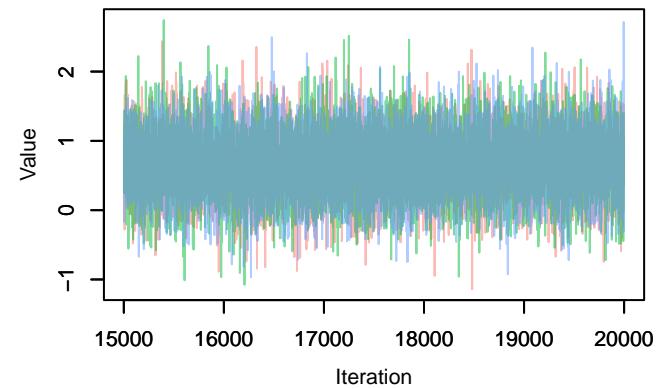
Trace – $B[\text{weight_kg (C3)}, \text{Strongyloides (S6)}]$



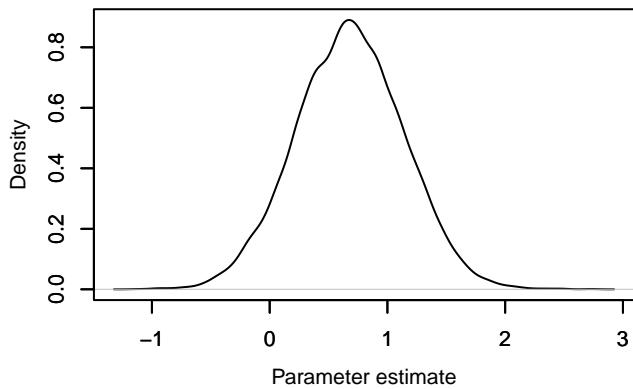
Density – $B[\text{weight_kg (C3)}, \text{Strongyloides (S6)}]$



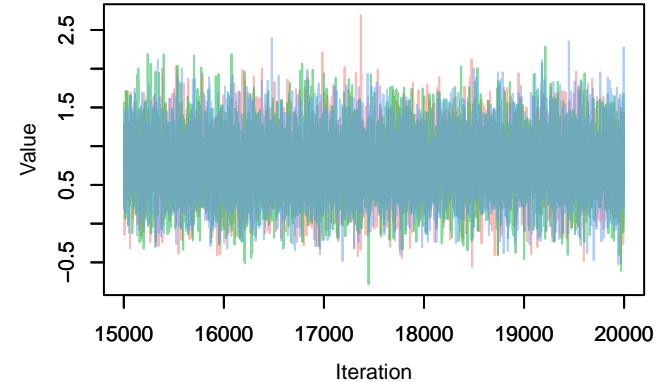
Trace – $B[\text{seasonspring (C4)}, \text{Strongyloides (S6)}]$



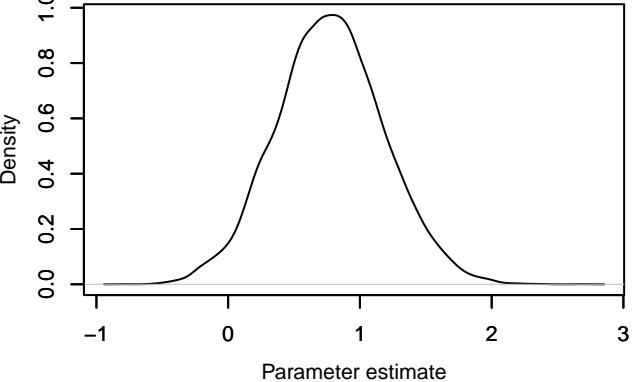
Density – $B[\text{seasonspring (C4)}, \text{Strongyloides (S6)}]$



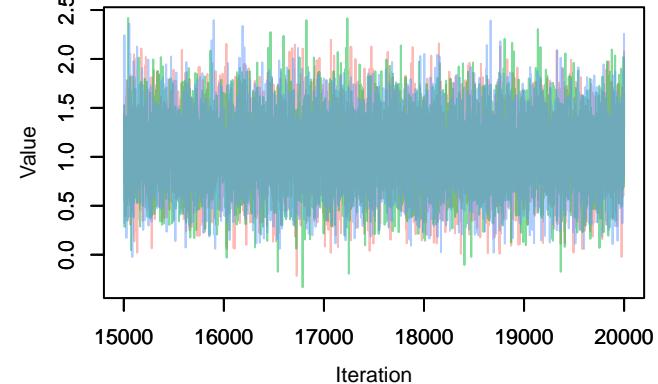
Trace – $B[\text{seasonwinter (C5), Strongyloides (S6)}$



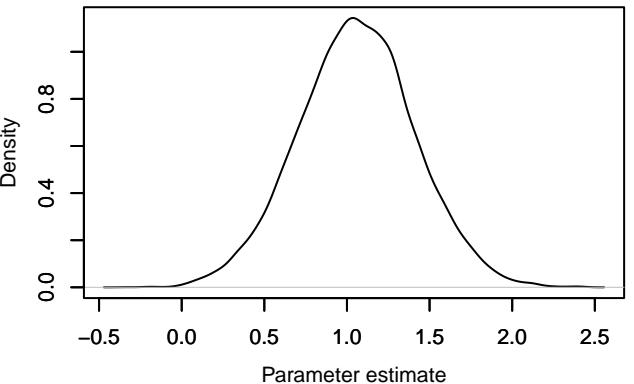
Density – $B[\text{seasonwinter (C5), Strongyloides (S6)}$



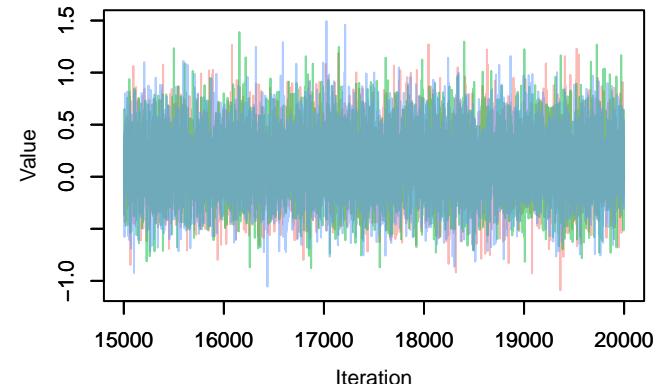
Trace – $B[\text{areaBrandenburg (C6), Strongyloides (S6)}$



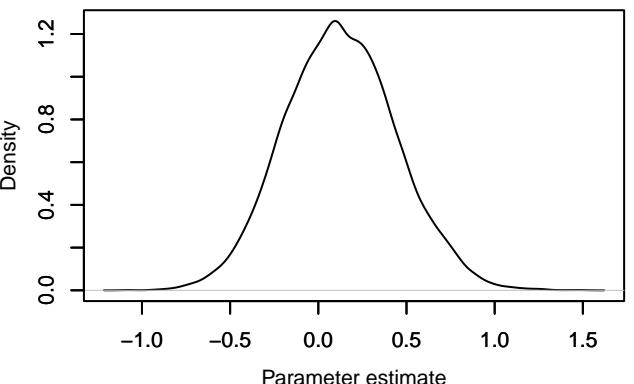
Density – $B[\text{areaBrandenburg (C6), Strongyloides (S6)}$

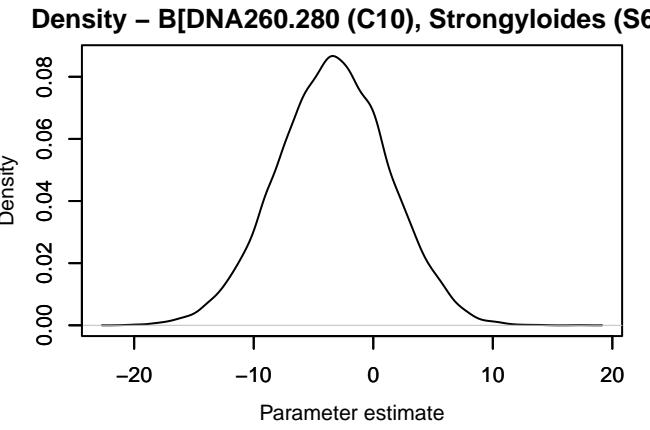
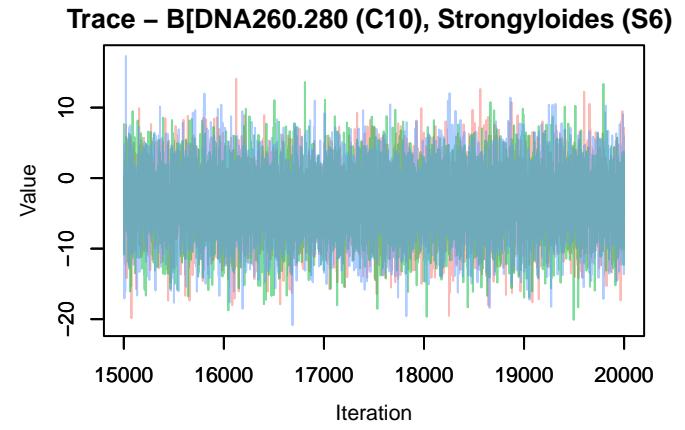
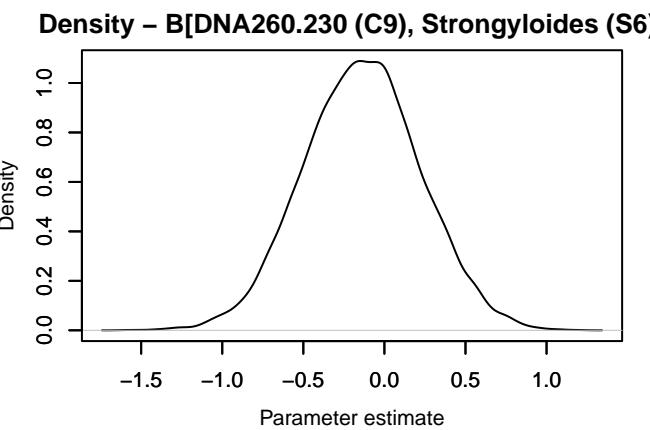
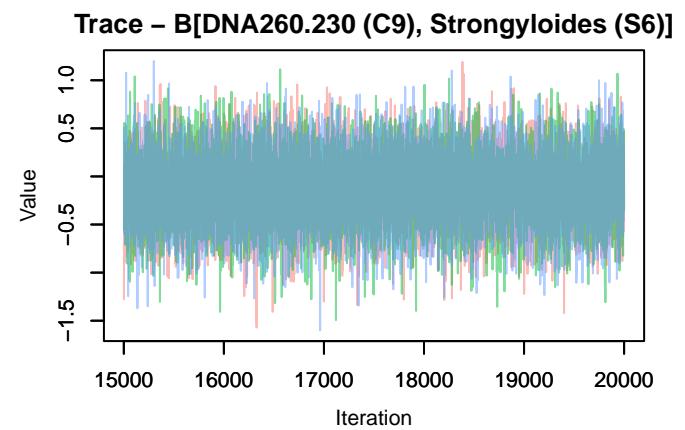
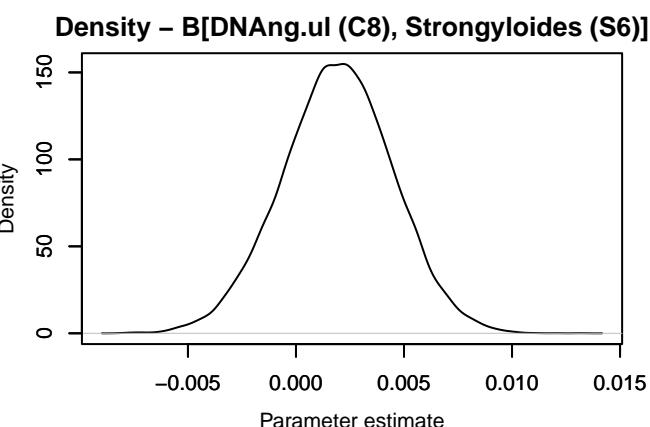
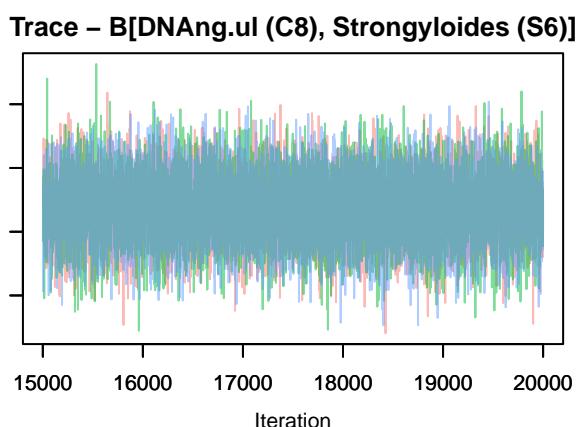


Trace – $B[\text{conditionexcellent (C7), Strongyloides (S6)}$

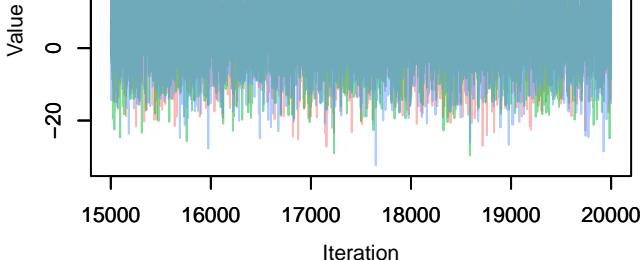


Density – $B[\text{conditionexcellent (C7), Strongyloides (S6)}$

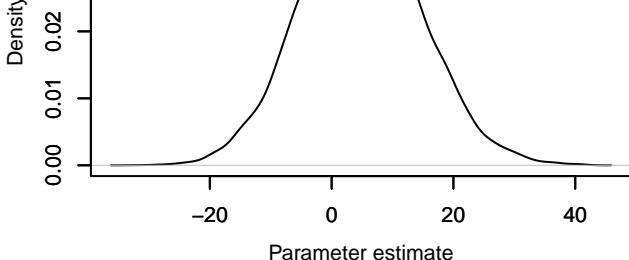




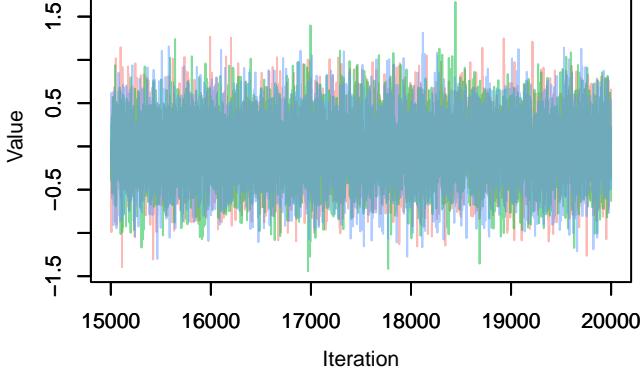
Trace – $B[(\text{Intercept}) (\text{C1}), \text{Pearsonema} (\text{S7})]$



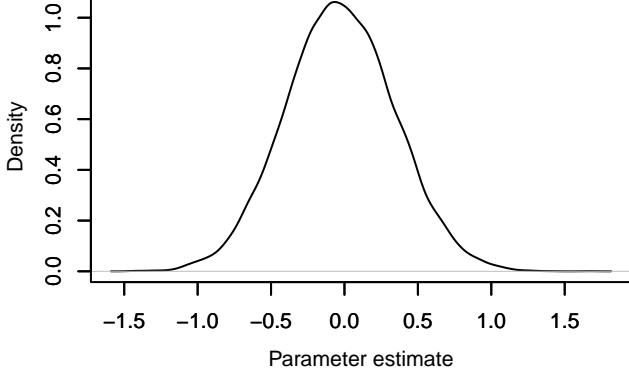
Density – $B[(\text{Intercept}) (\text{C1}), \text{Pearsonema} (\text{S7})]$



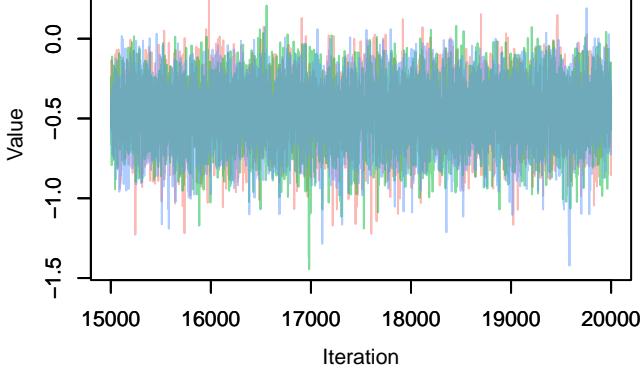
Trace – $B[\text{sexmale} (\text{C2}), \text{Pearsonema} (\text{S7})]$



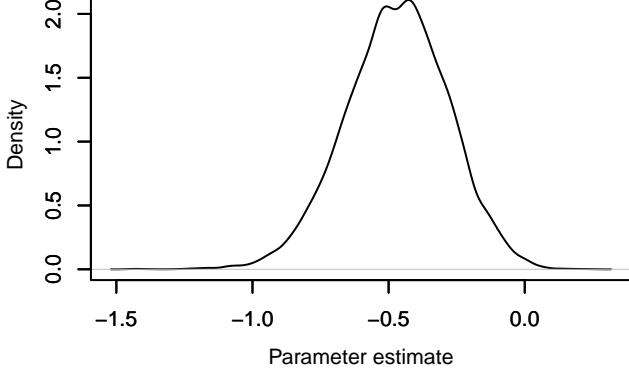
Density – $B[\text{sexmale} (\text{C2}), \text{Pearsonema} (\text{S7})]$



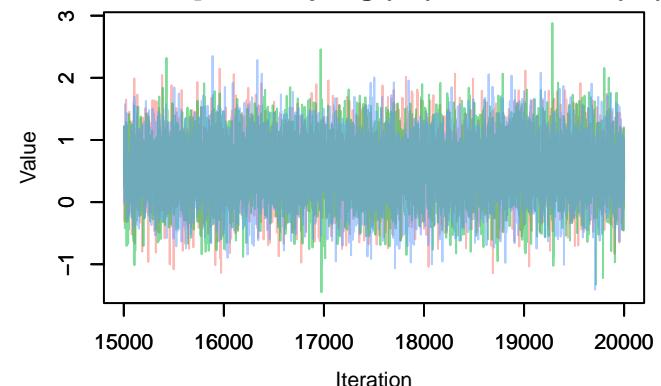
Trace – $B[\text{weight_kg} (\text{C3}), \text{Pearsonema} (\text{S7})]$



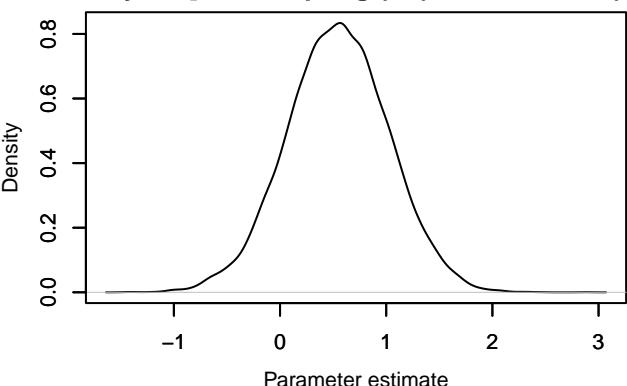
Density – $B[\text{weight_kg} (\text{C3}), \text{Pearsonema} (\text{S7})]$



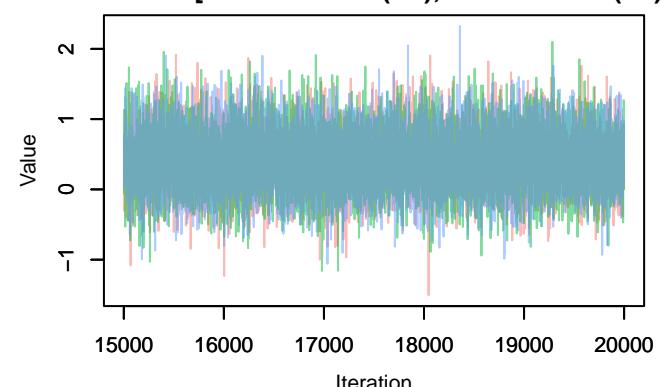
Trace – $B[\text{seasonspring (C4)}, \text{Pearsonema (S7)}]$



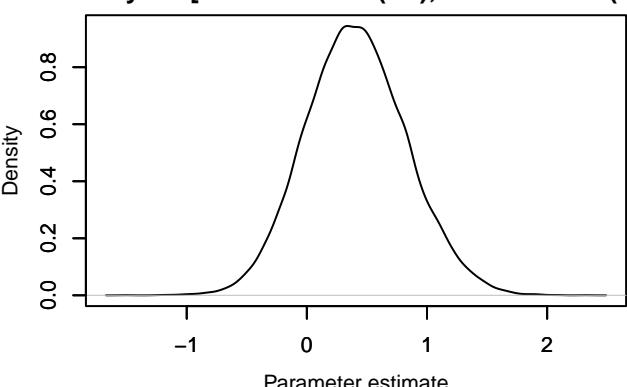
Density – $B[\text{seasonspring (C4)}, \text{Pearsonema (S7)}]$



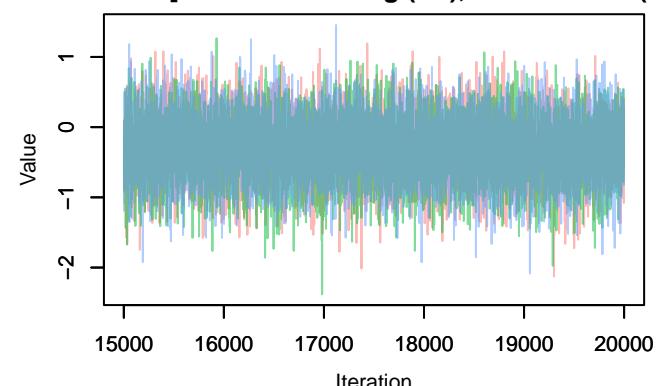
Trace – $B[\text{seasonwinter (C5)}, \text{Pearsonema (S7)}]$



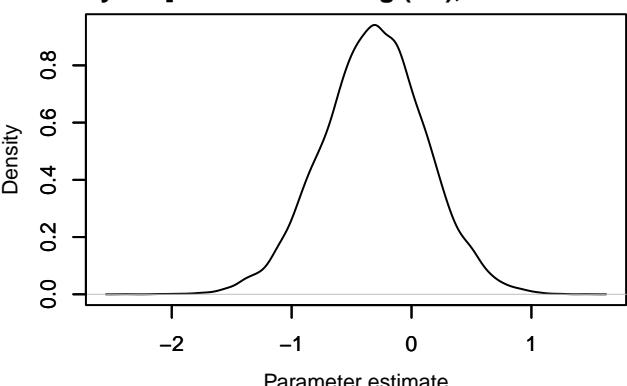
Density – $B[\text{seasonwinter (C5)}, \text{Pearsonema (S7)}]$



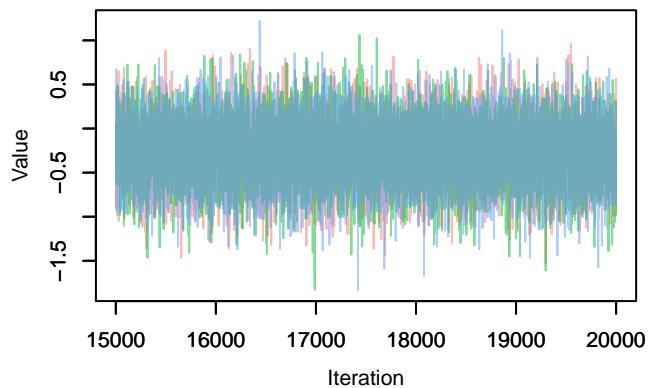
Trace – $B[\text{areaBrandenburg (C6)}, \text{Pearsonema (S7)}]$



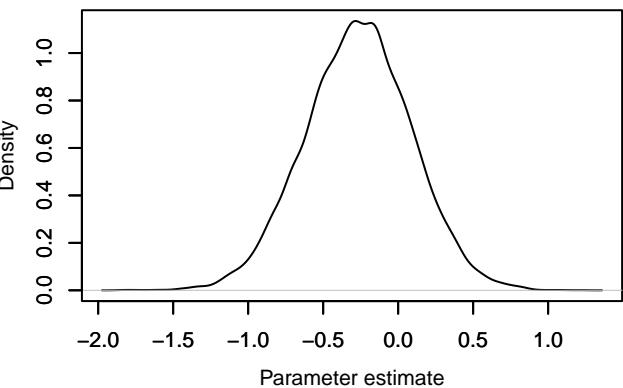
Density – $B[\text{areaBrandenburg (C6)}, \text{Pearsonema (S7)}]$



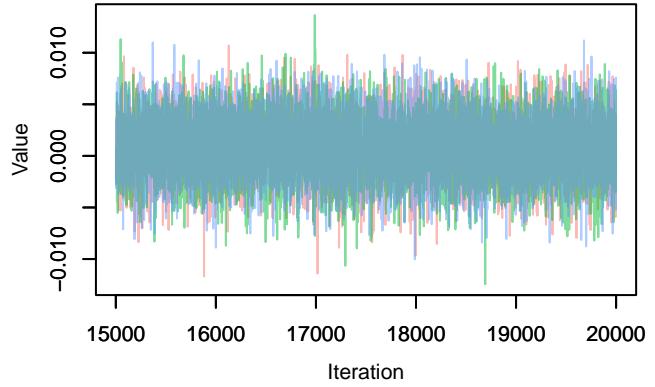
Trace – $B[\text{conditionexcellent (C7)}, \text{Pearsonema (S)}$



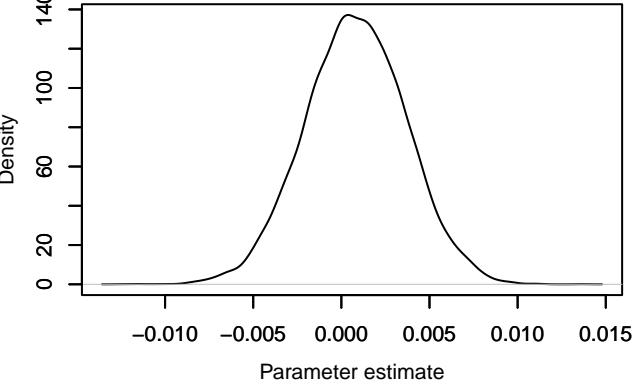
Density – $B[\text{conditionexcellent (C7)}, \text{Pearsonema (S)}$



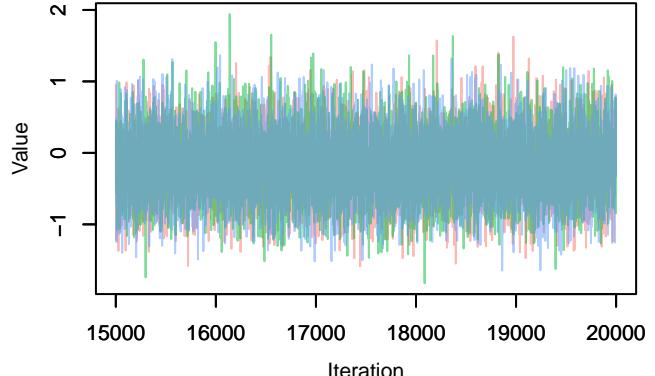
Trace – $B[\text{DNAng.ul (C8)}, \text{Pearsonema (S7)}]$



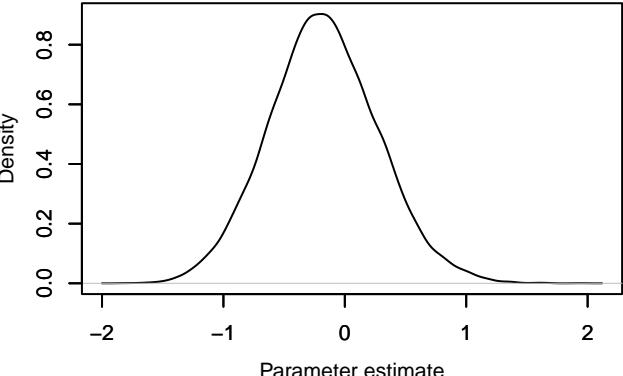
Density – $B[\text{DNAng.ul (C8)}, \text{Pearsonema (S7)}]$



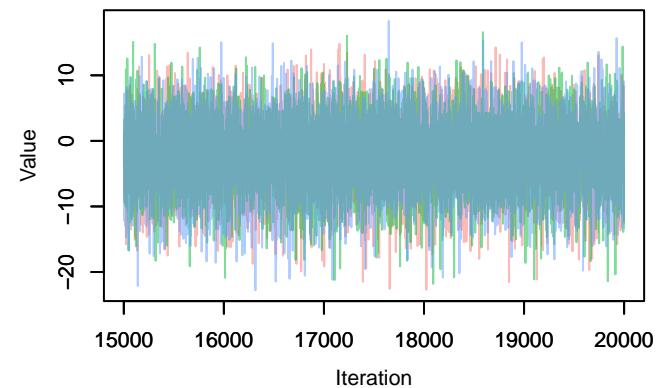
Trace – $B[\text{DNA260.230 (C9)}, \text{Pearsonema (S7)}]$



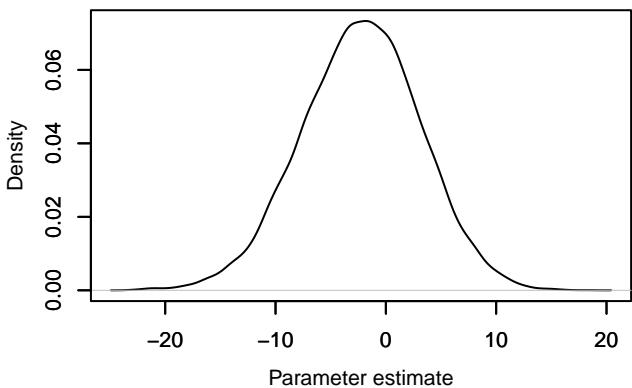
Density – $B[\text{DNA260.230 (C9)}, \text{Pearsonema (S7)}]$



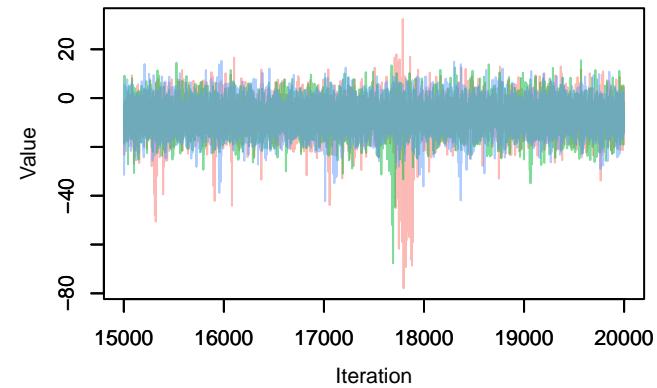
Trace – $B[\text{DNA260.280 (C10), Pearsonema (S7)}]$



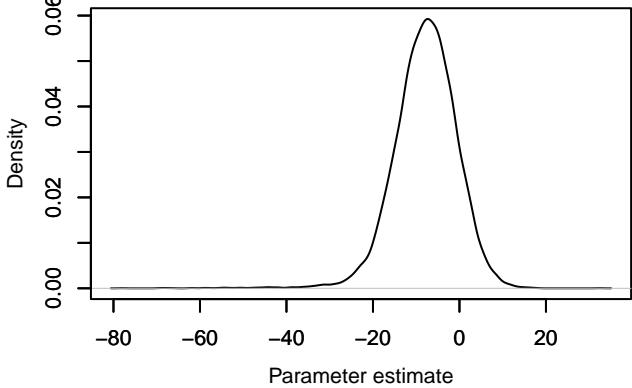
Density – $B[\text{DNA260.280 (C10), Pearsonema (S7)}]$



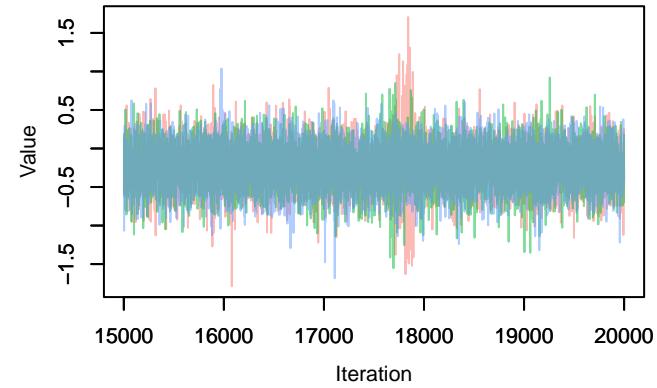
Trace – $B[(\text{Intercept}) (\text{C1}), \text{Mesocestoides (S8)}]$



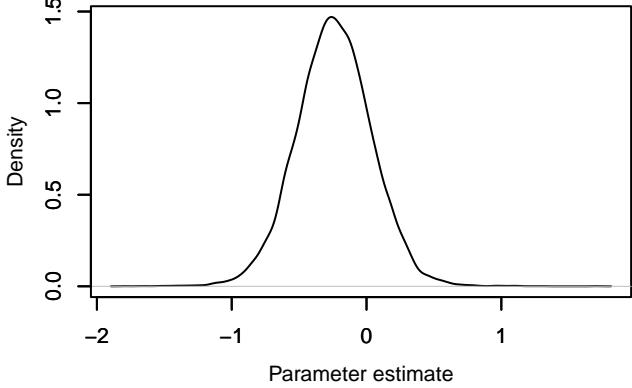
Density – $B[(\text{Intercept}) (\text{C1}), \text{Mesocestoides (S8)}]$



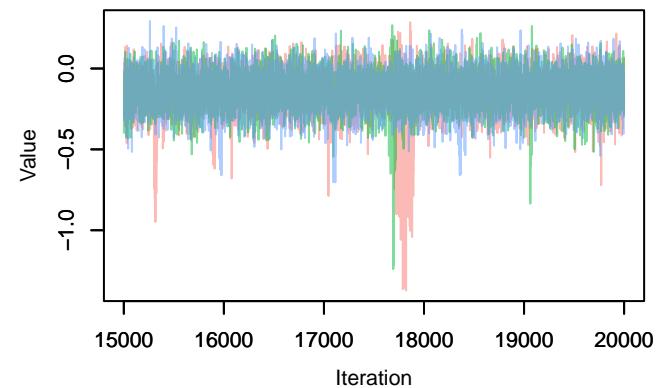
Trace – $B[\text{sexmale (C2), Mesocestoides (S8)}]$



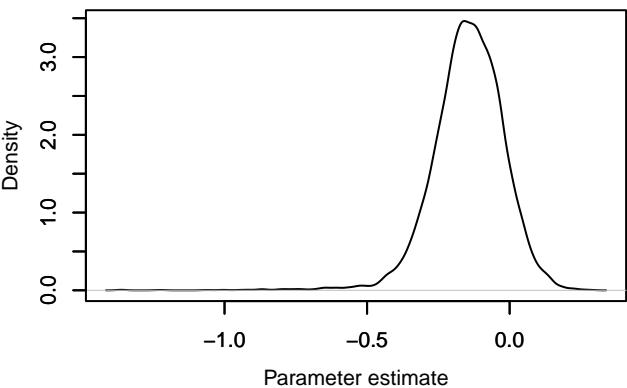
Density – $B[\text{sexmale (C2), Mesocestoides (S8)}]$



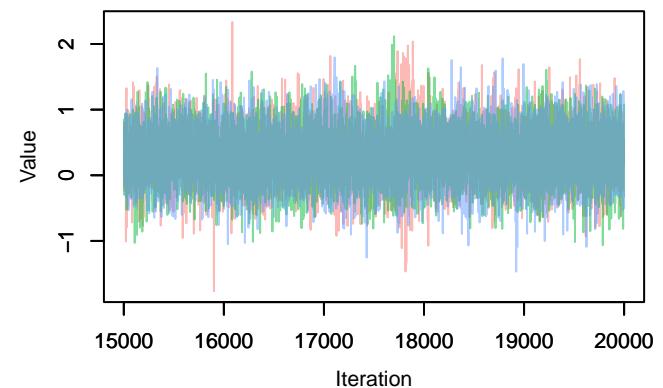
Trace – B[weight_kg (C3), Mesocestoides (S8)]



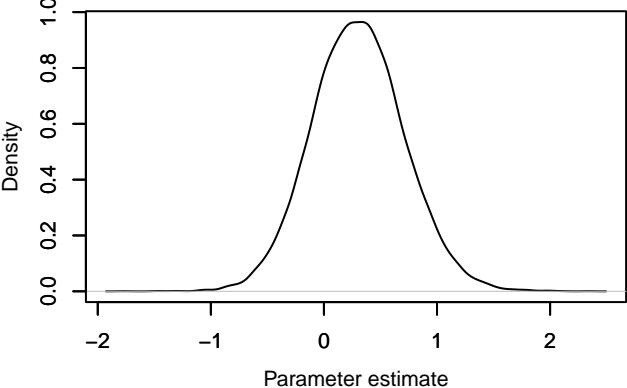
Density – B[weight_kg (C3), Mesocestoides (S8)]



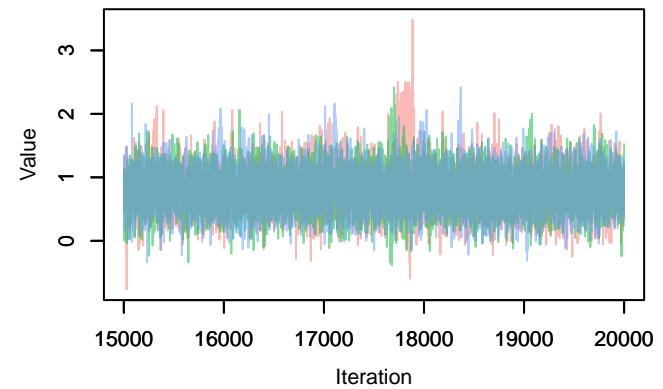
Trace – B[seasonspring (C4), Mesocestoides (S8)]



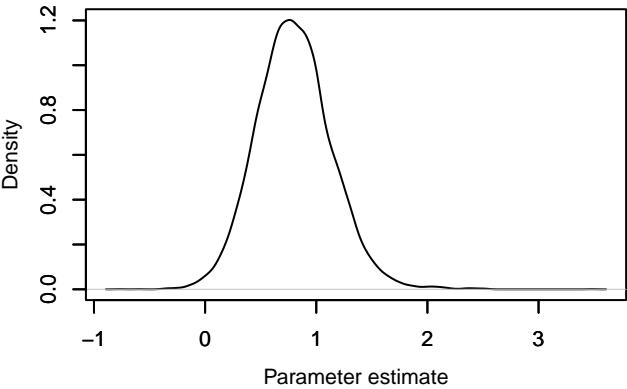
Density – B[seasonspring (C4), Mesocestoides (S8)]

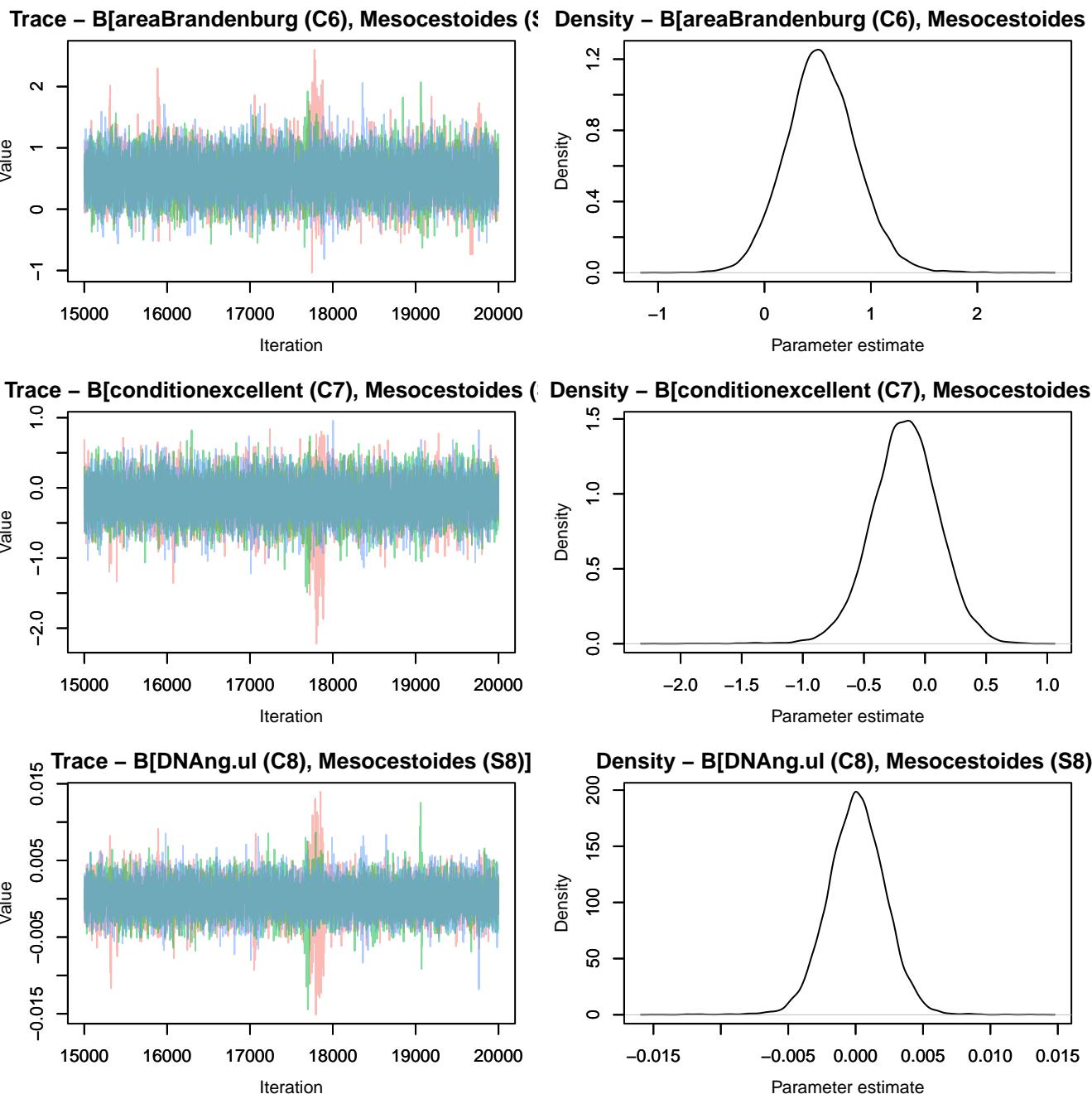


Trace – B[seasonwinter (C5), Mesocestoides (S8)]

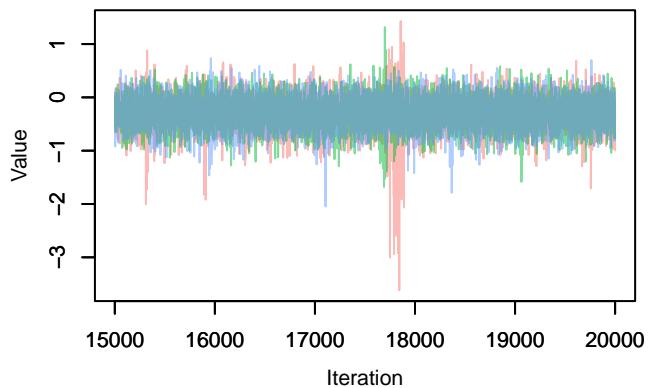


Density – B[seasonwinter (C5), Mesocestoides (S8)]

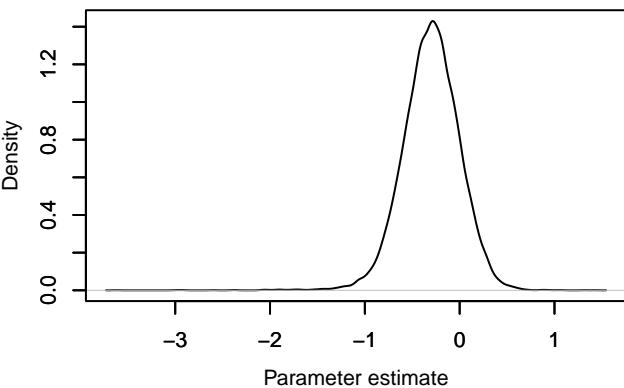




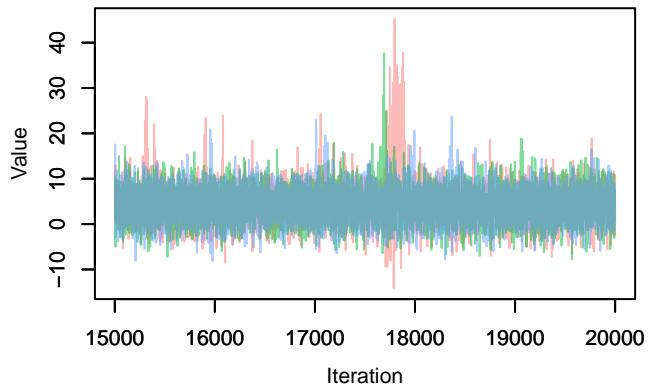
Trace – $B[\text{DNA260.230 (C9), Mesocestoides (S8)}]$



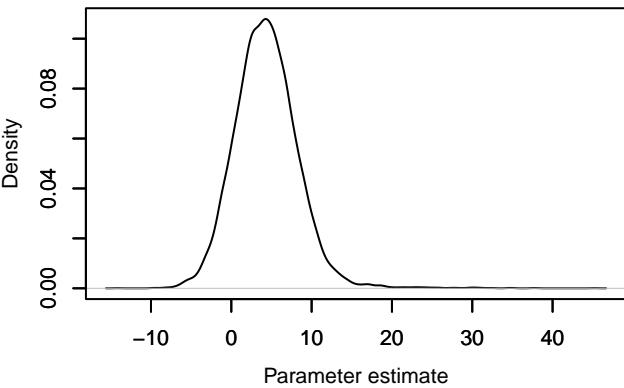
Density – $B[\text{DNA260.230 (C9), Mesocestoides (S8)}]$



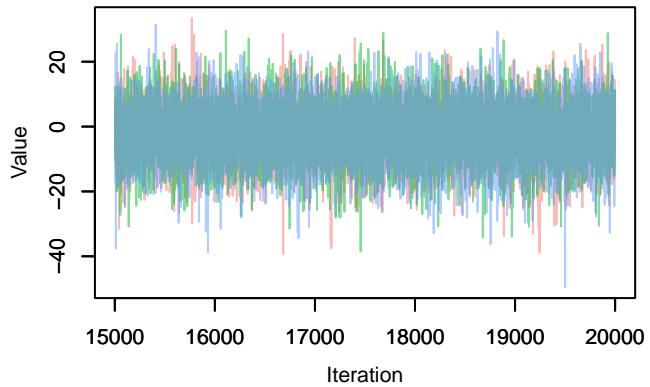
Trace – $B[\text{DNA260.280 (C10), Mesocestoides (S8)}]$



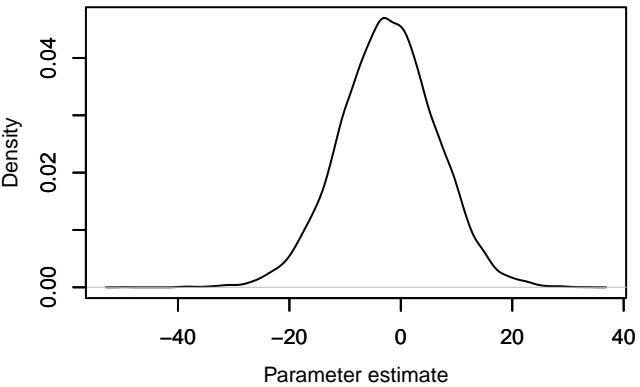
Density – $B[\text{DNA260.280 (C10), Mesocestoides (S8)}]$



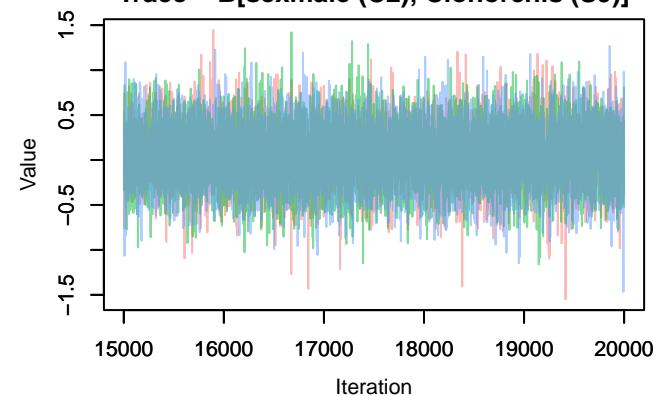
Trace – $B[(\text{Intercept}) (\text{C1}), \text{Clonorchis (S9)}]$



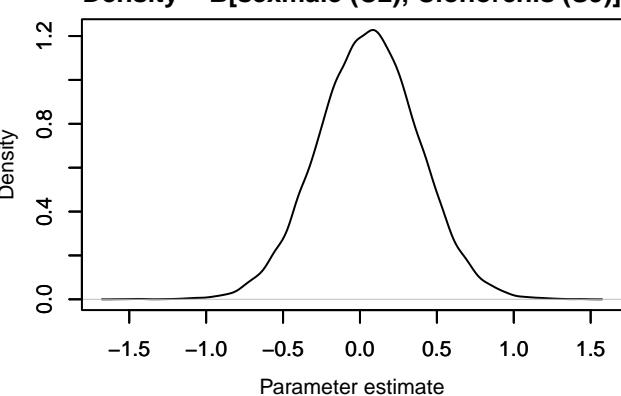
Density – $B[(\text{Intercept}) (\text{C1}), \text{Clonorchis (S9)}]$



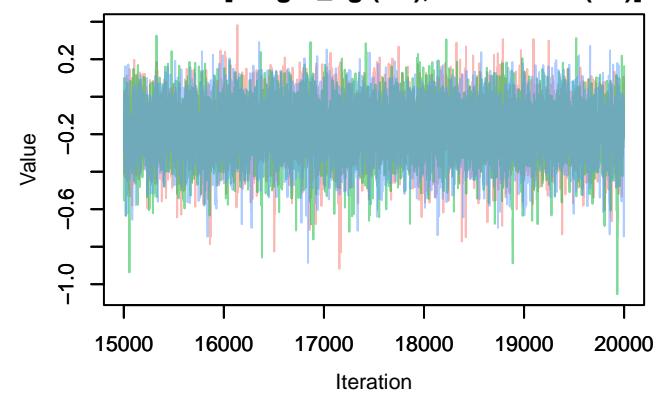
Trace – B[sexmale (C2), Clonorchis (S9)]



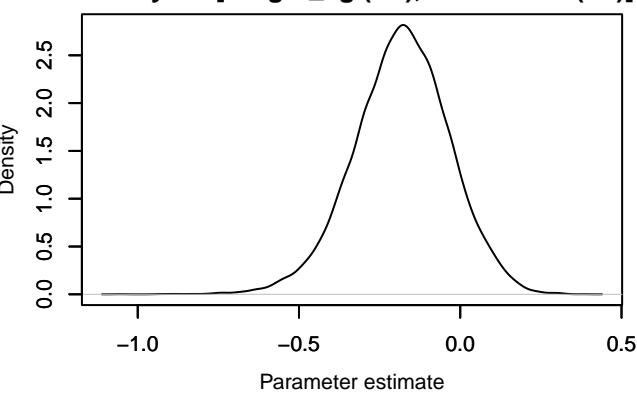
Density – B[sexmale (C2), Clonorchis (S9)]



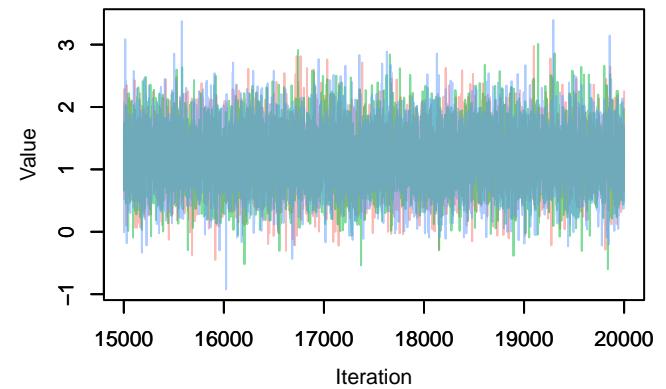
Trace – B[weight_kg (C3), Clonorchis (S9)]



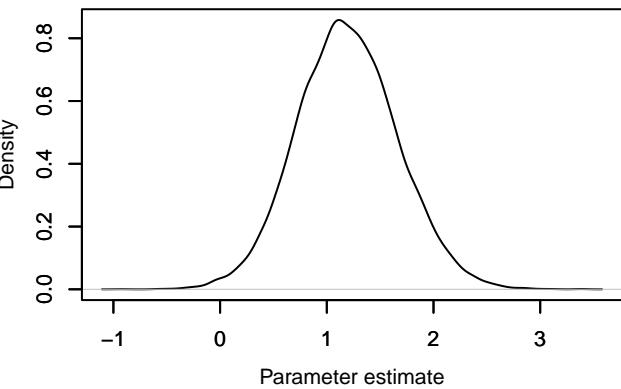
Density – B[weight_kg (C3), Clonorchis (S9)]



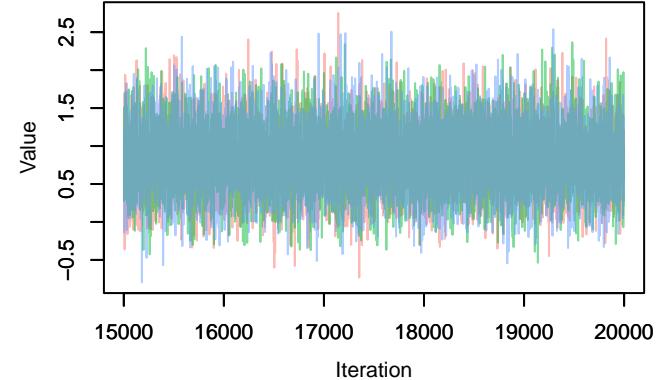
Trace – B[seasonspring (C4), Clonorchis (S9)]



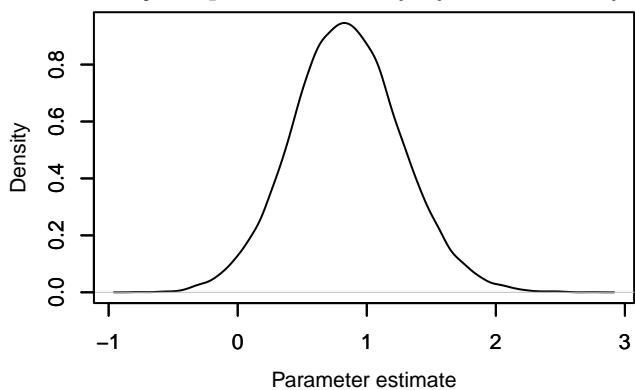
Density – B[seasonspring (C4), Clonorchis (S9)]



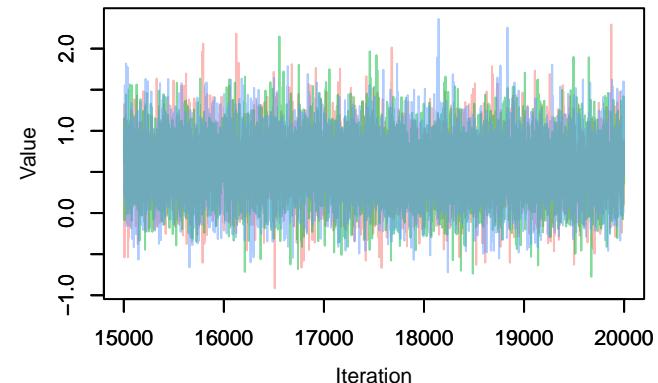
Trace – B[seasonwinter (C5), Clonorchis (S9)]



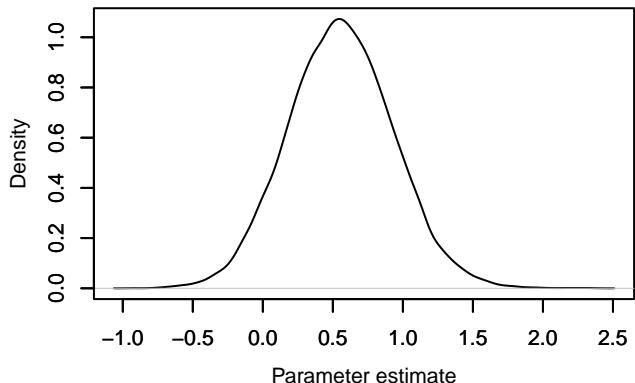
Density – B[seasonwinter (C5), Clonorchis (S9)]



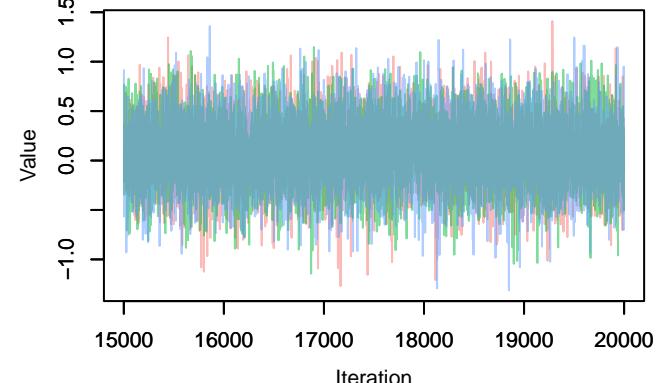
Trace – B[areaBrandenburg (C6), Clonorchis (S9)]



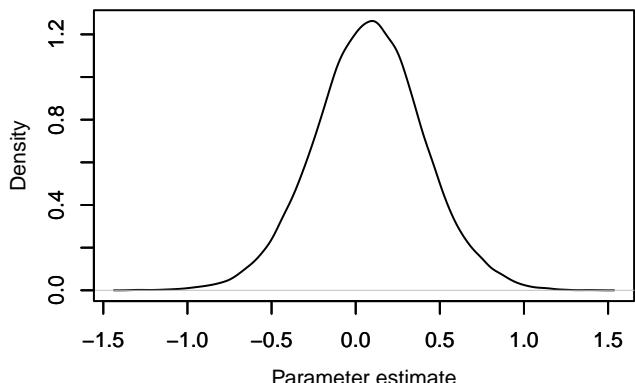
Density – B[areaBrandenburg (C6), Clonorchis (S9)]



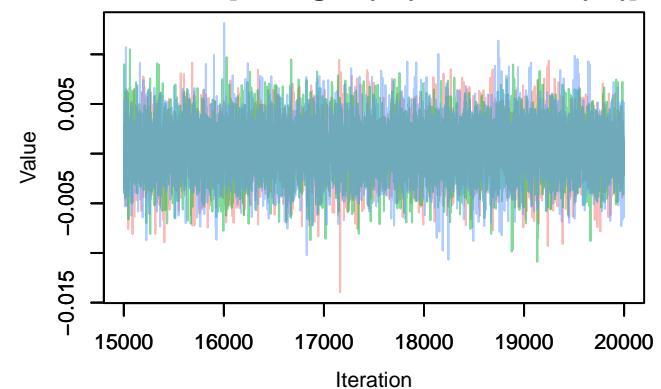
Trace – B[conditionexcellent (C7), Clonorchis (S9)]



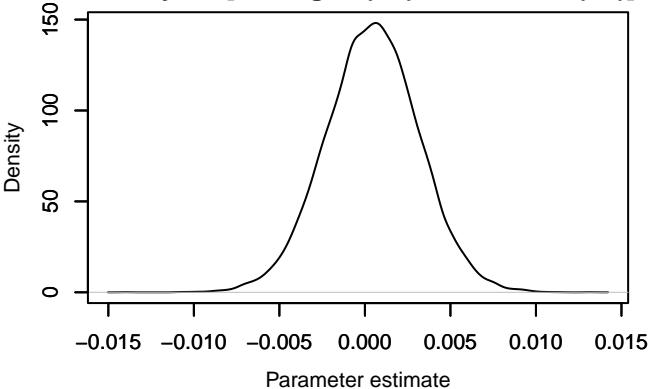
Density – B[conditionexcellent (C7), Clonorchis (S9)]



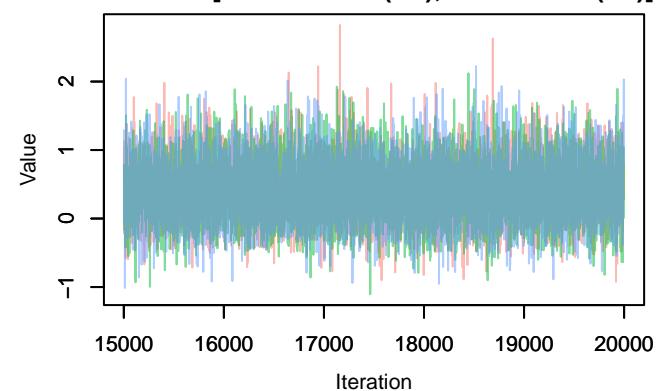
Trace – B[DNAng.ul (C8), Clonorchis (S9)]



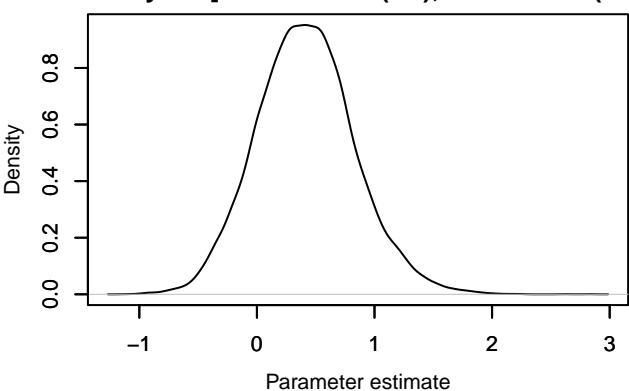
Density – B[DNAng.ul (C8), Clonorchis (S9)]



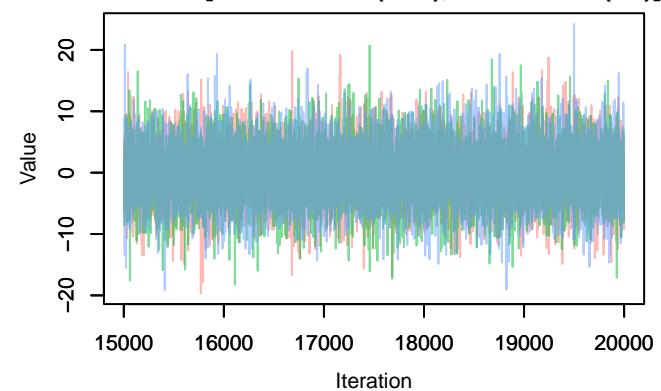
Trace – B[DNA260.230 (C9), Clonorchis (S9)]



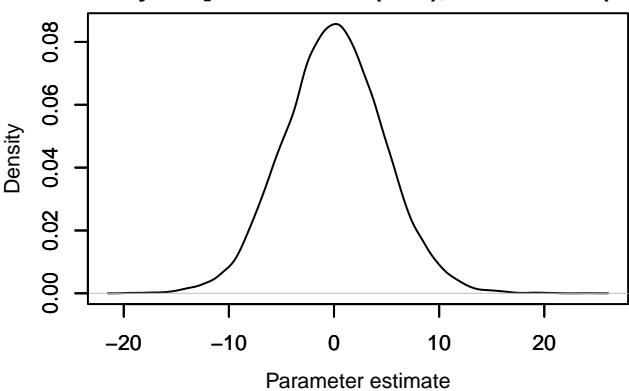
Density – B[DNA260.230 (C9), Clonorchis (S9)]



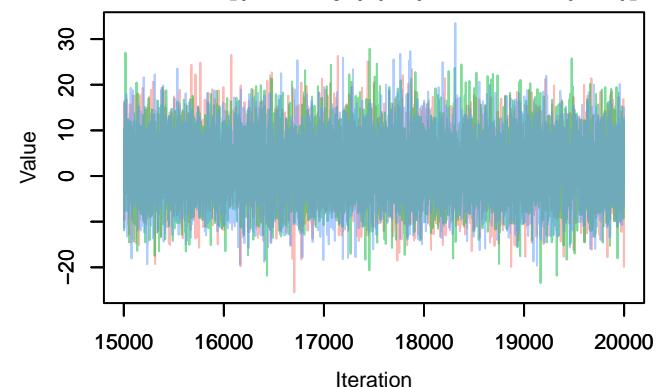
Trace – B[DNA260.280 (C10), Clonorchis (S9)]



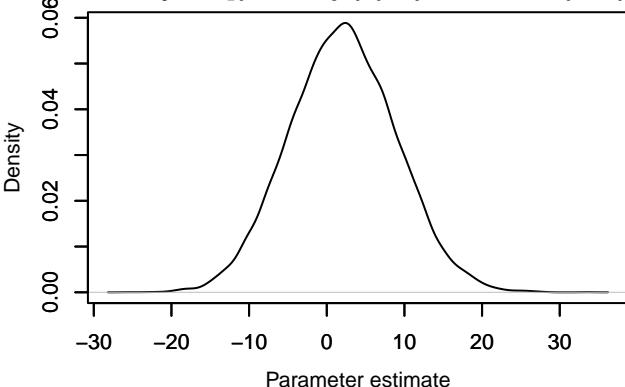
Density – B[DNA260.280 (C10), Clonorchis (S9)]



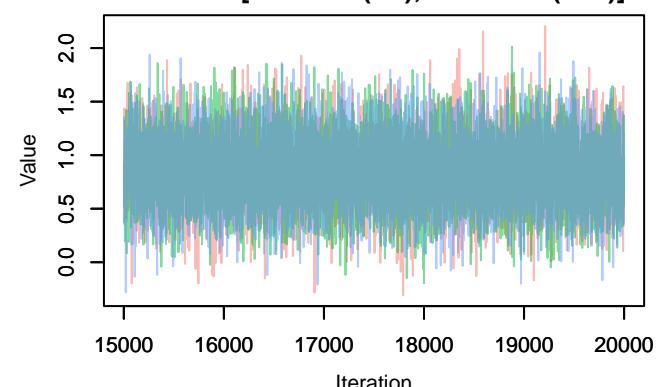
Trace – $B[(\text{Intercept}) (\text{C1}), \text{Toxocara} (\text{S10})]$



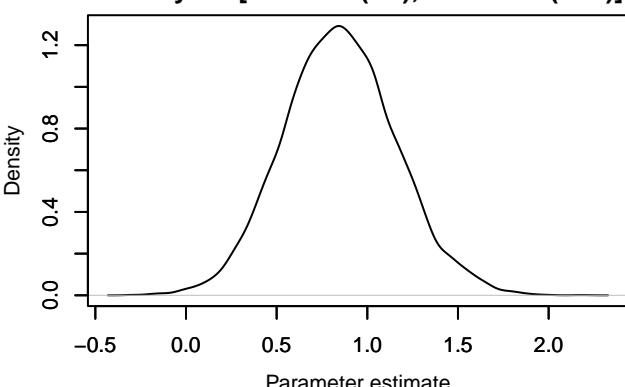
Density – $B[(\text{Intercept}) (\text{C1}), \text{Toxocara} (\text{S10})]$



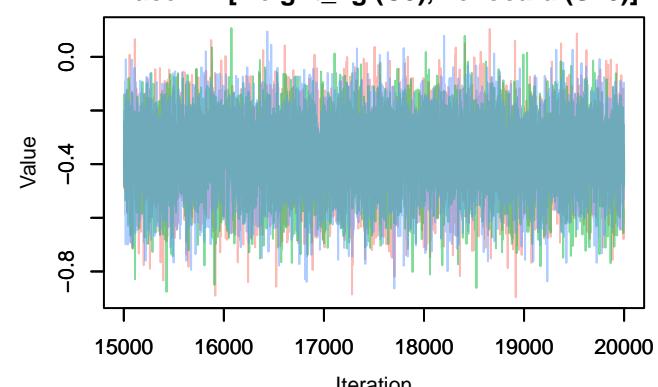
Trace – $B[\text{sexmale} (\text{C2}), \text{Toxocara} (\text{S10})]$



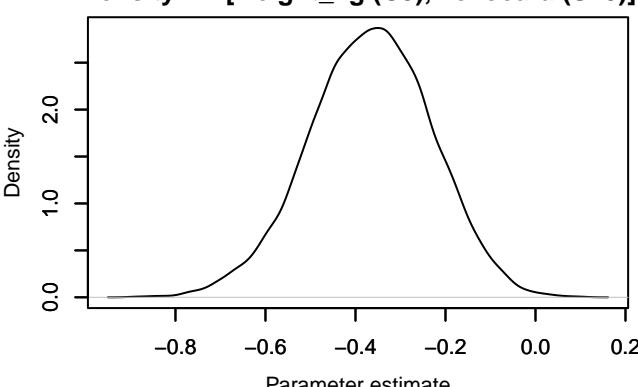
Density – $B[\text{sexmale} (\text{C2}), \text{Toxocara} (\text{S10})]$

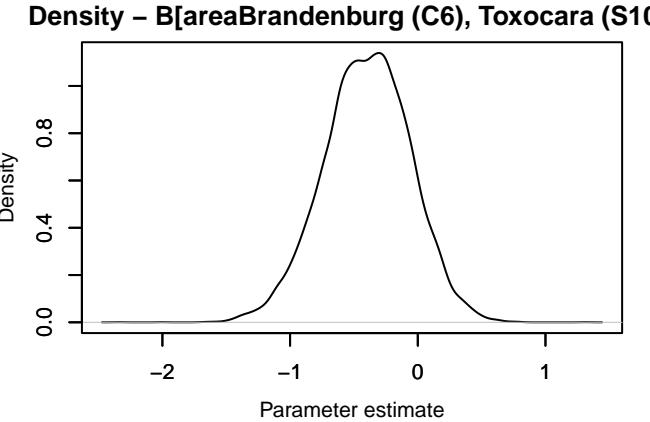
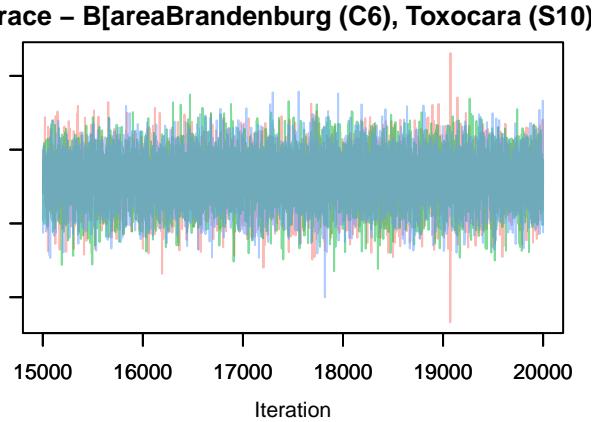
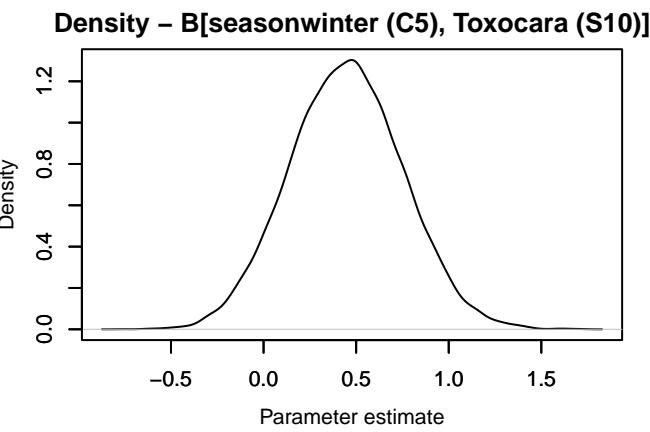
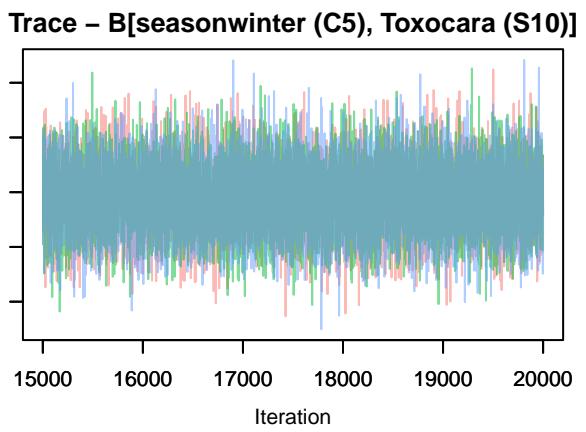
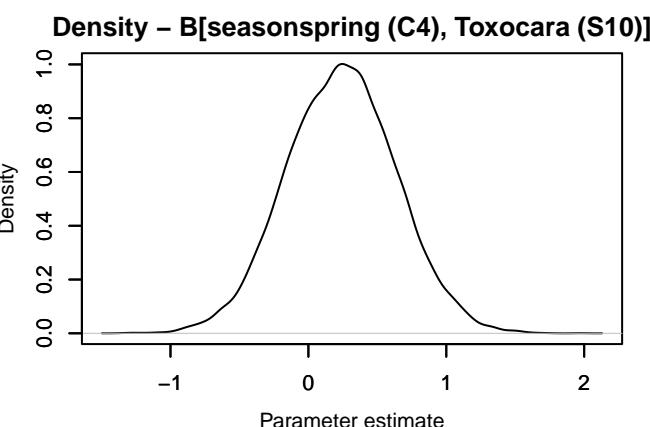
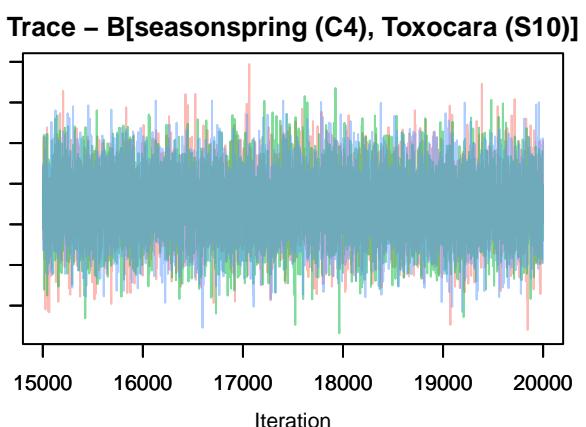


Trace – $B[\text{weight_kg} (\text{C3}), \text{Toxocara} (\text{S10})]$

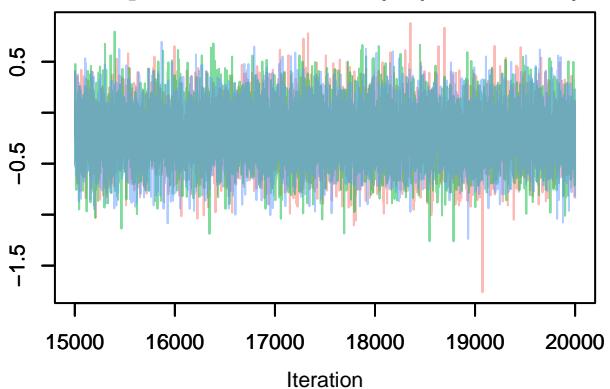


Density – $B[\text{weight_kg} (\text{C3}), \text{Toxocara} (\text{S10})]$

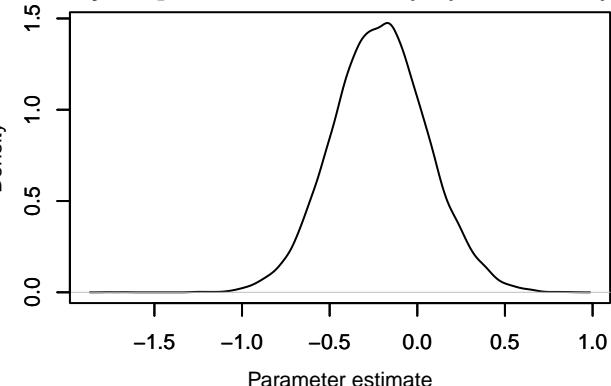




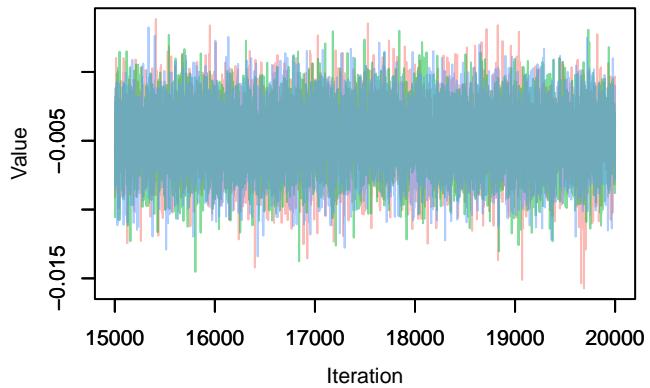
Trace – $B[\text{conditionexcellent (C7), Toxocara (S10)}]$



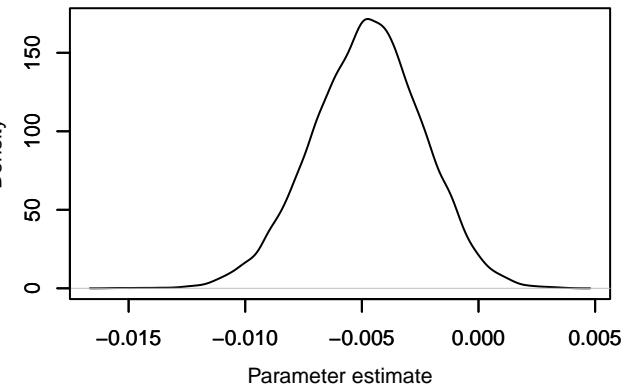
Density – $B[\text{conditionexcellent (C7), Toxocara (S10)}]$



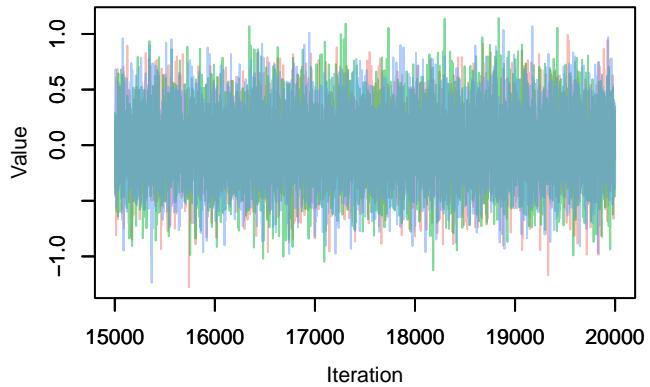
Trace – $B[\text{DNAng.ul (C8), Toxocara (S10)}]$



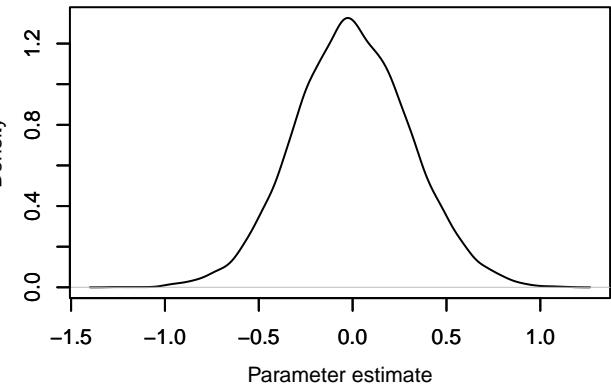
Density – $B[\text{DNAng.ul (C8), Toxocara (S10)}]$



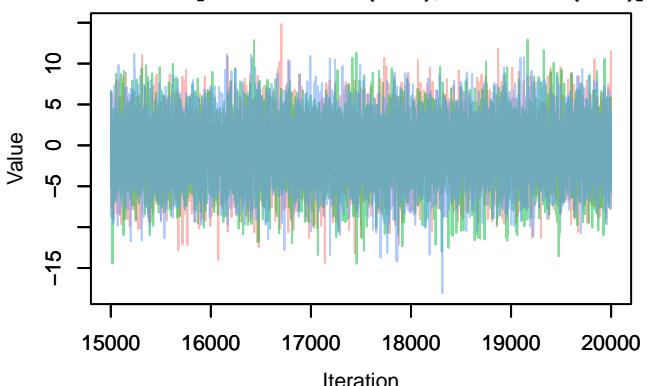
Trace – $B[\text{DNA260.230 (C9), Toxocara (S10)}]$



Density – $B[\text{DNA260.230 (C9), Toxocara (S10)}]$



Trace – B[DNA260.280 (C10), Toxocara (S10)]



Density – B[DNA260.280 (C10), Toxocara (S10)]

