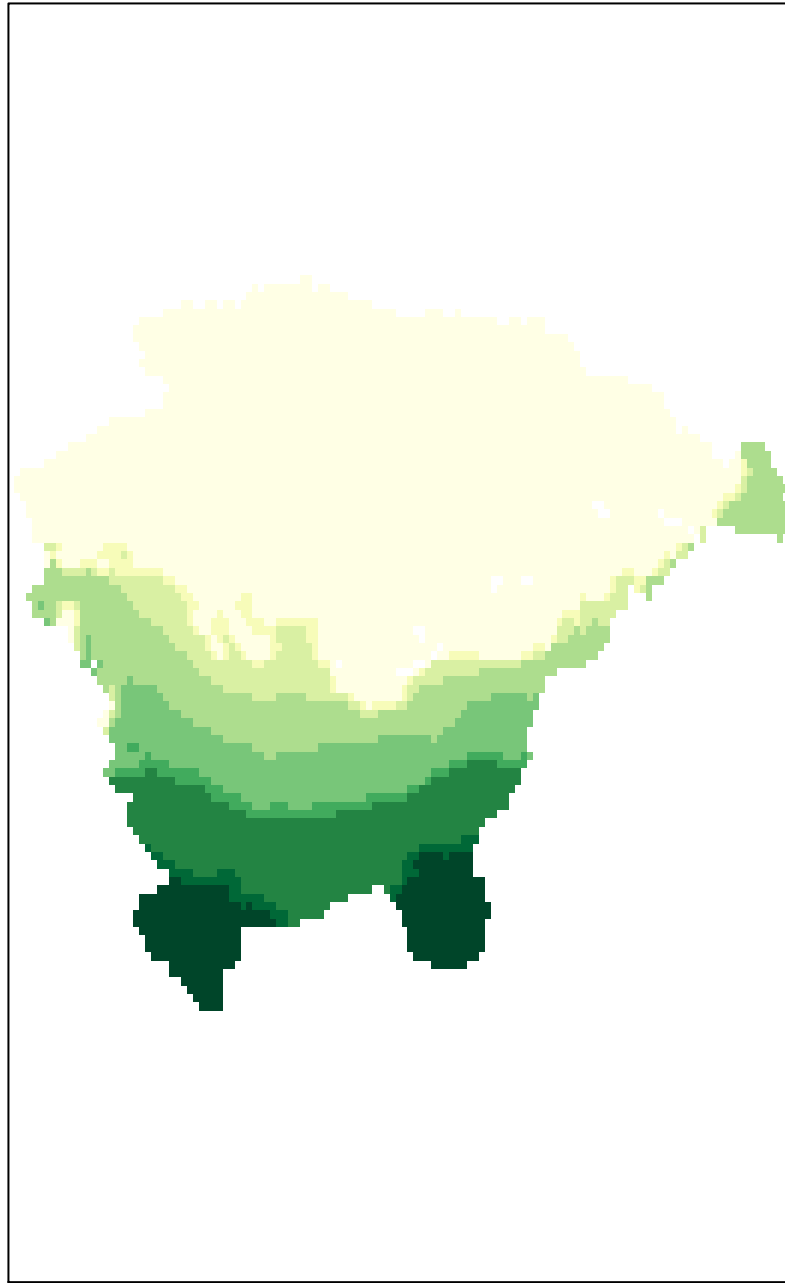
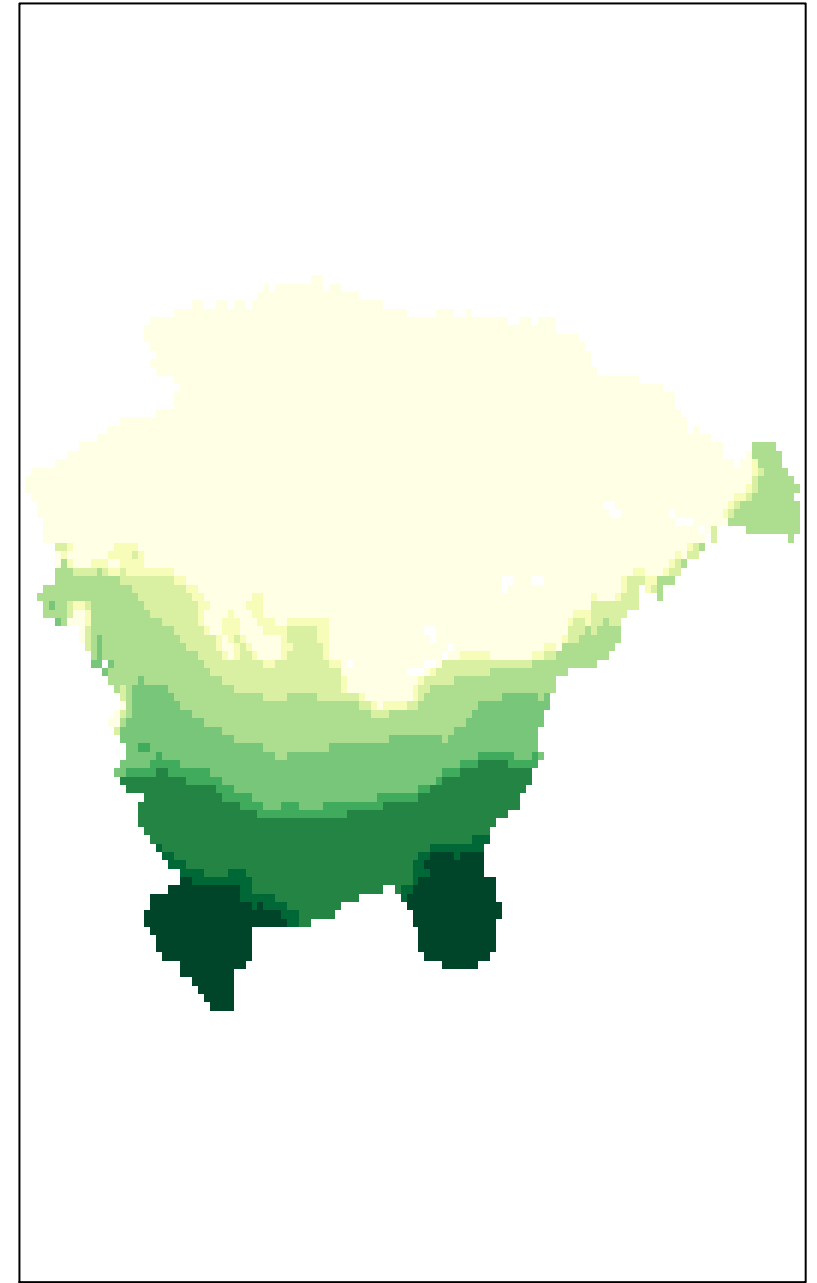


MAX, X21000.ybp

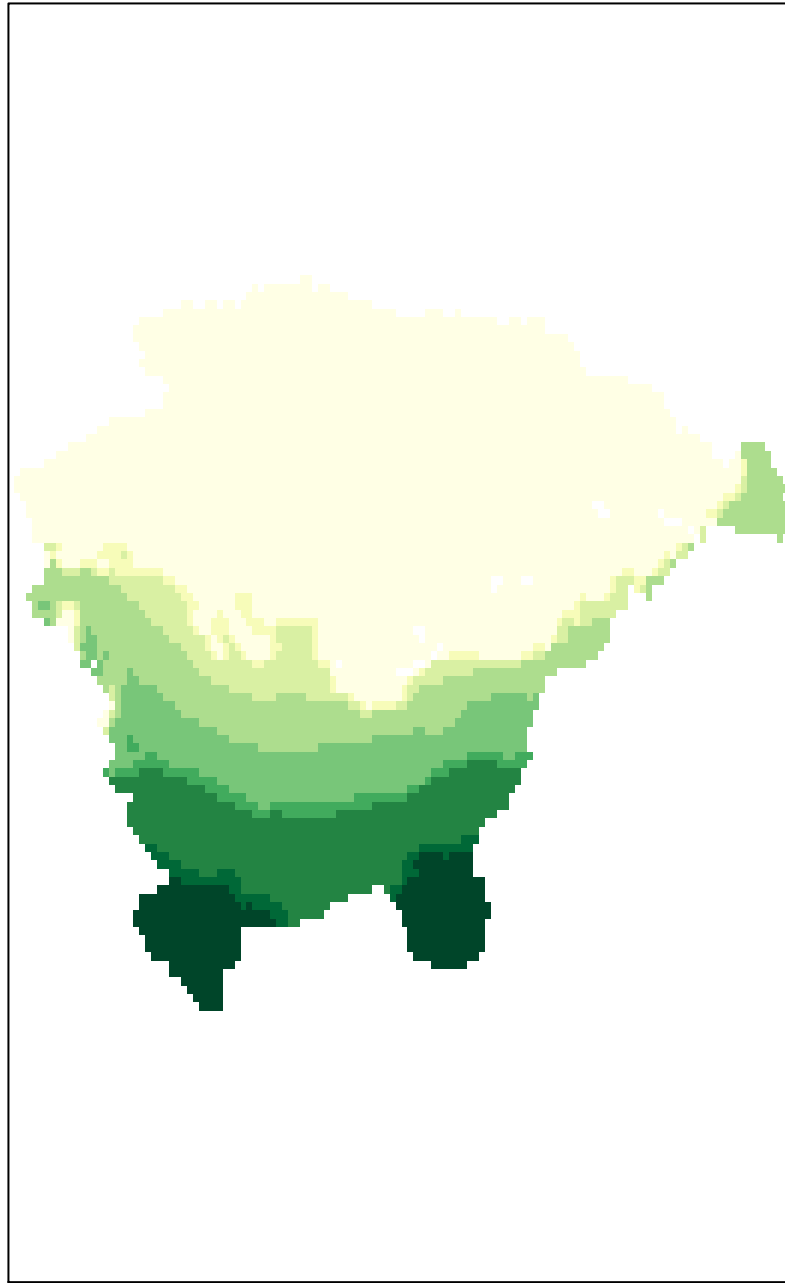


MAX, X21000.ybp

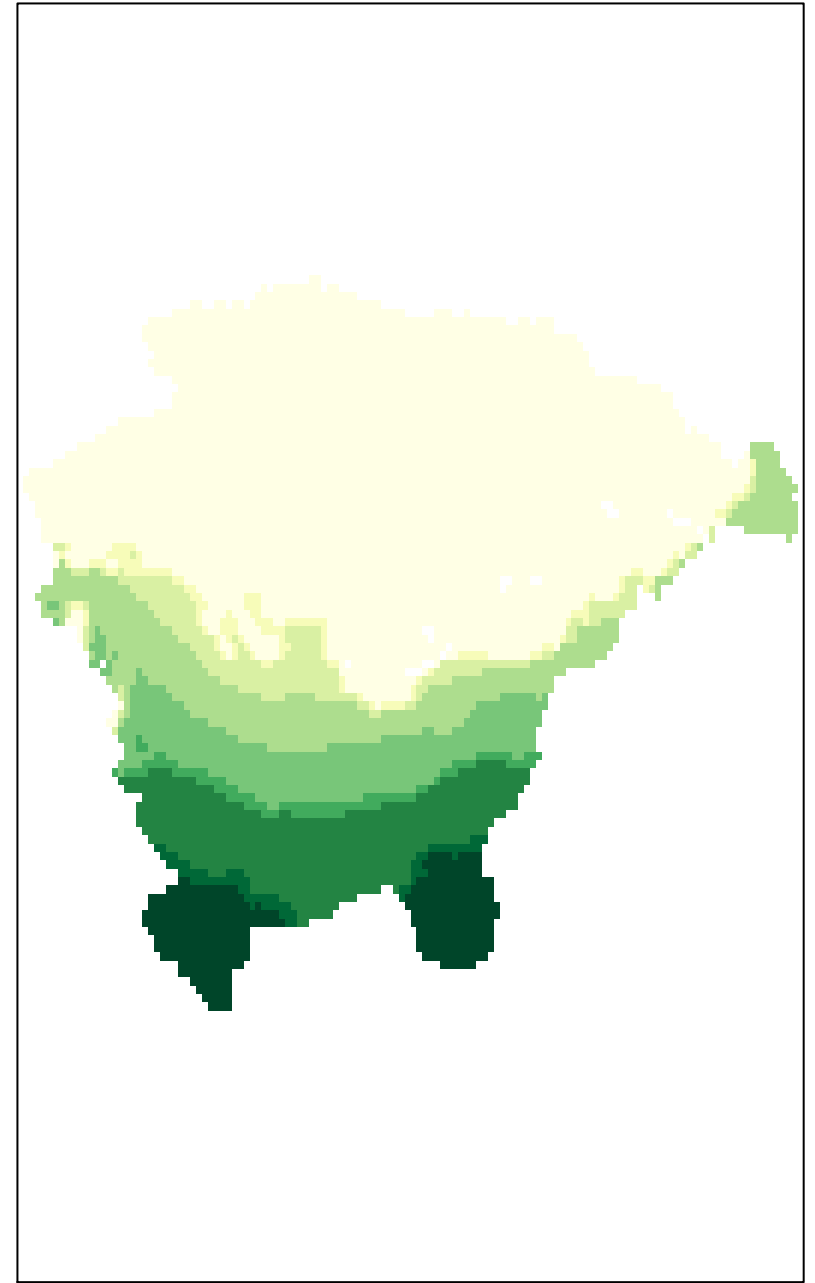


Species skipped = *Fraxinus caroliniana*, GCM = Lorenz_ccsm

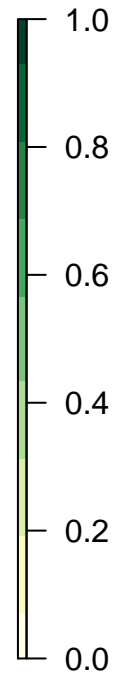
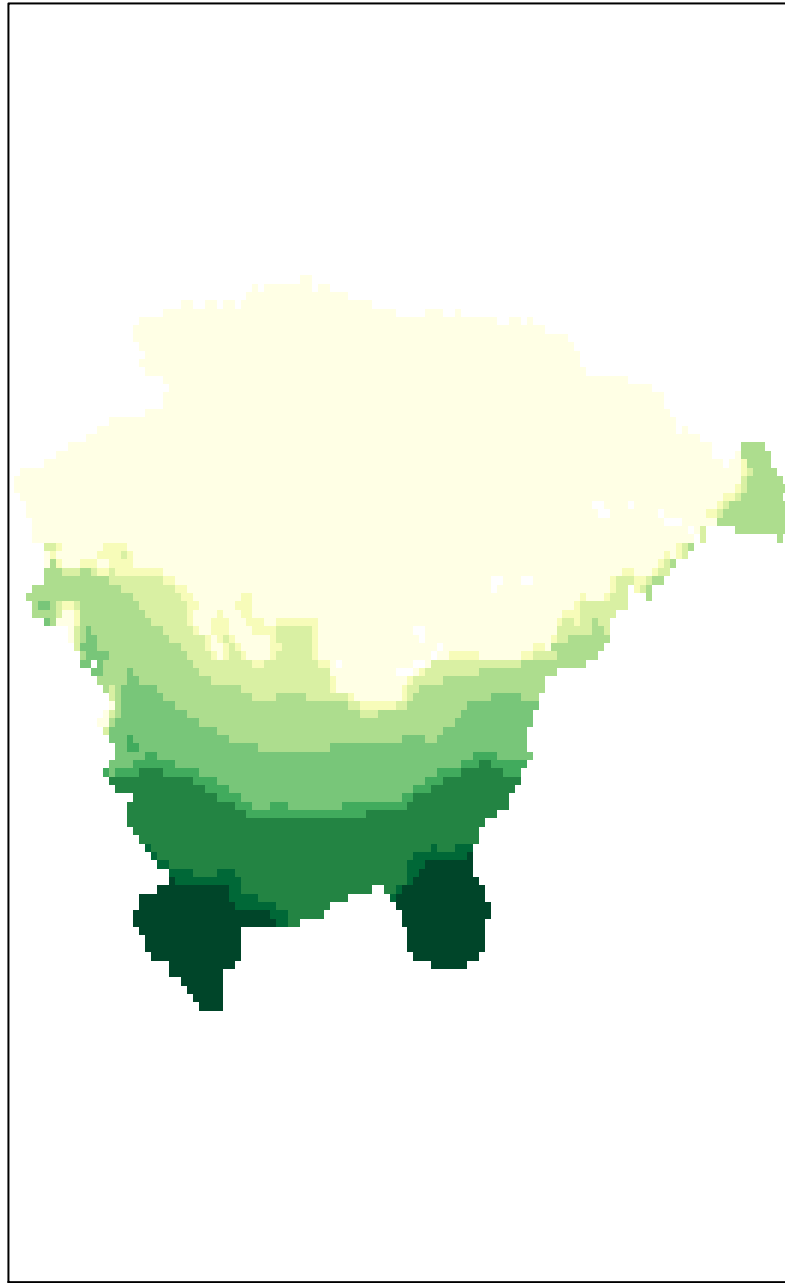
MAX, X20000.ybp



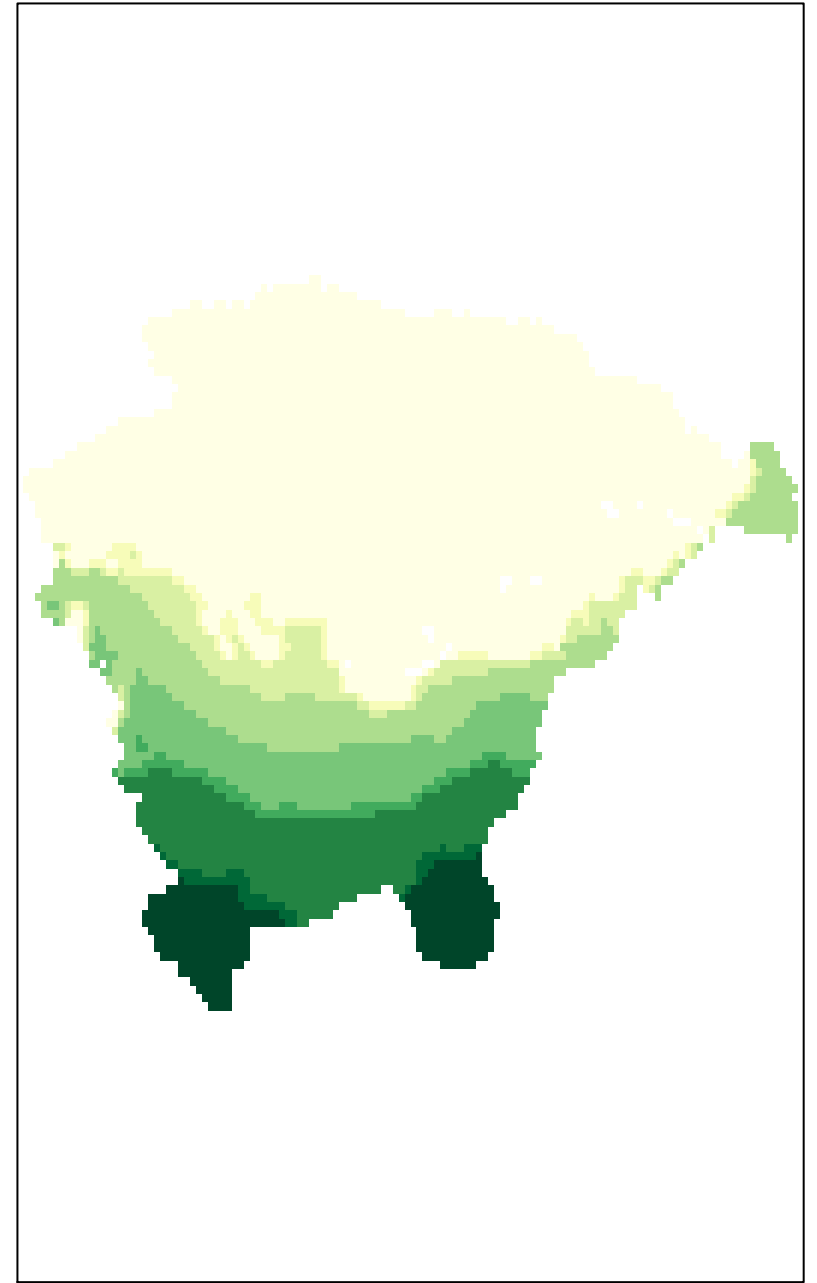
MAX, X20000.ybp



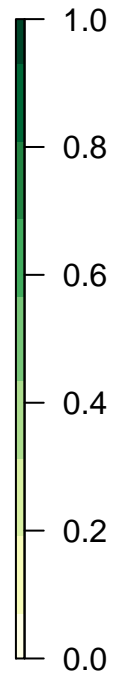
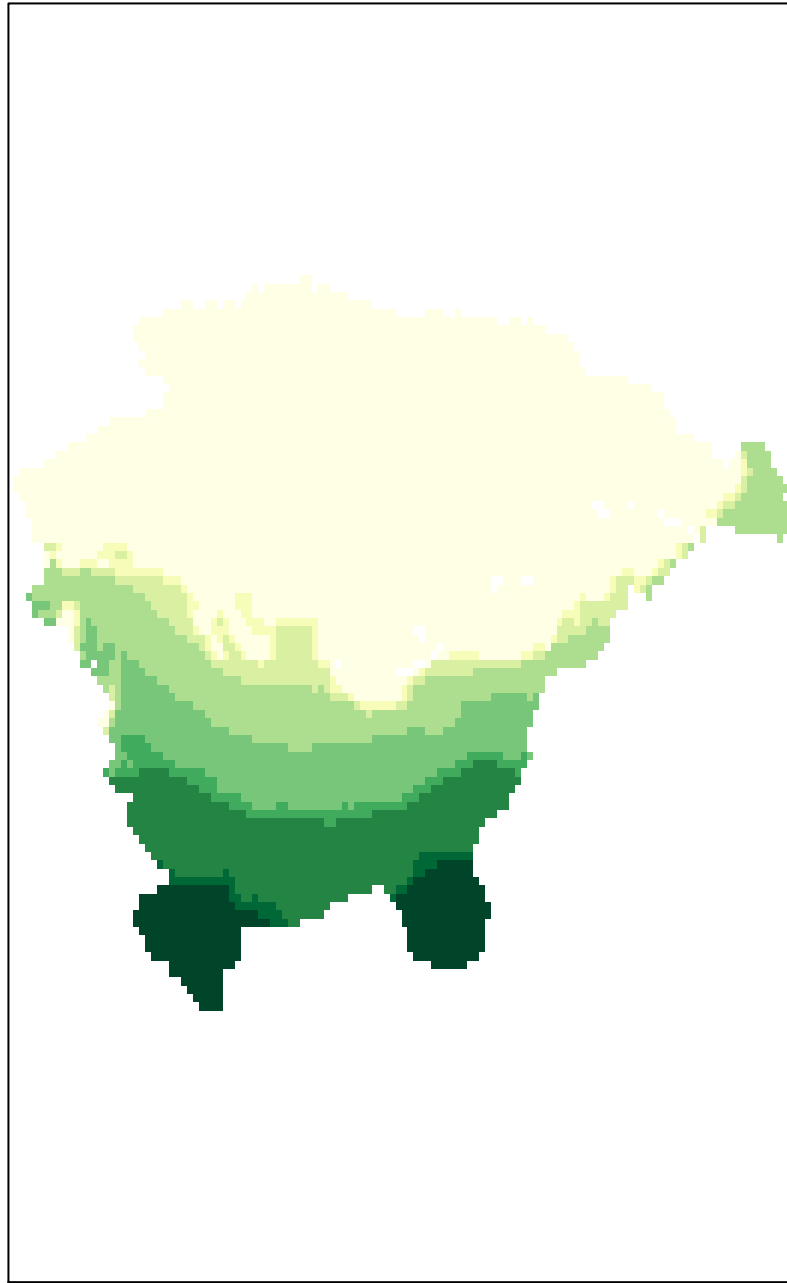
MAX, X19000.ybp



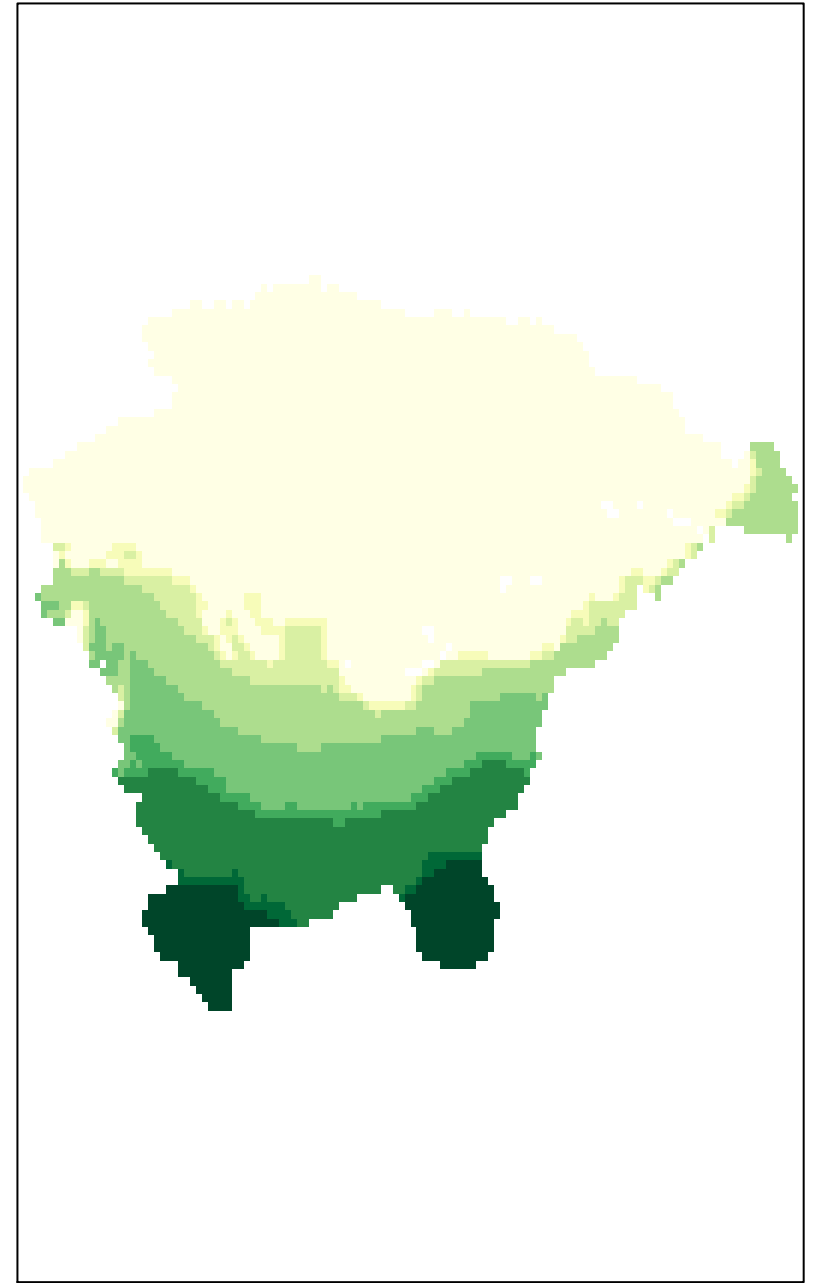
MAX, X19000.ybp



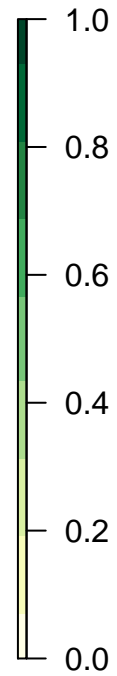
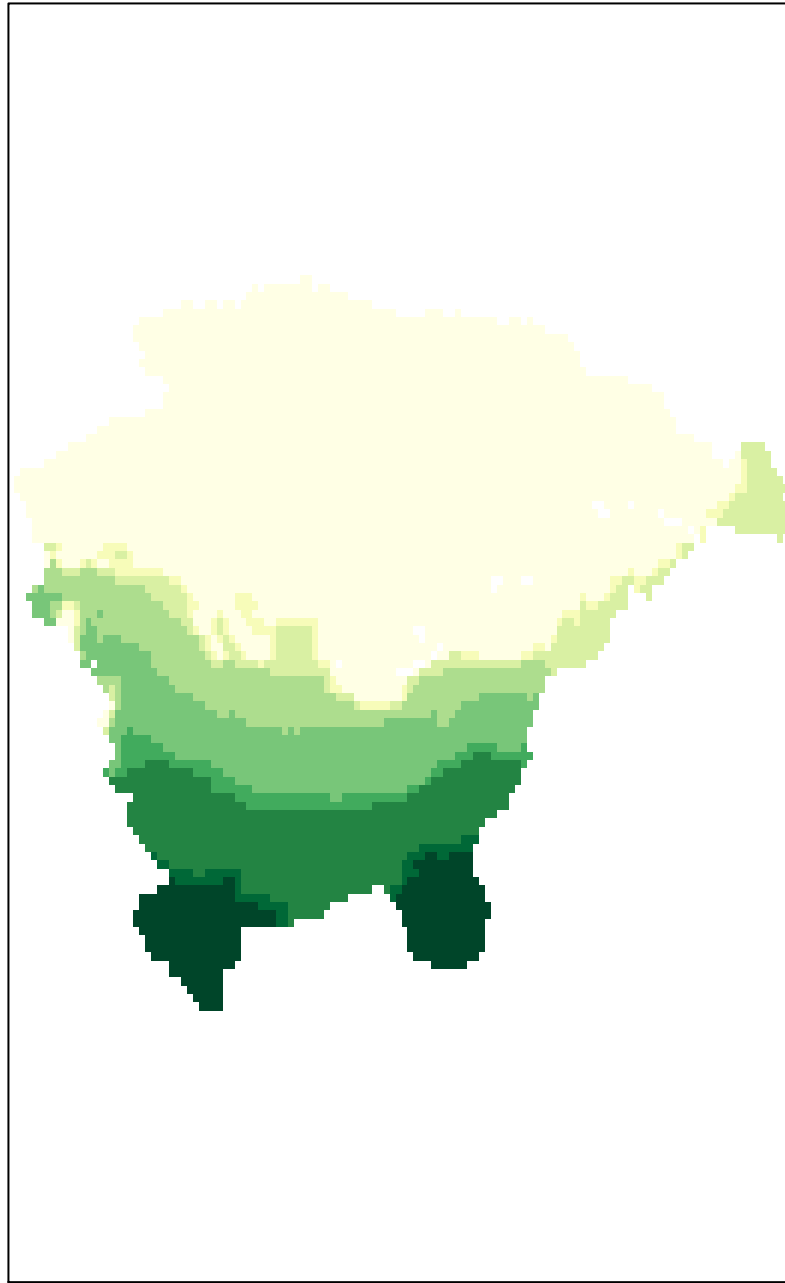
MAX, X18000.ybp



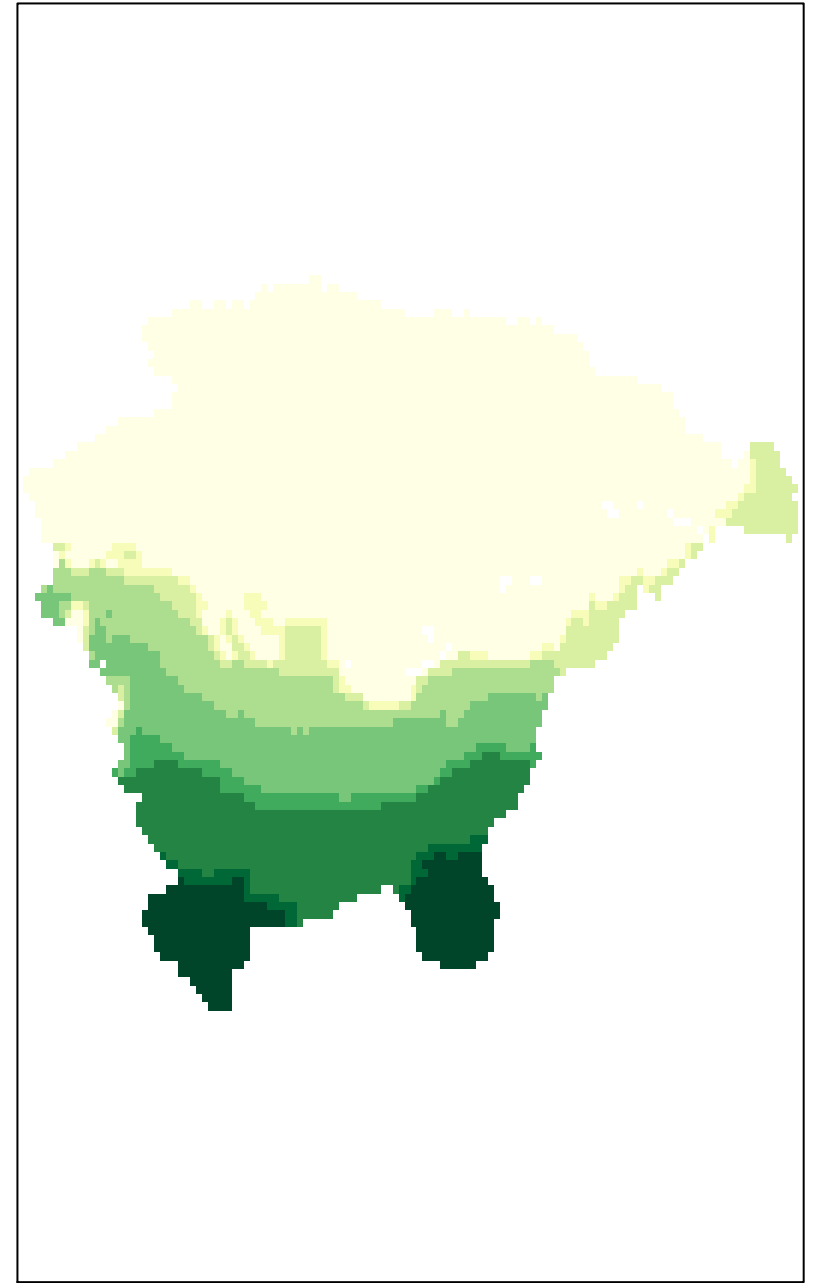
MAX, X18000.ybp



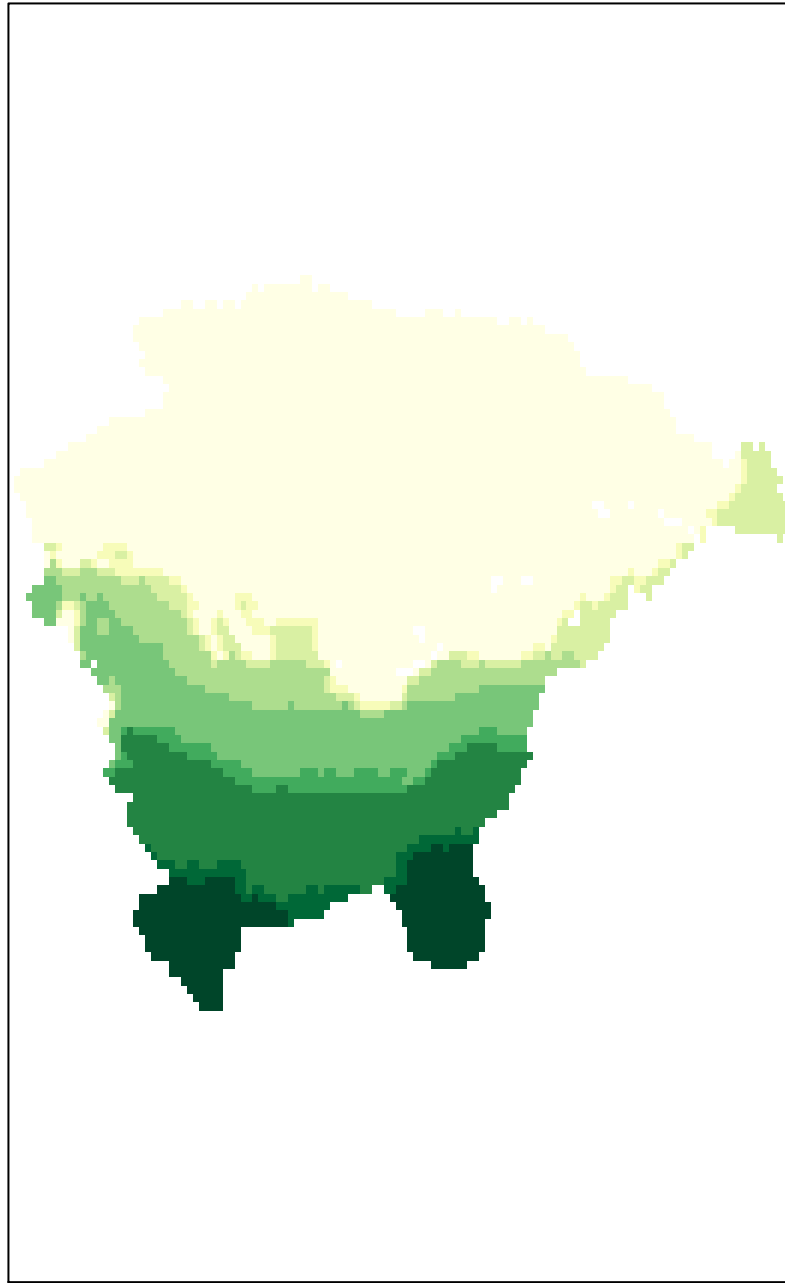
MAX, X17000.ybp



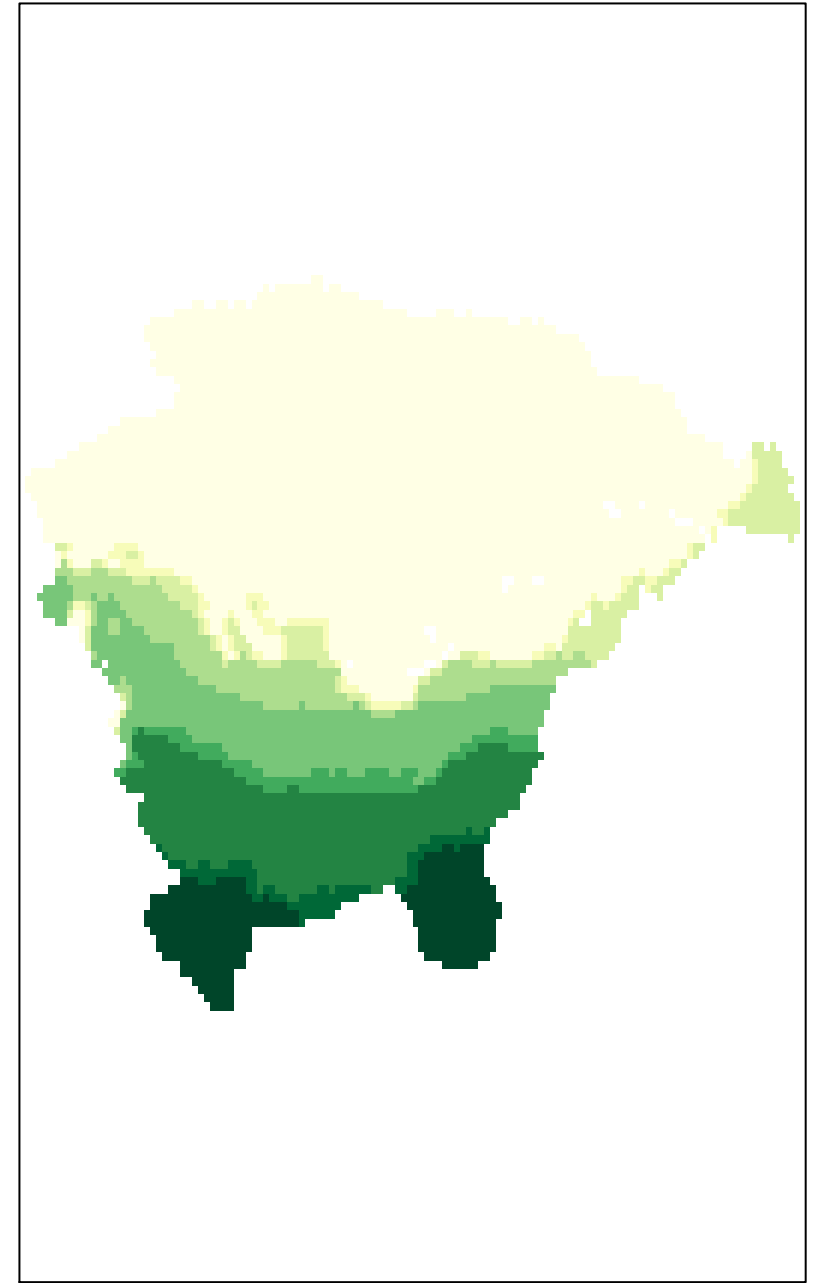
MAX, X17000.ybp



MAX, X16000.ybp

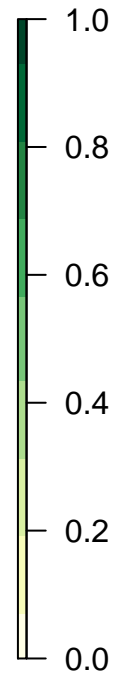
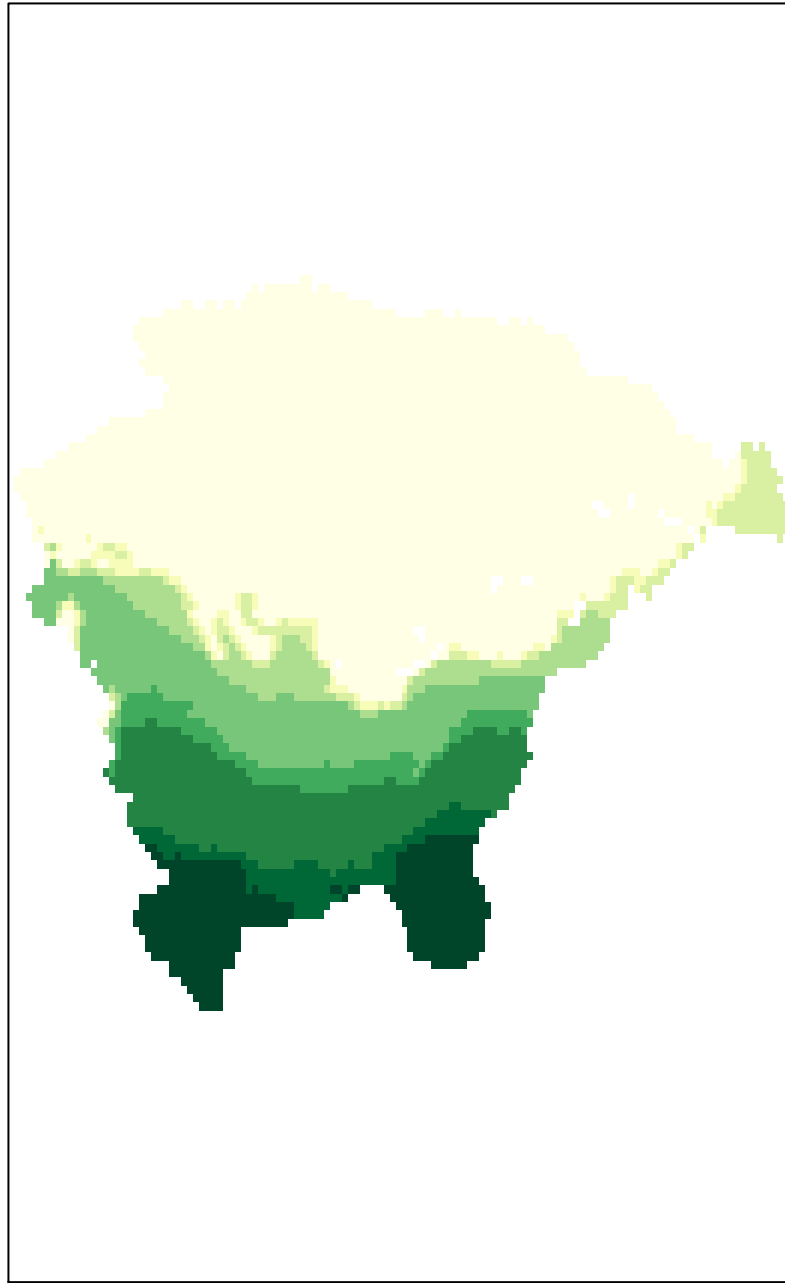


MAX, X16000.ybp

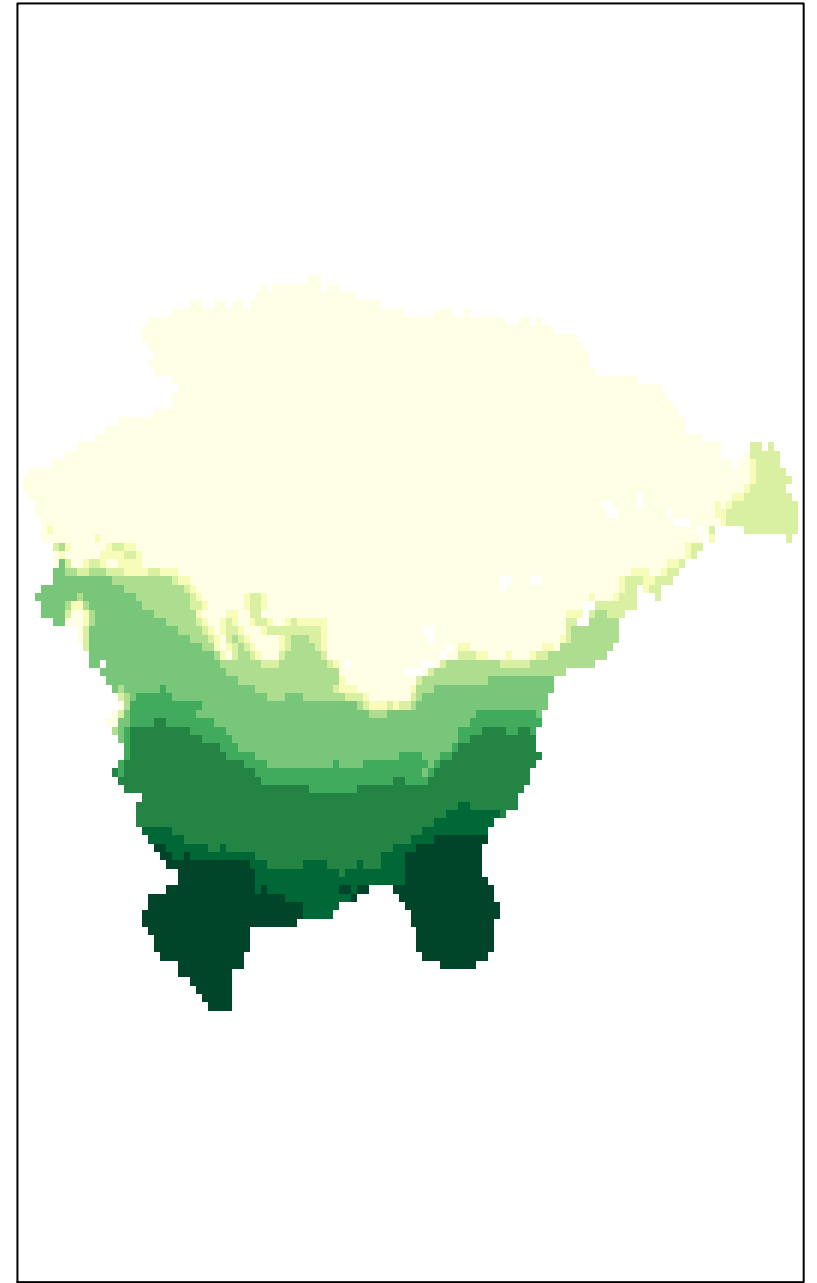


Species skipped = *Fraxinus caroliniana*, GCM = Lorenz_ccsm

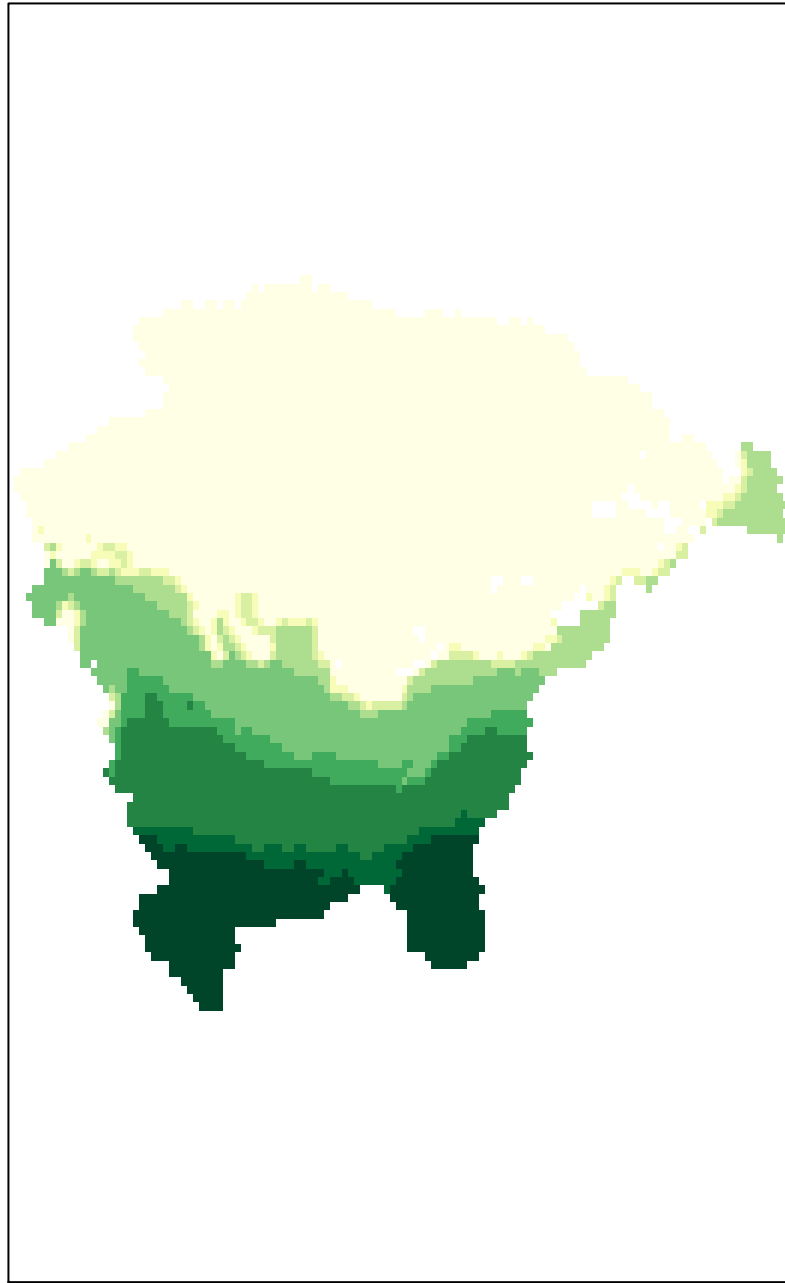
MAX, X15000.ybp



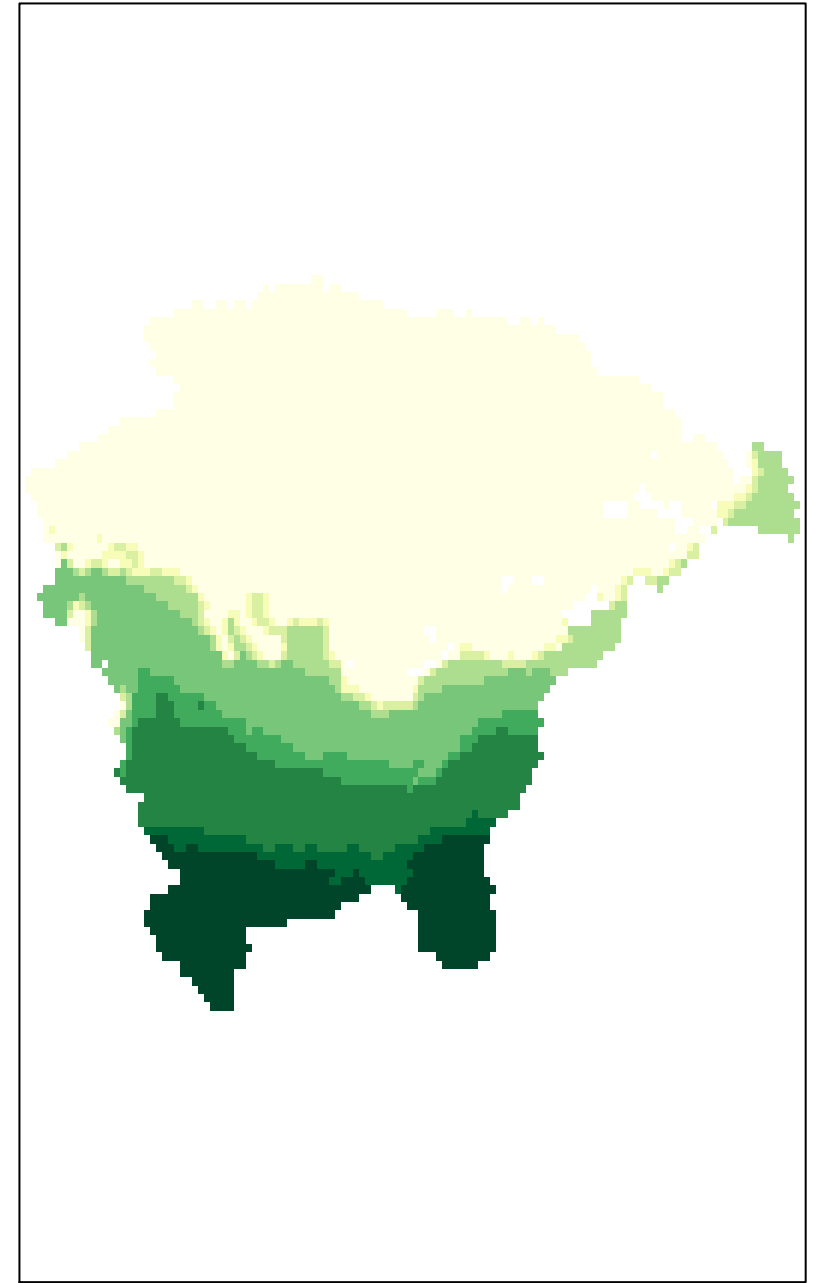
MAX, X15000.ybp



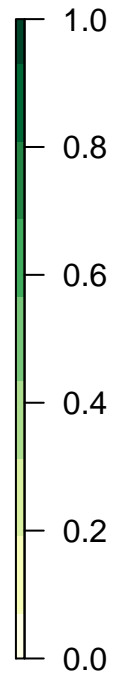
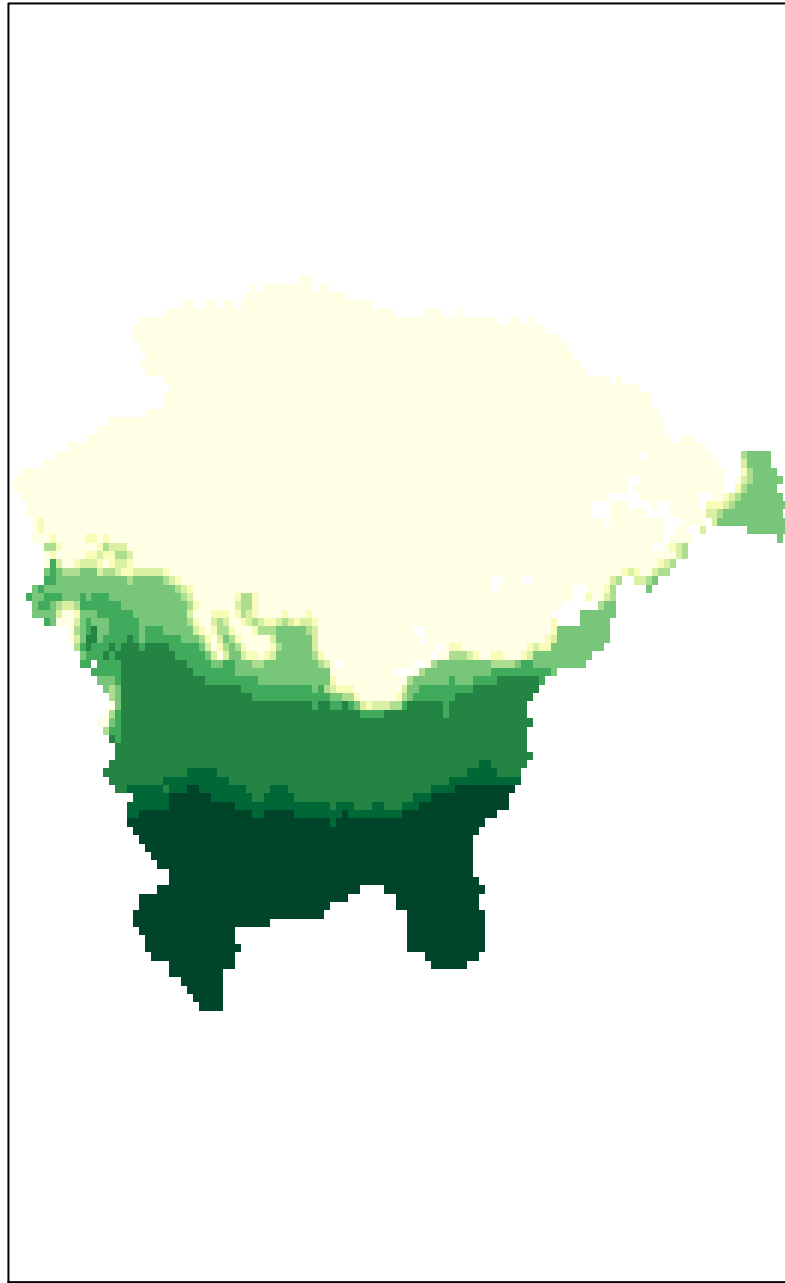
MAX, X14000.ybp



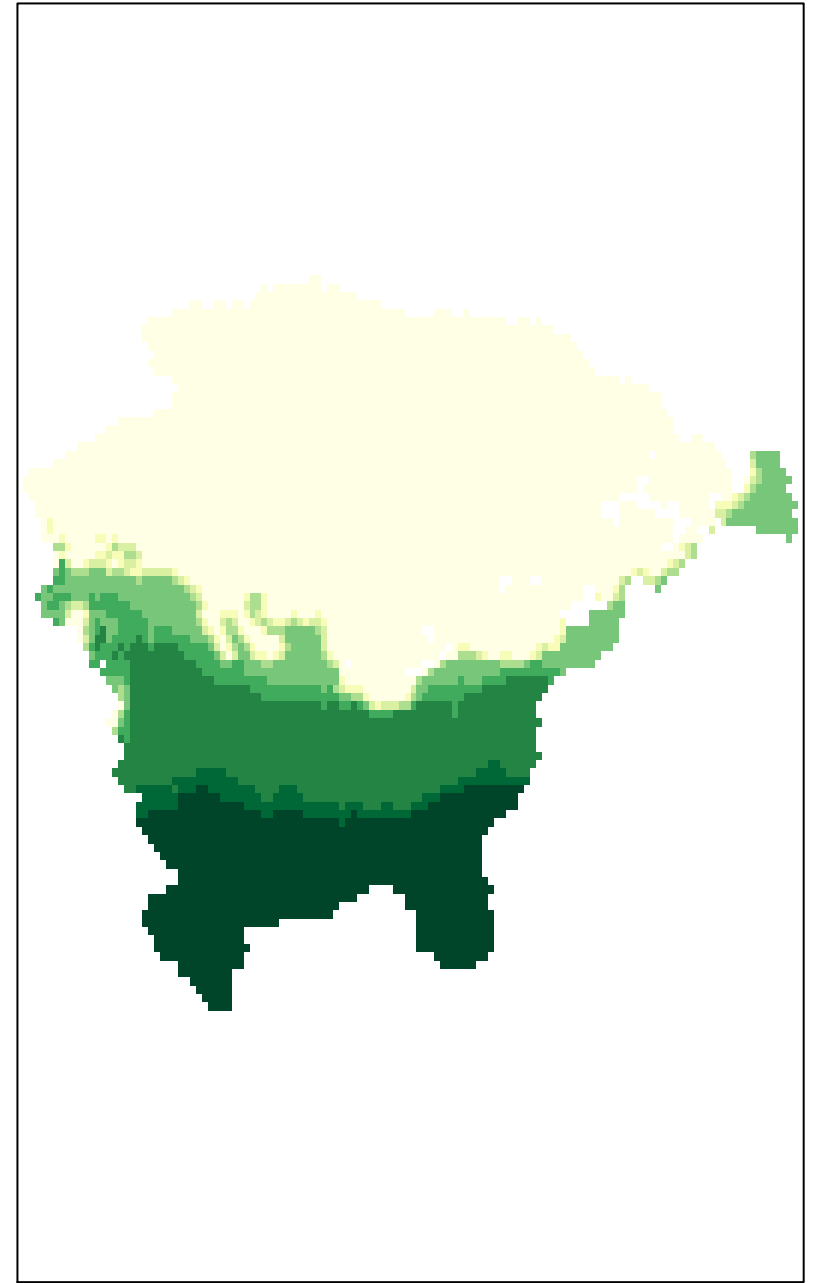
MAX, X14000.ybp



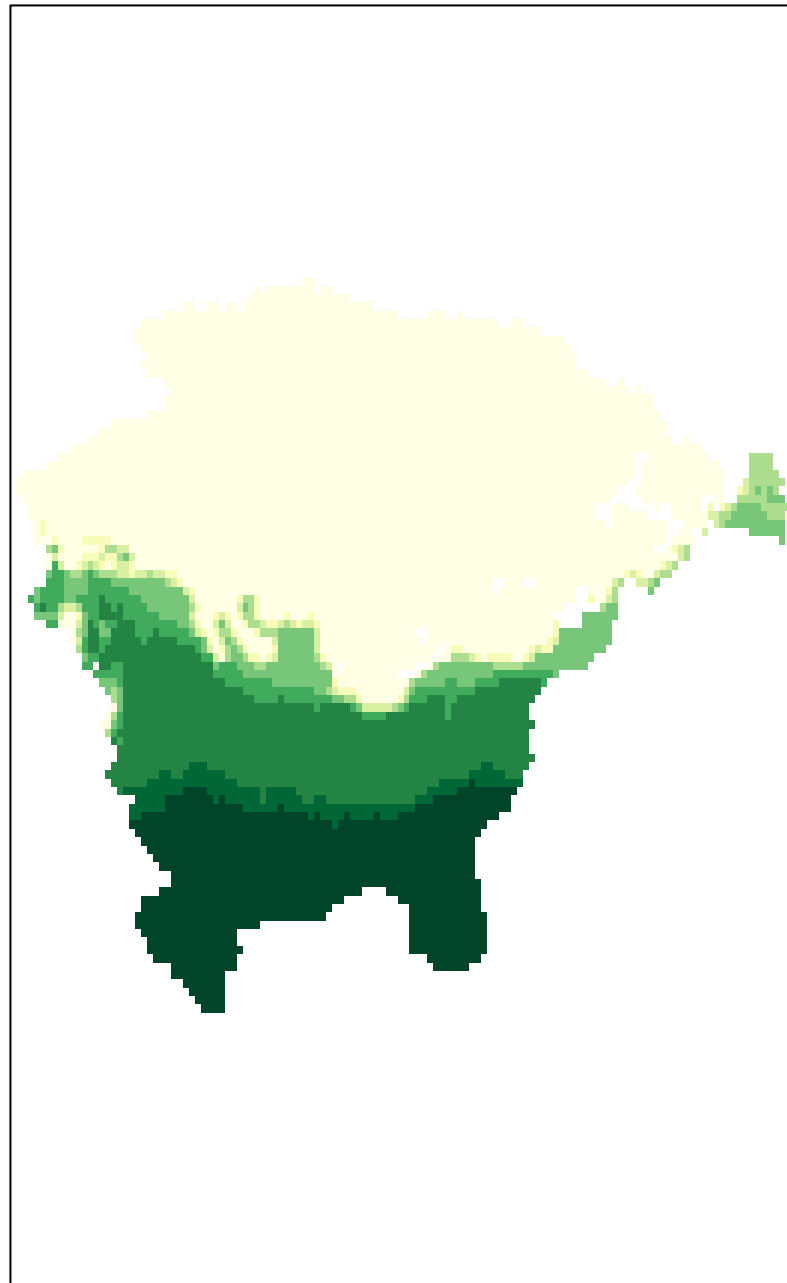
MAX, X13000.ybp



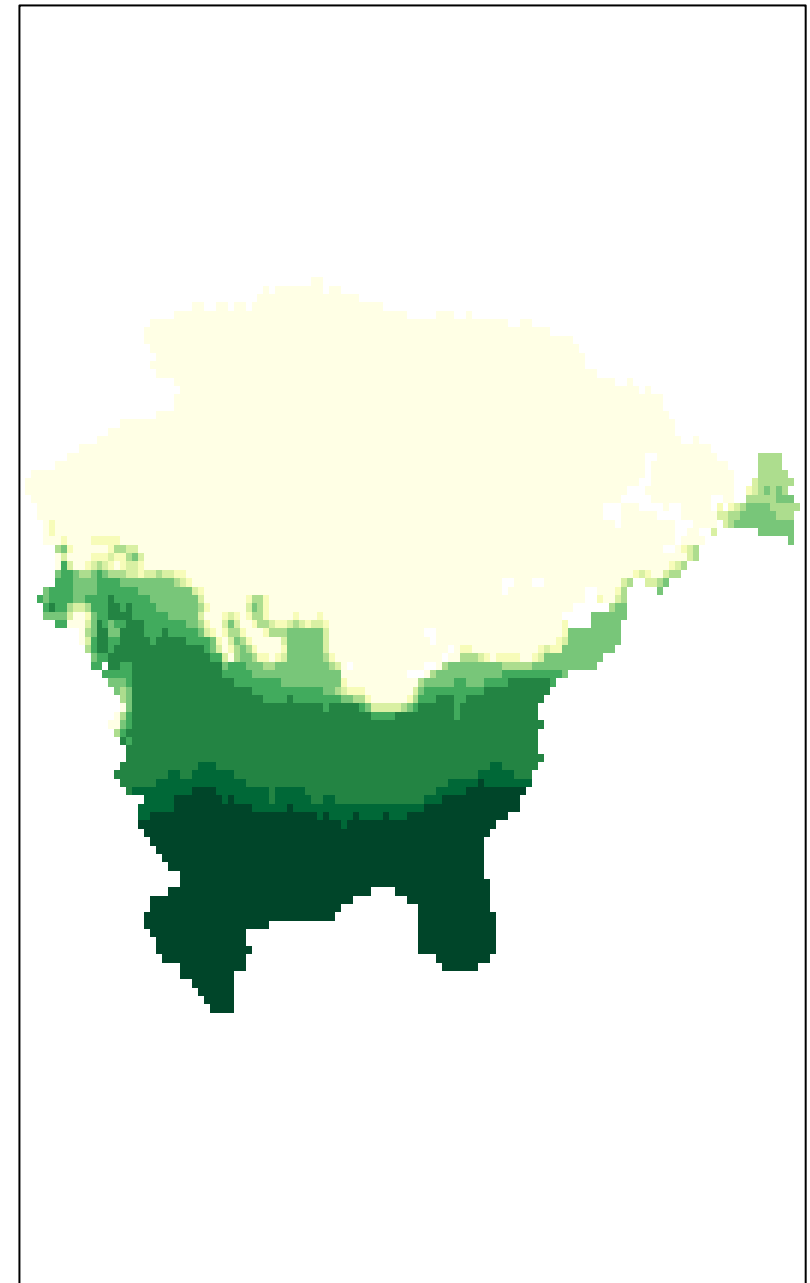
MAX, X13000.ybp



MAX, X12000.ybp

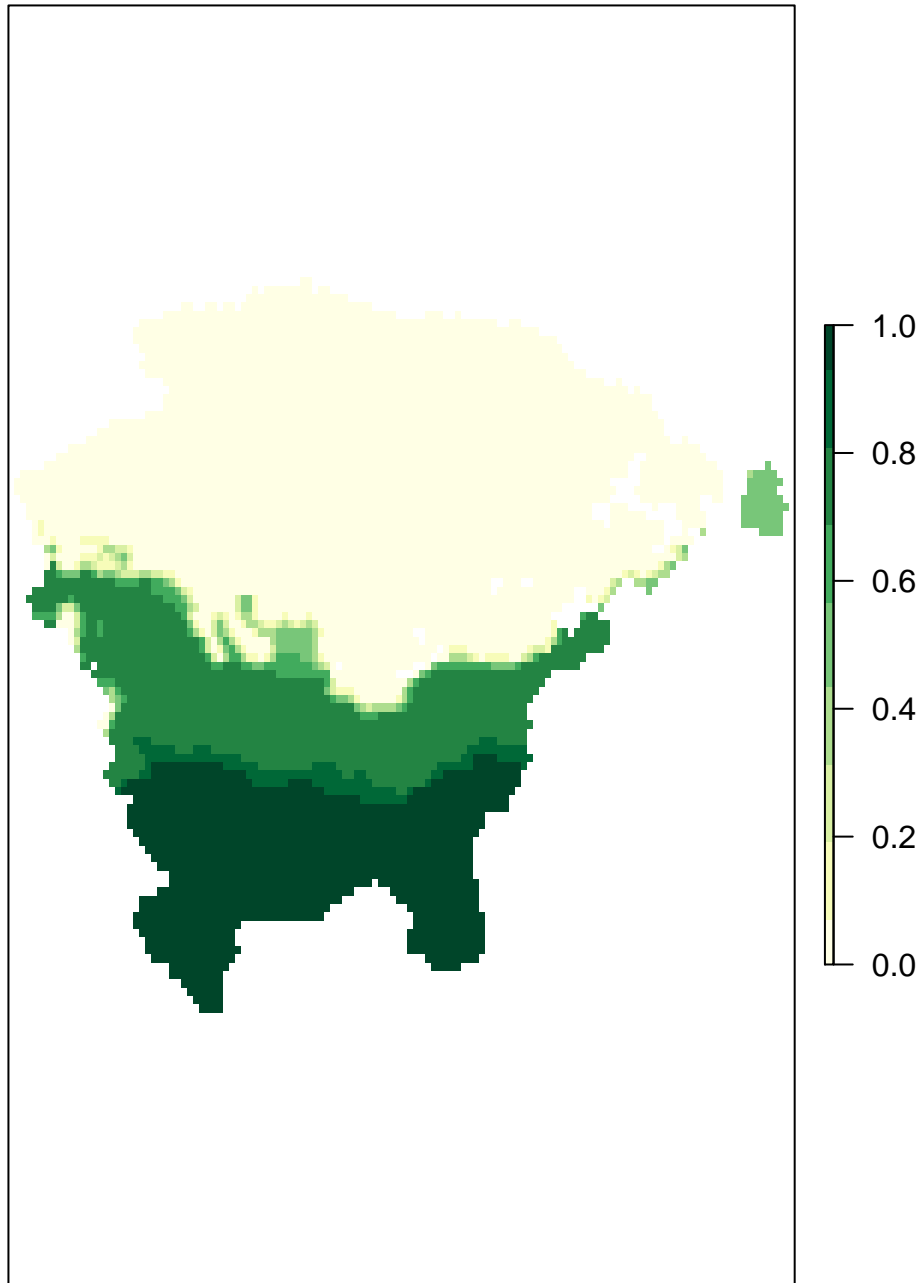


MAX, X12000.ybp

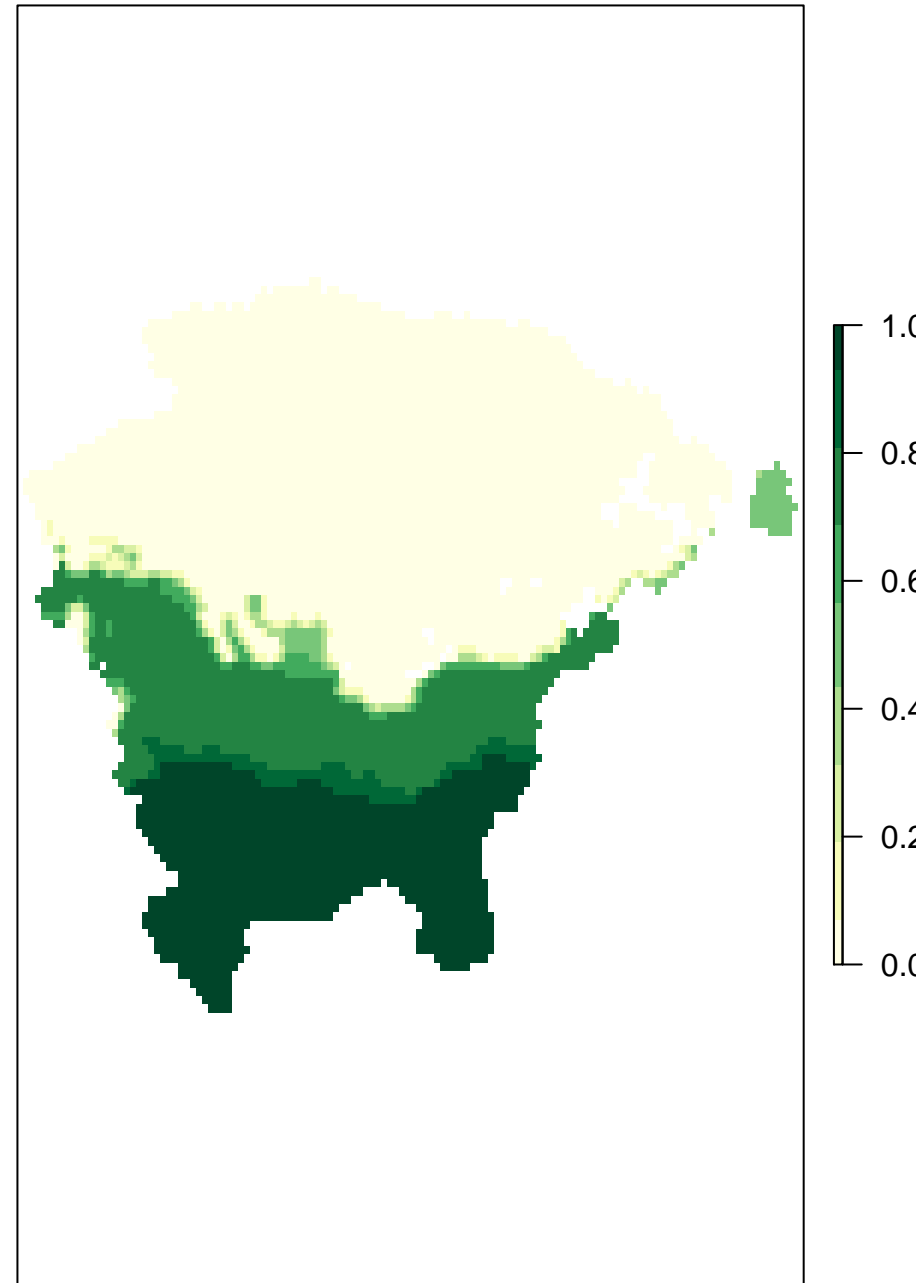


Species skipped = *Fraxinus caroliniana*, GCM = Lorenz_ccsm

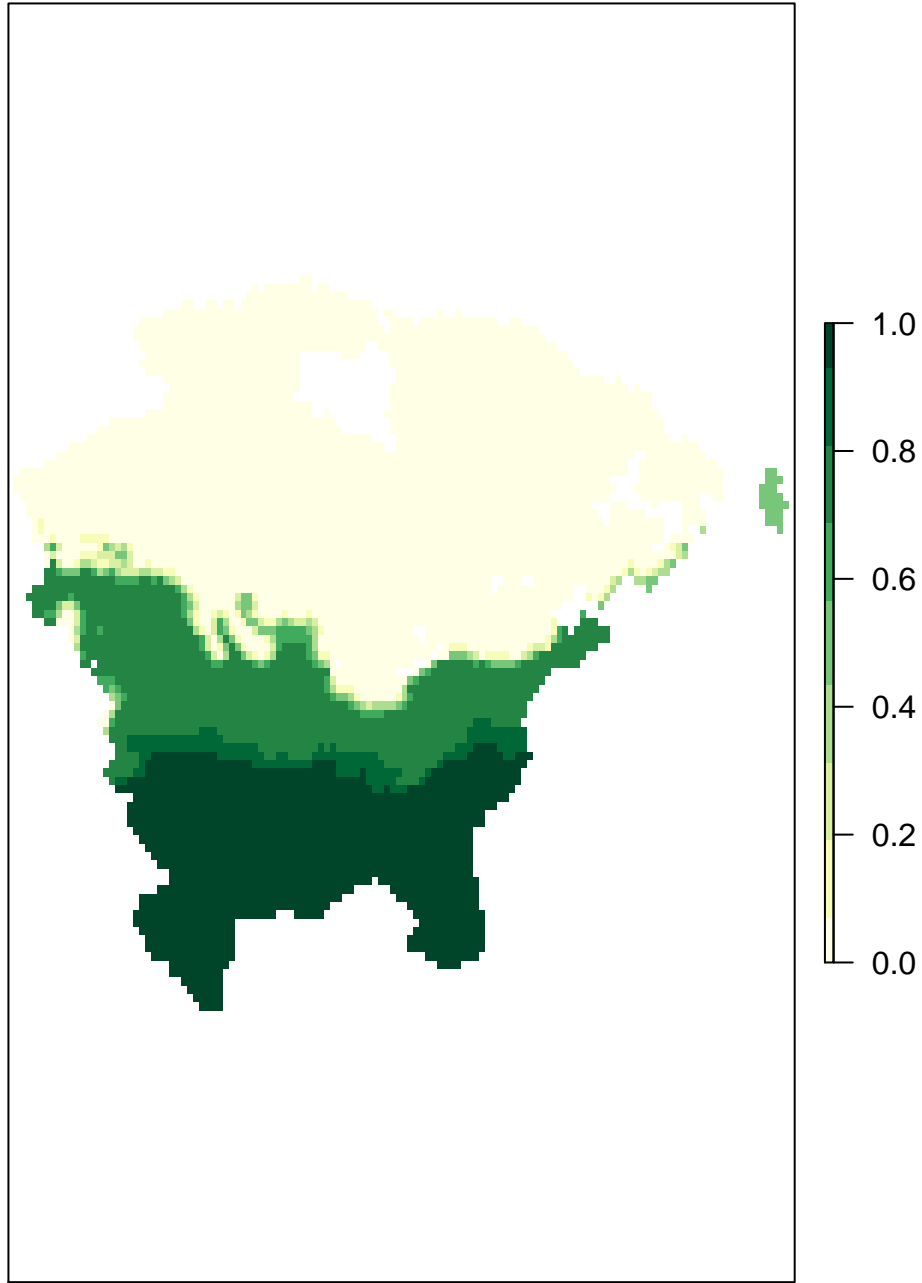
MAX, X11000.ybp



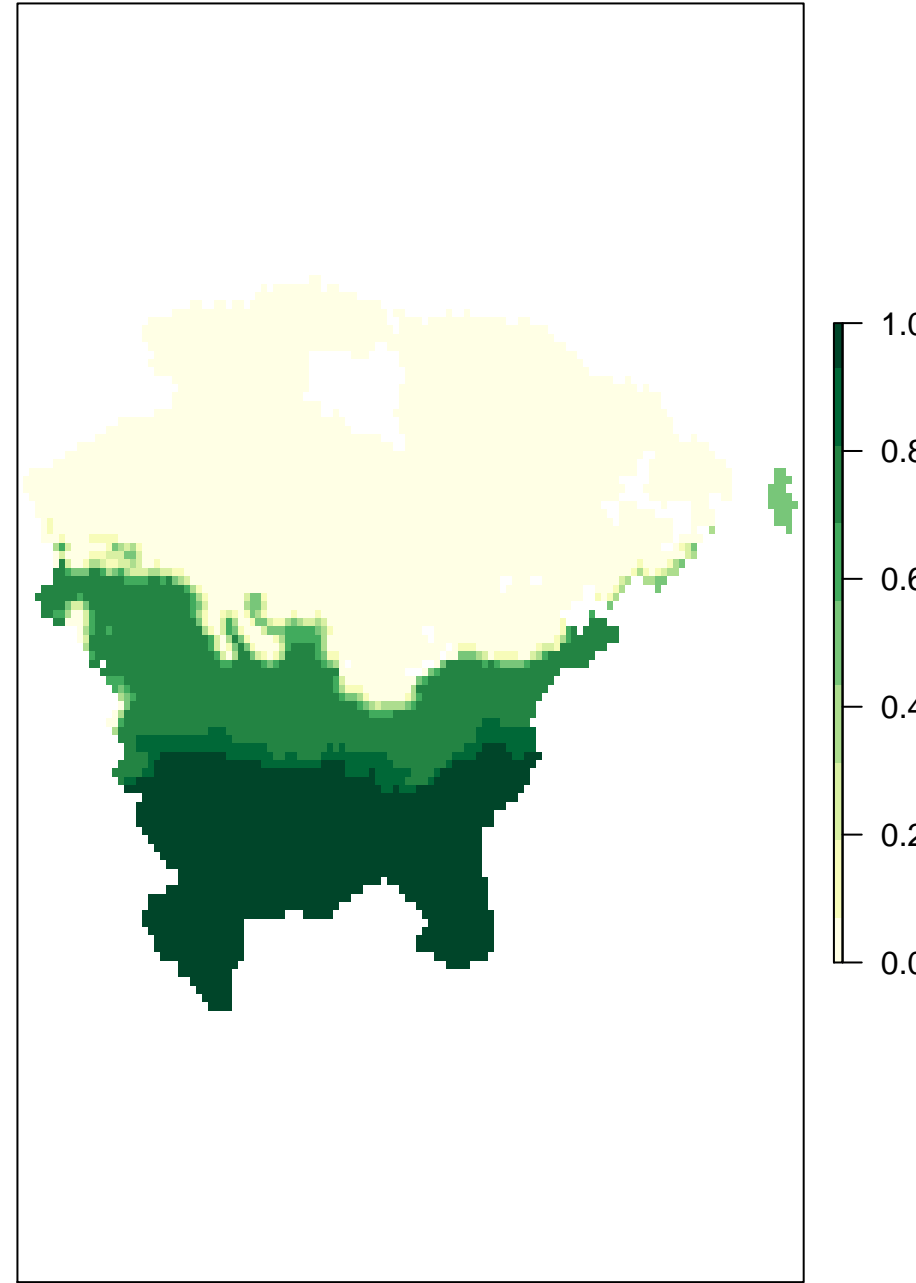
MAX, X11000.ybp



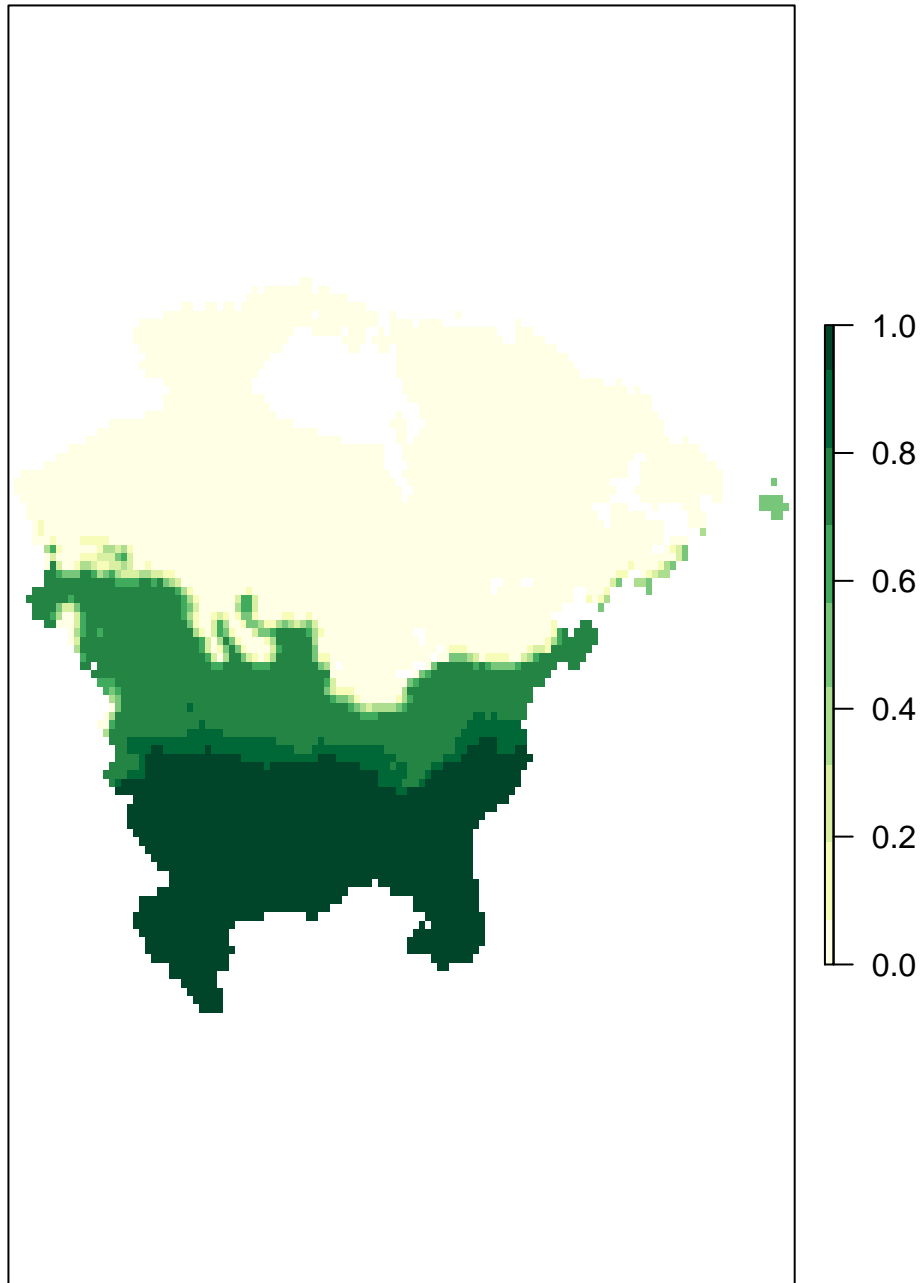
MAX, X10000.ybp



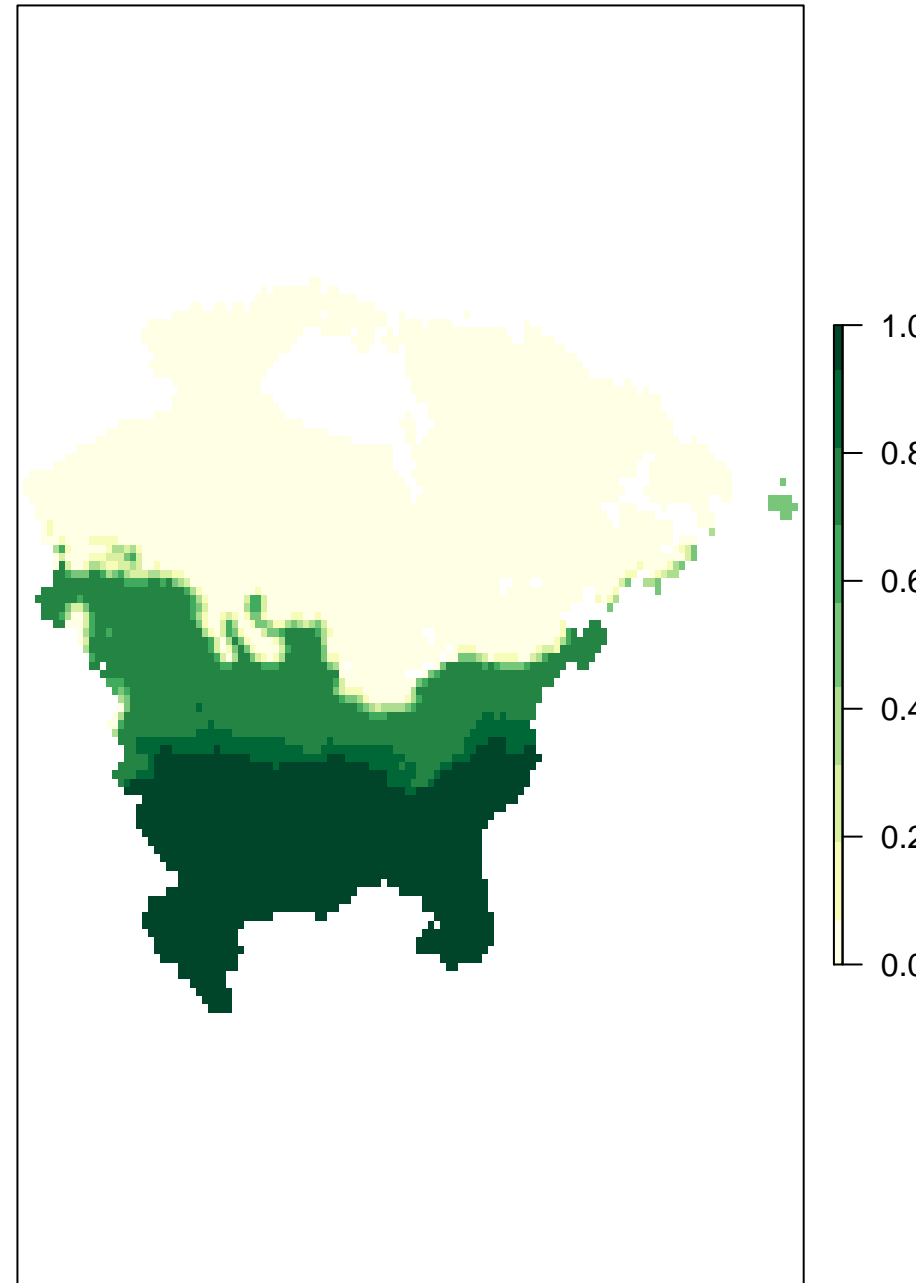
MAX, X10000.ybp



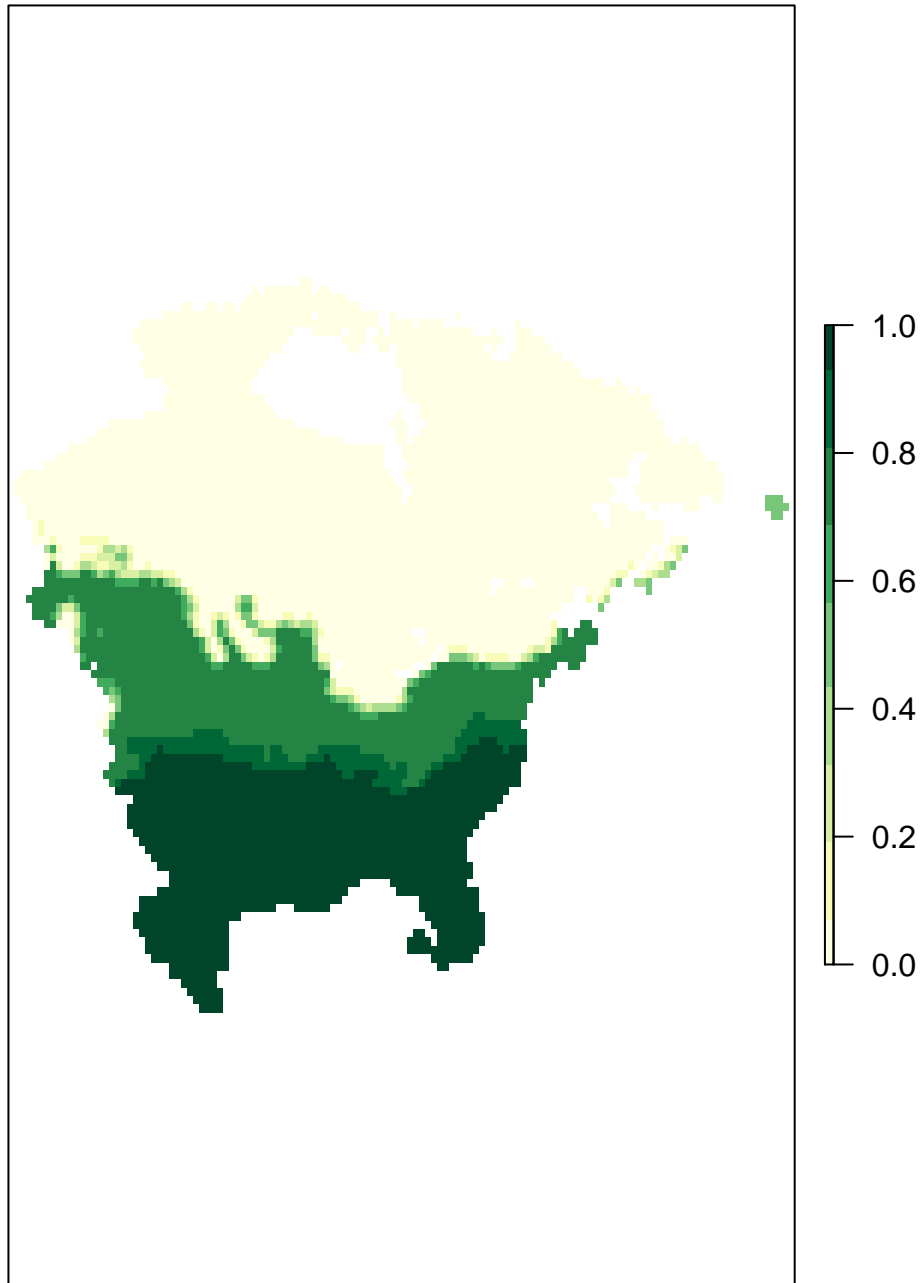
MAX, X9000.ybp



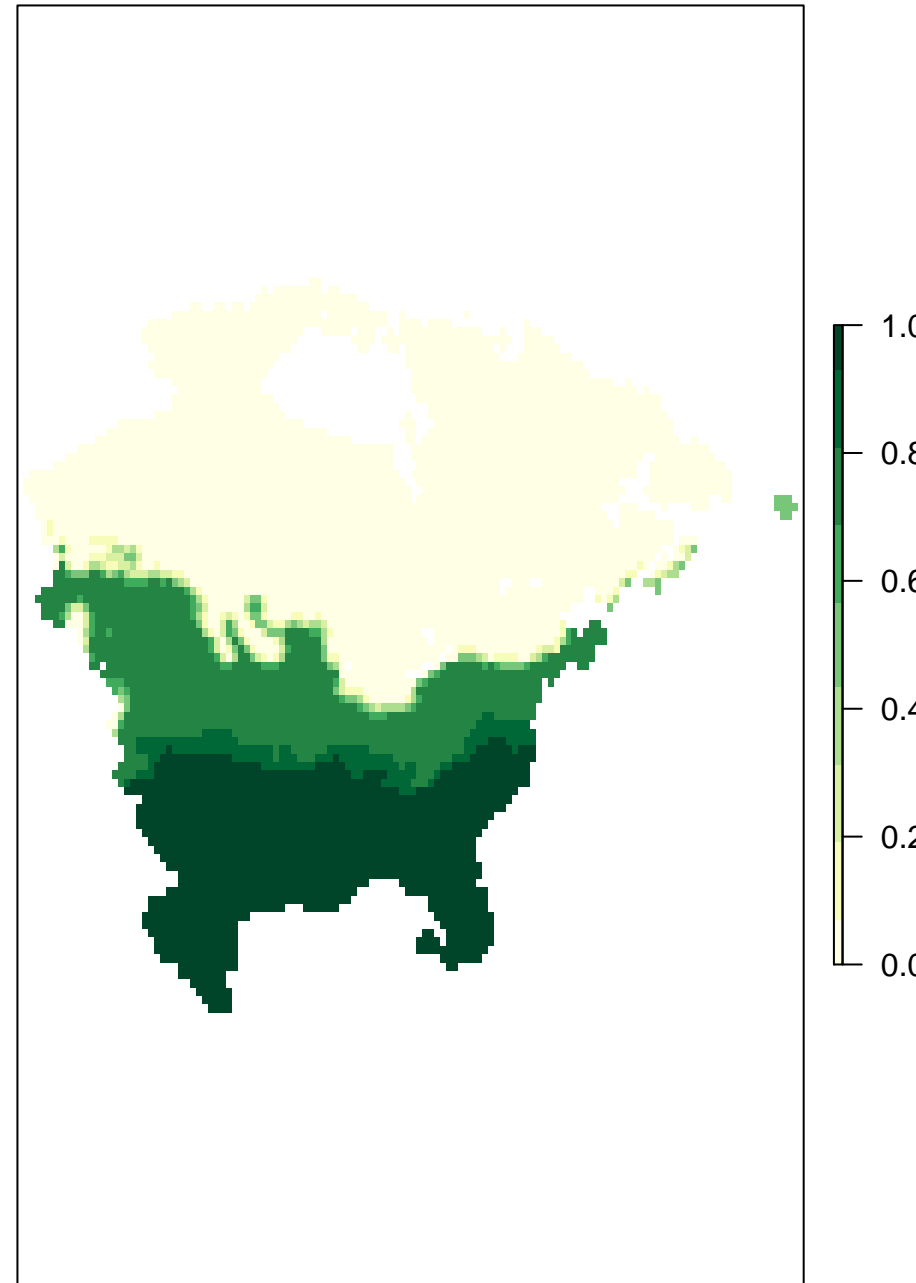
MAX, X9000.ybp



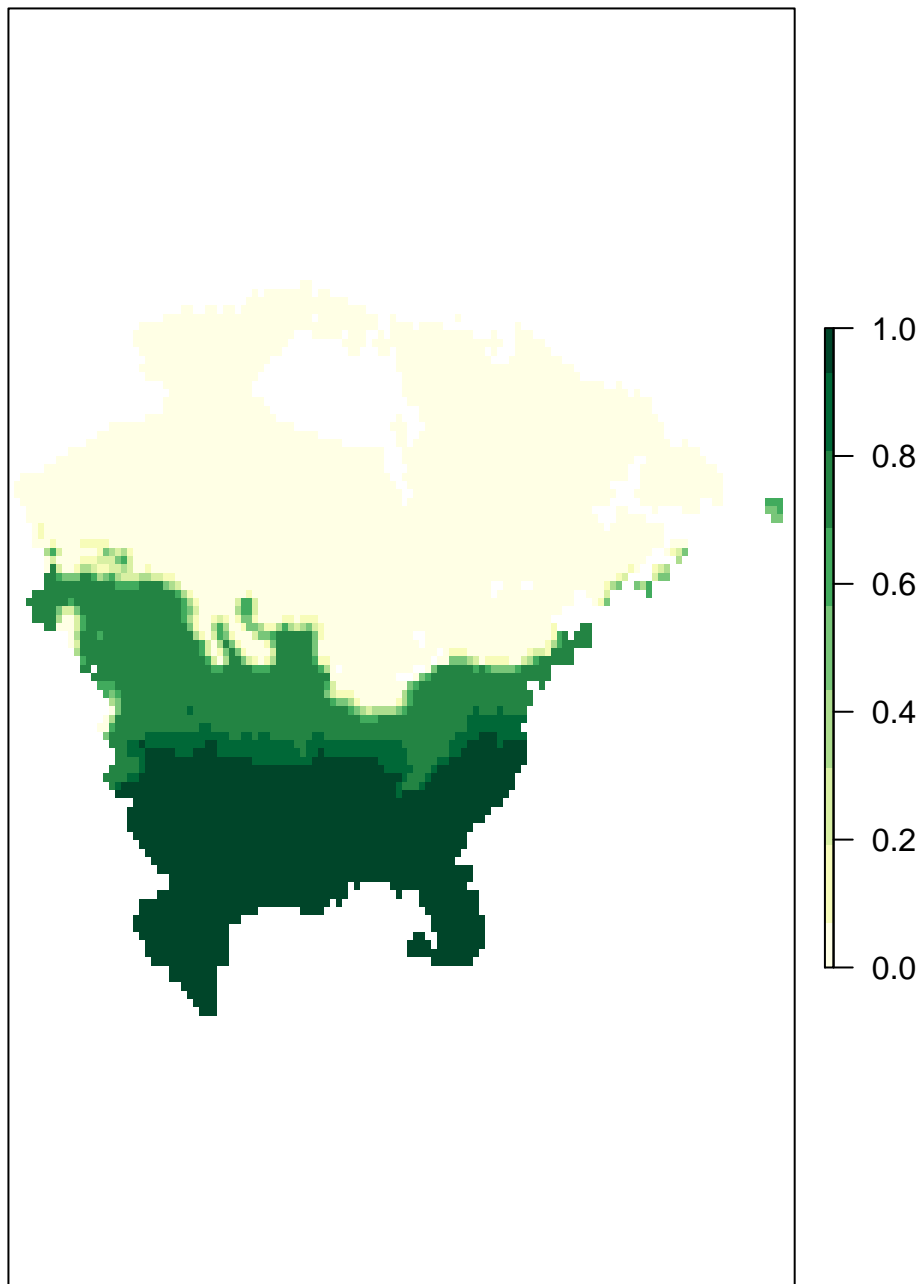
MAX, X8000.ybp



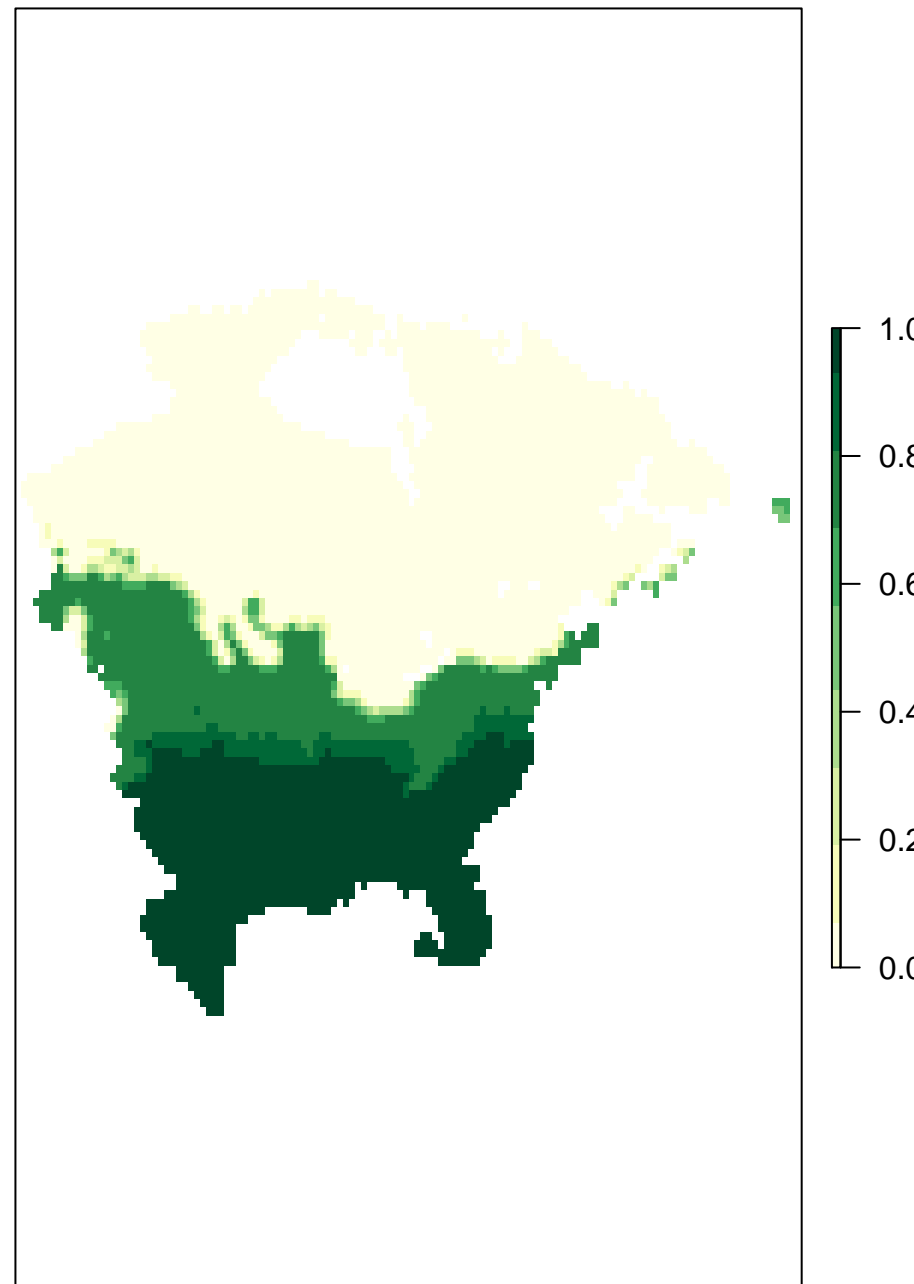
MAX, X8000.ybp



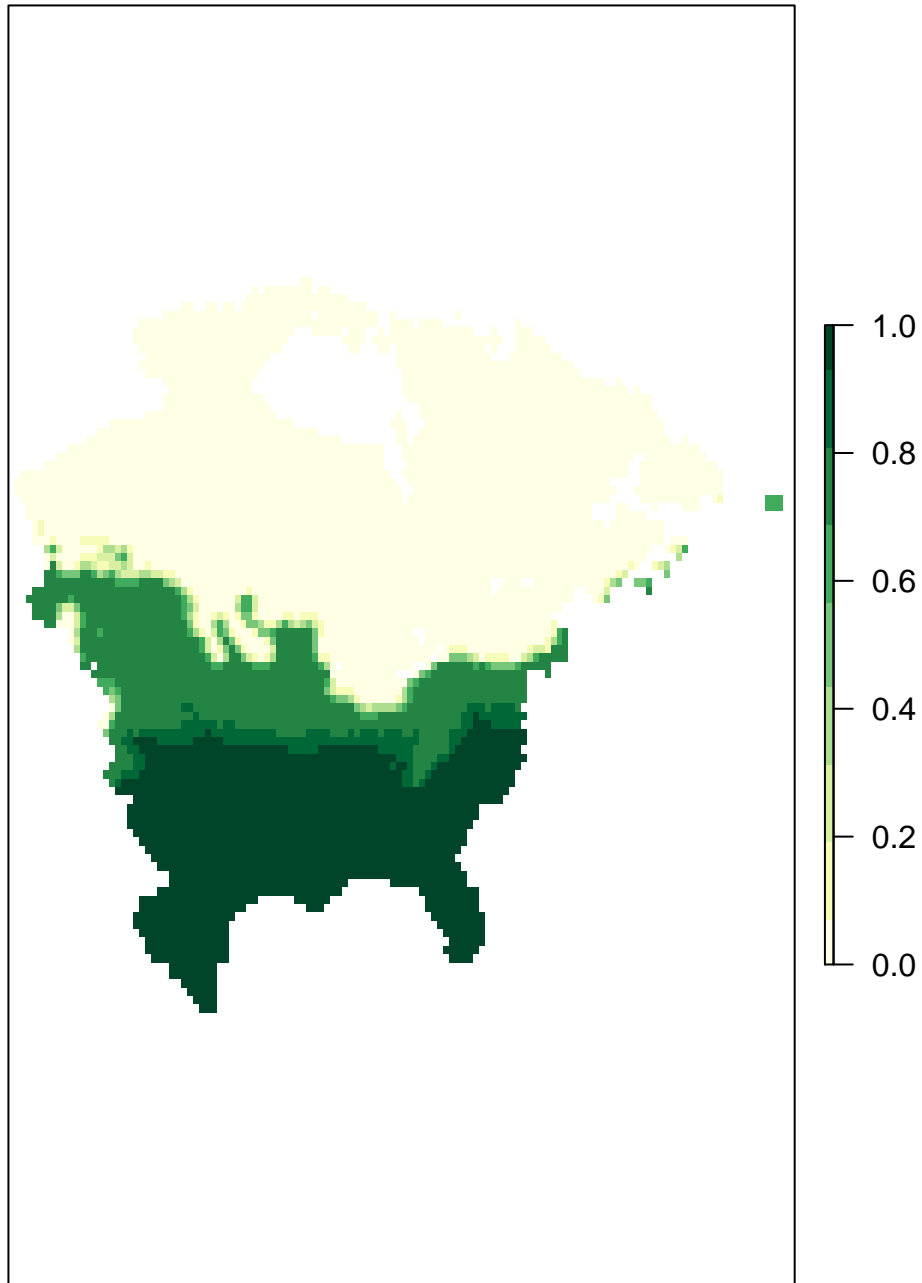
MAX, X7000.ybp



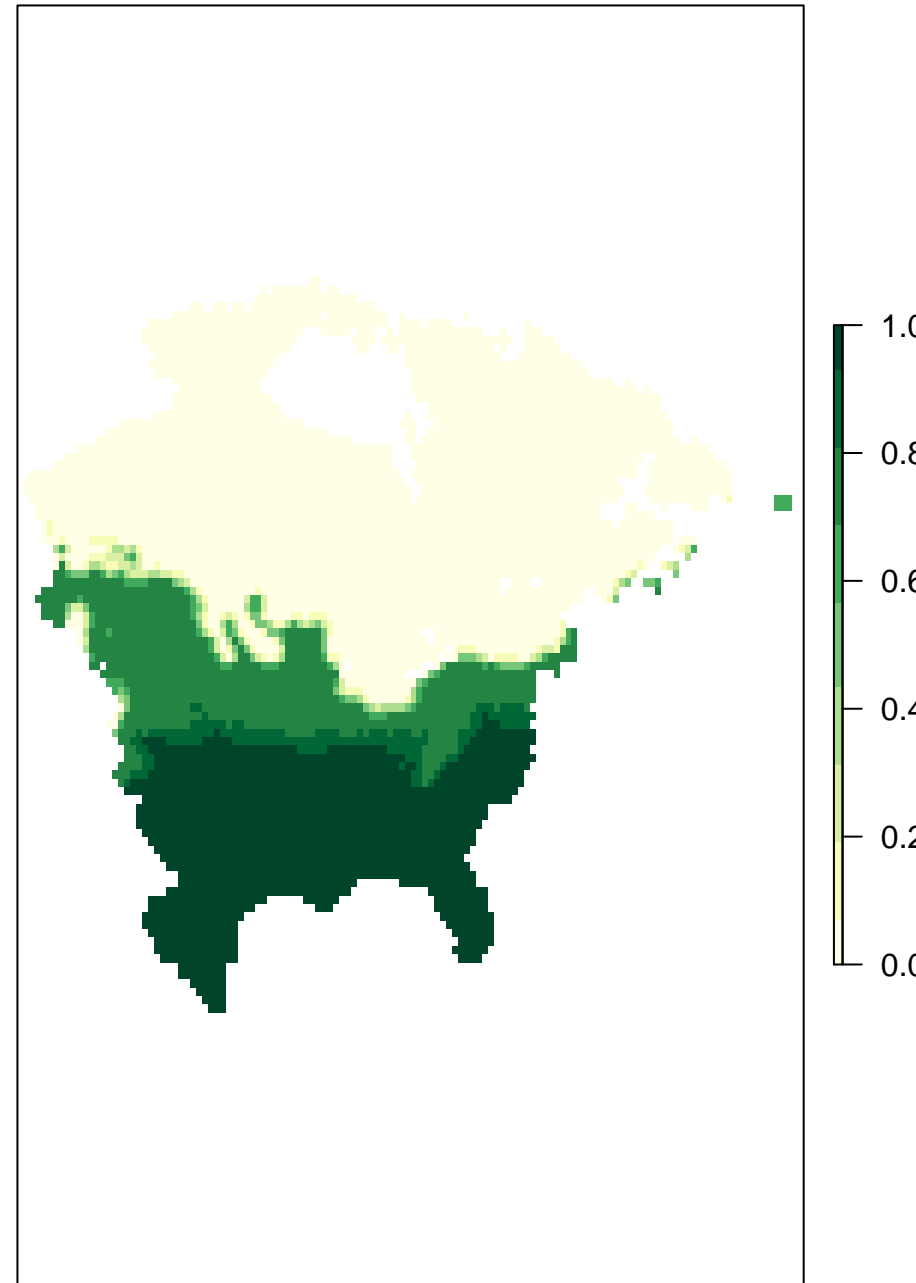
MAX, X7000.ybp



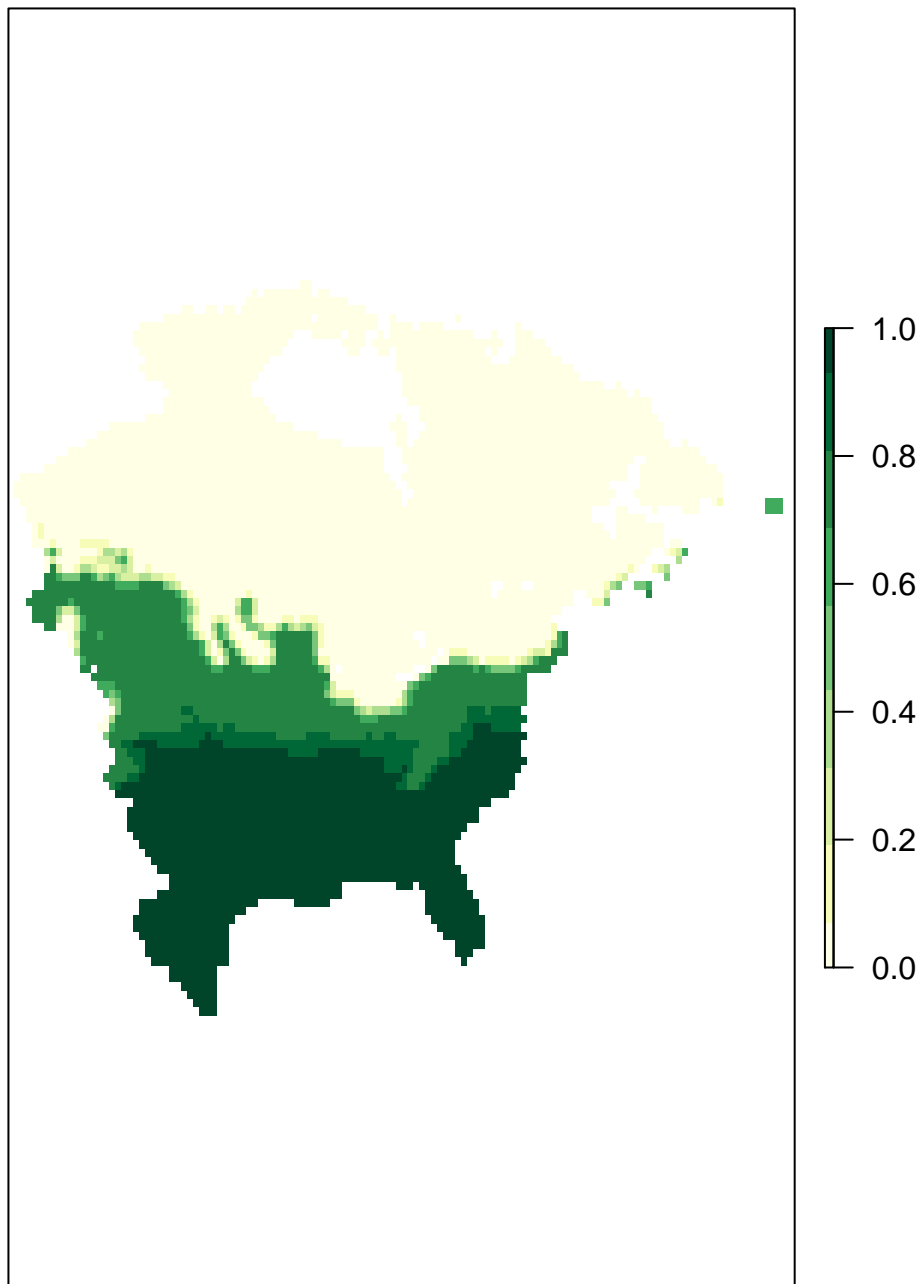
MAX, X6000.ybp



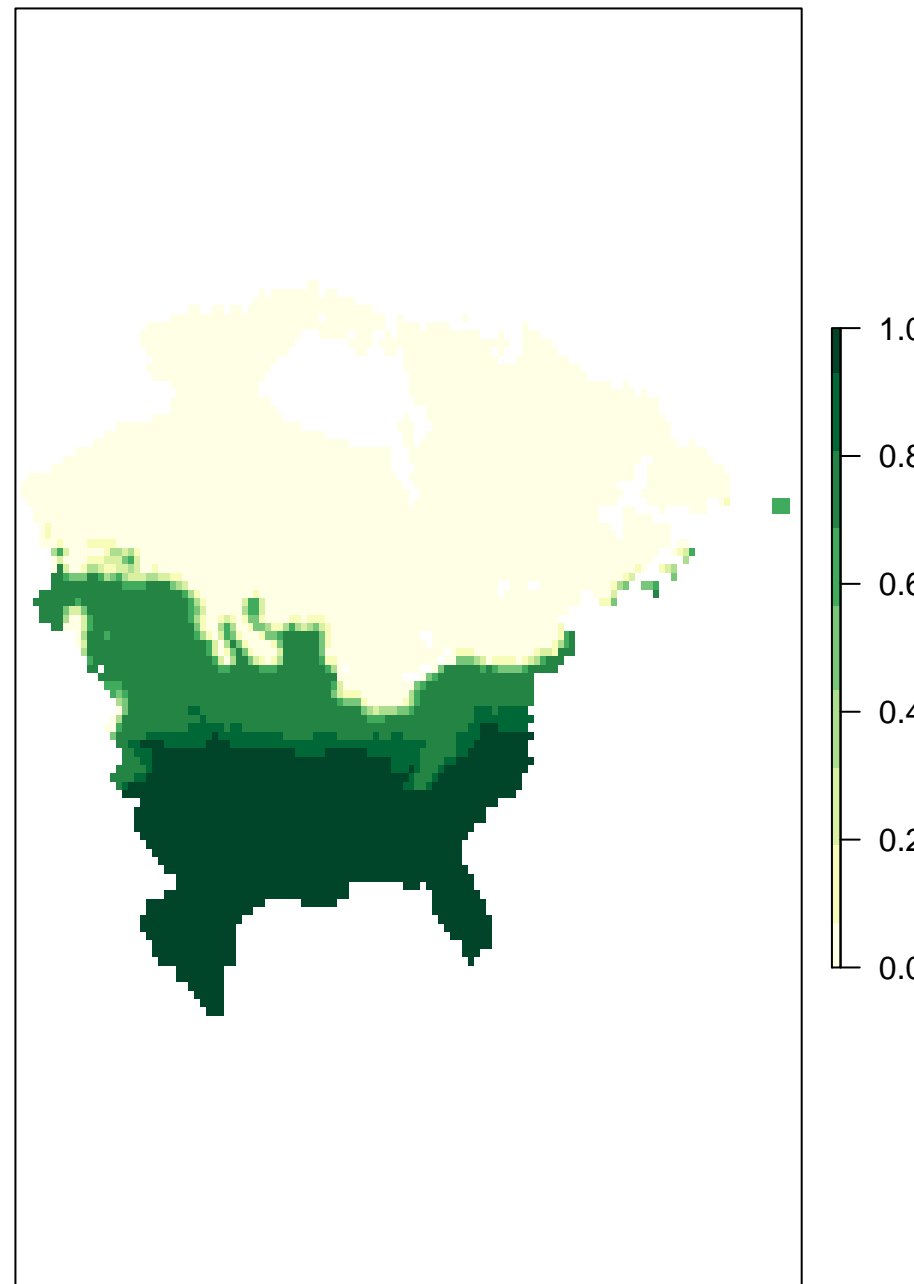
MAX, X6000.ybp



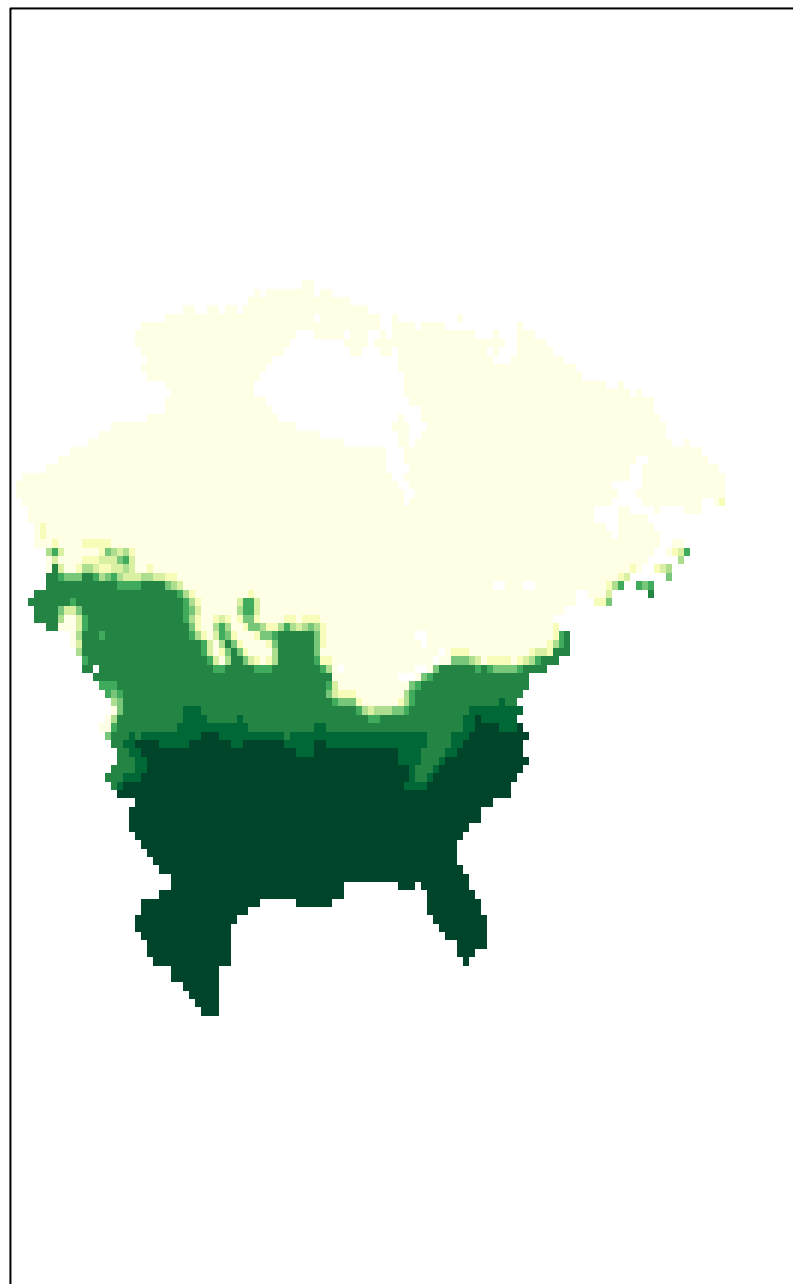
MAX, X5000.ybp



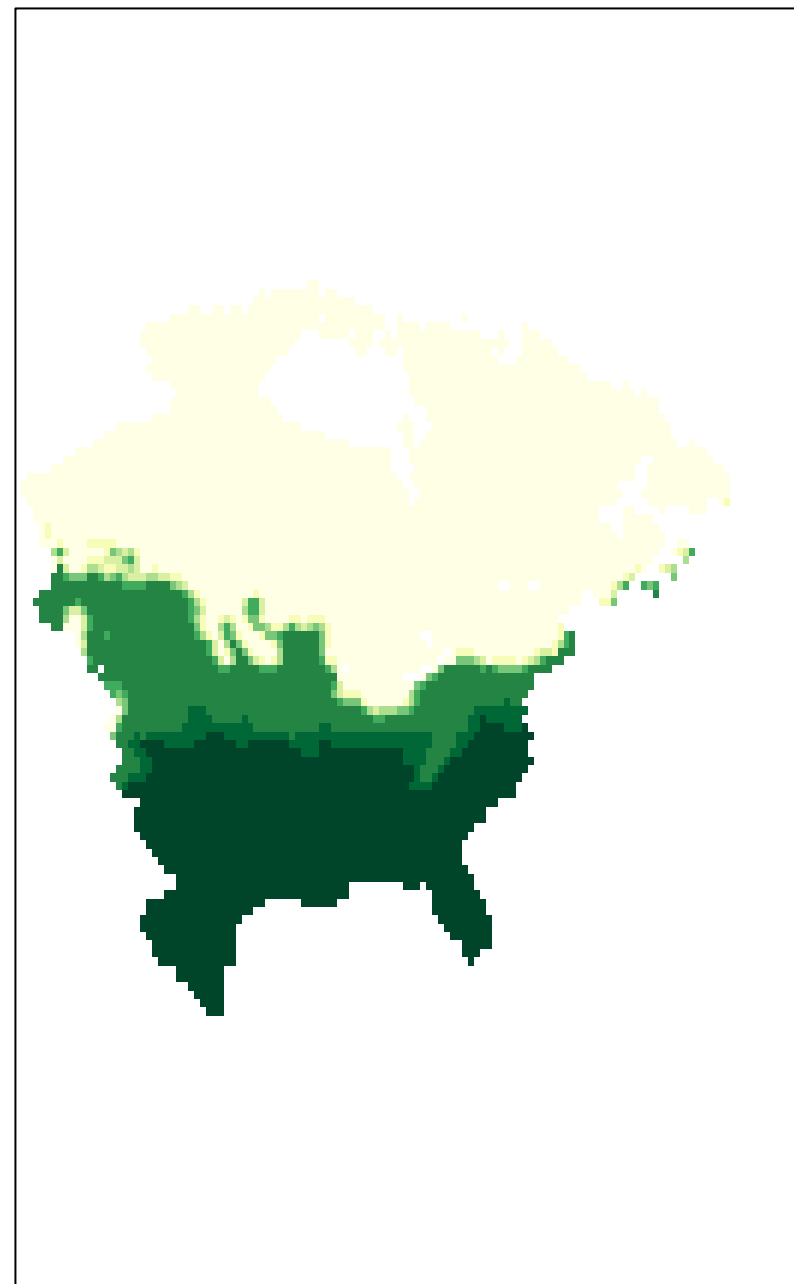
MAX, X5000.ybp



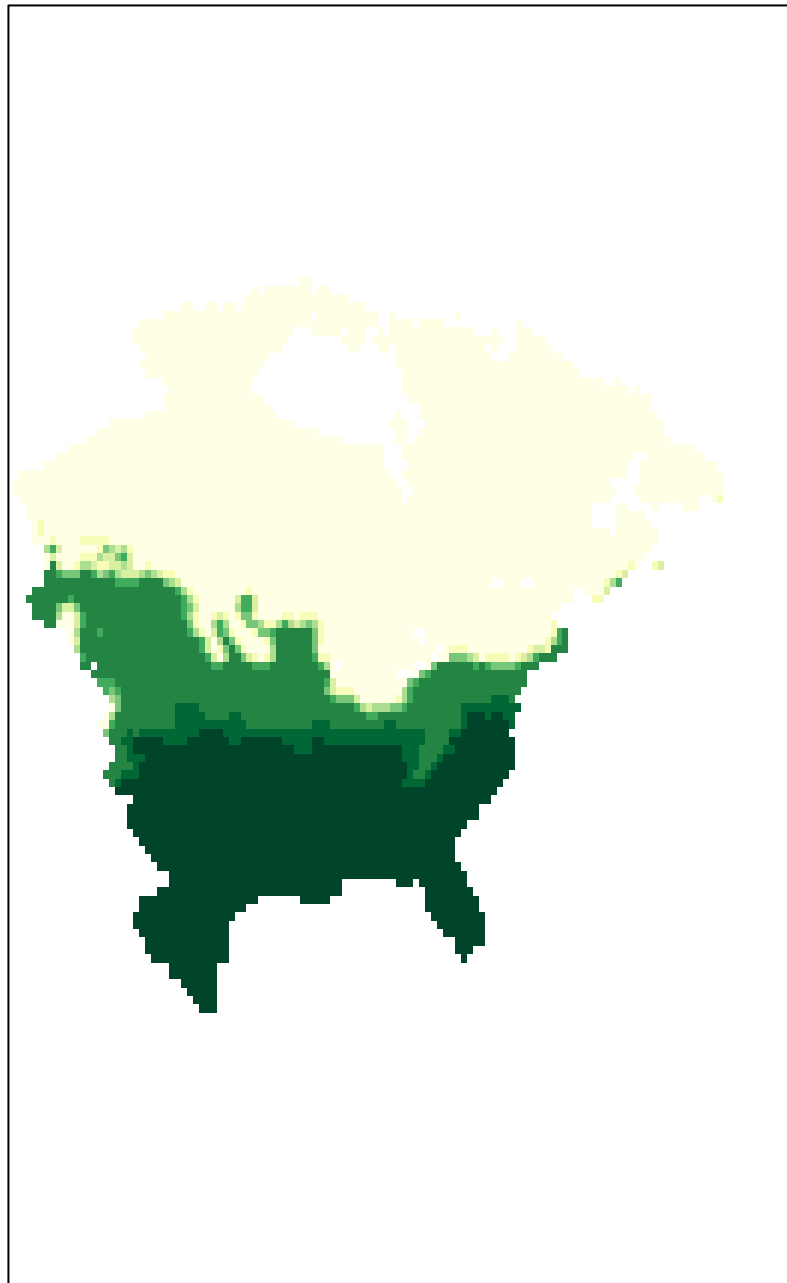
MAX, X4000.ybp



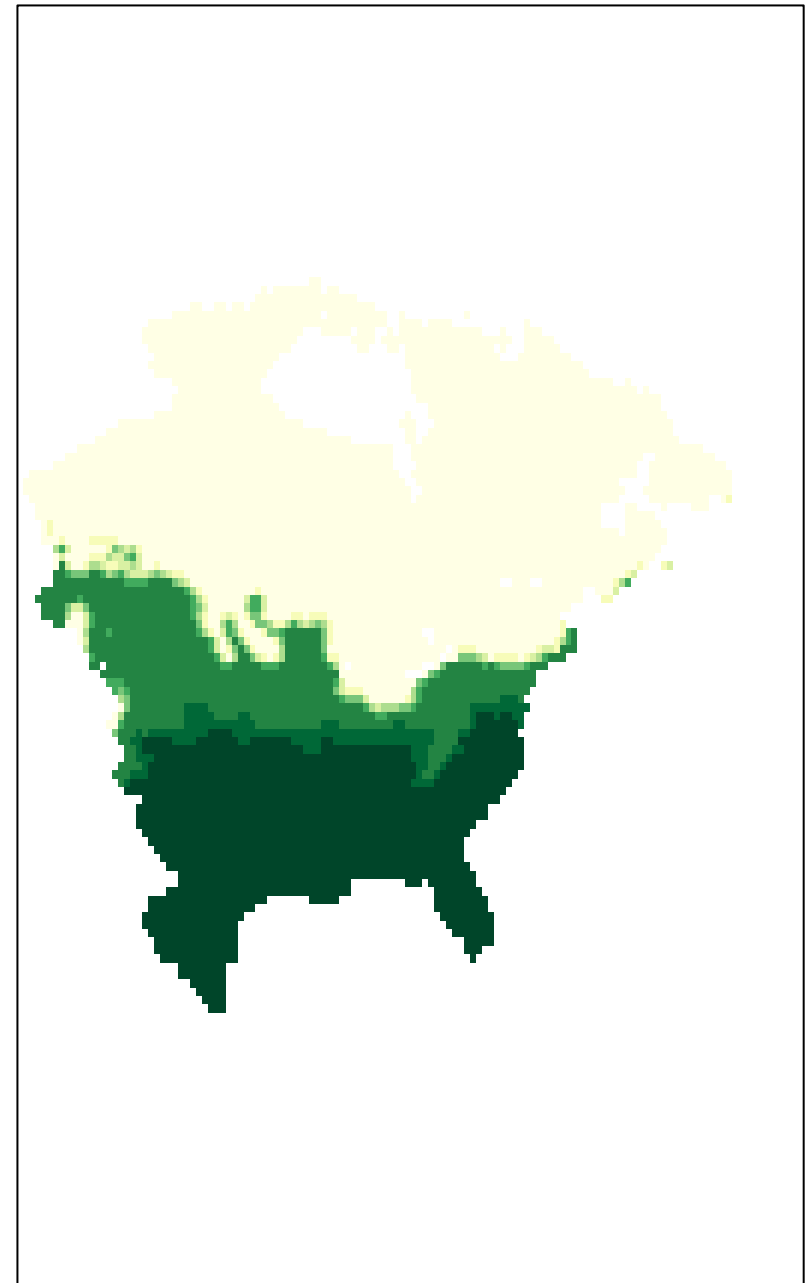
MAX, X4000.ybp



MAX, X3000.ybp

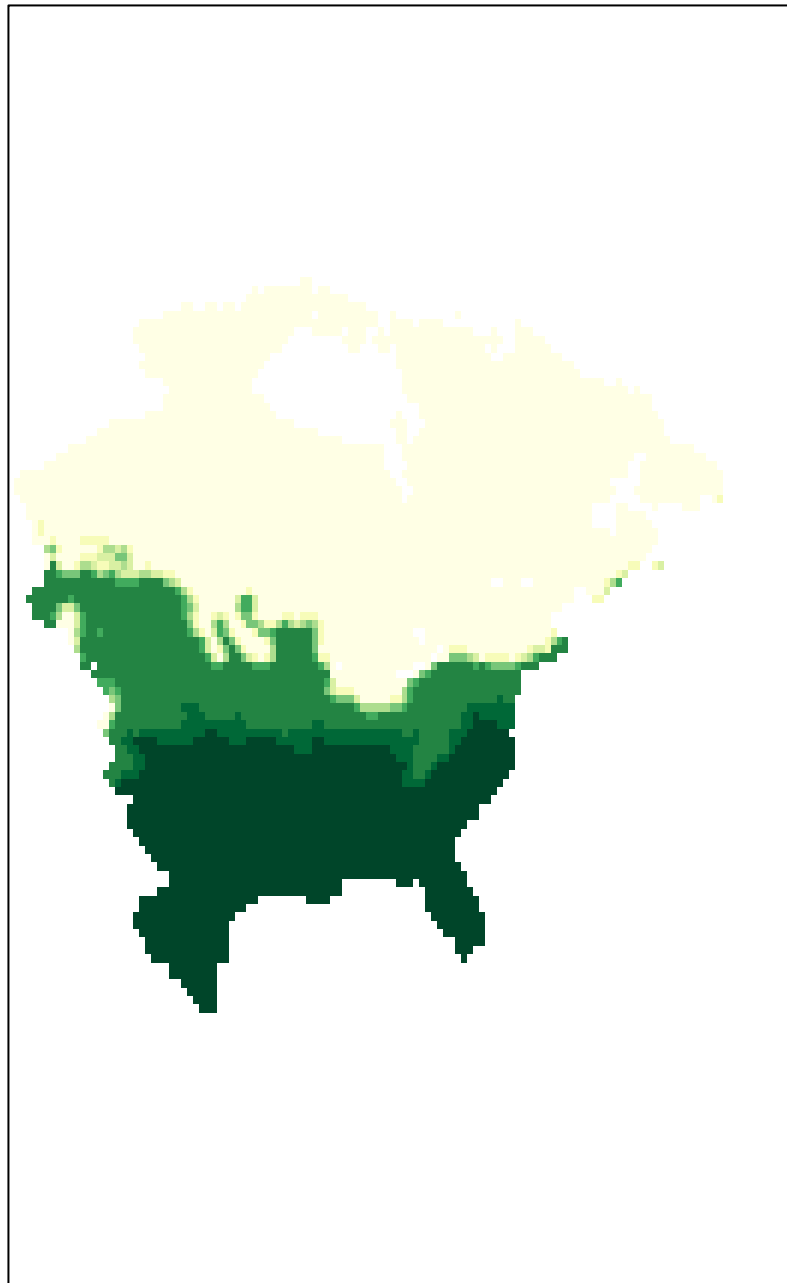


MAX, X3000.ybp

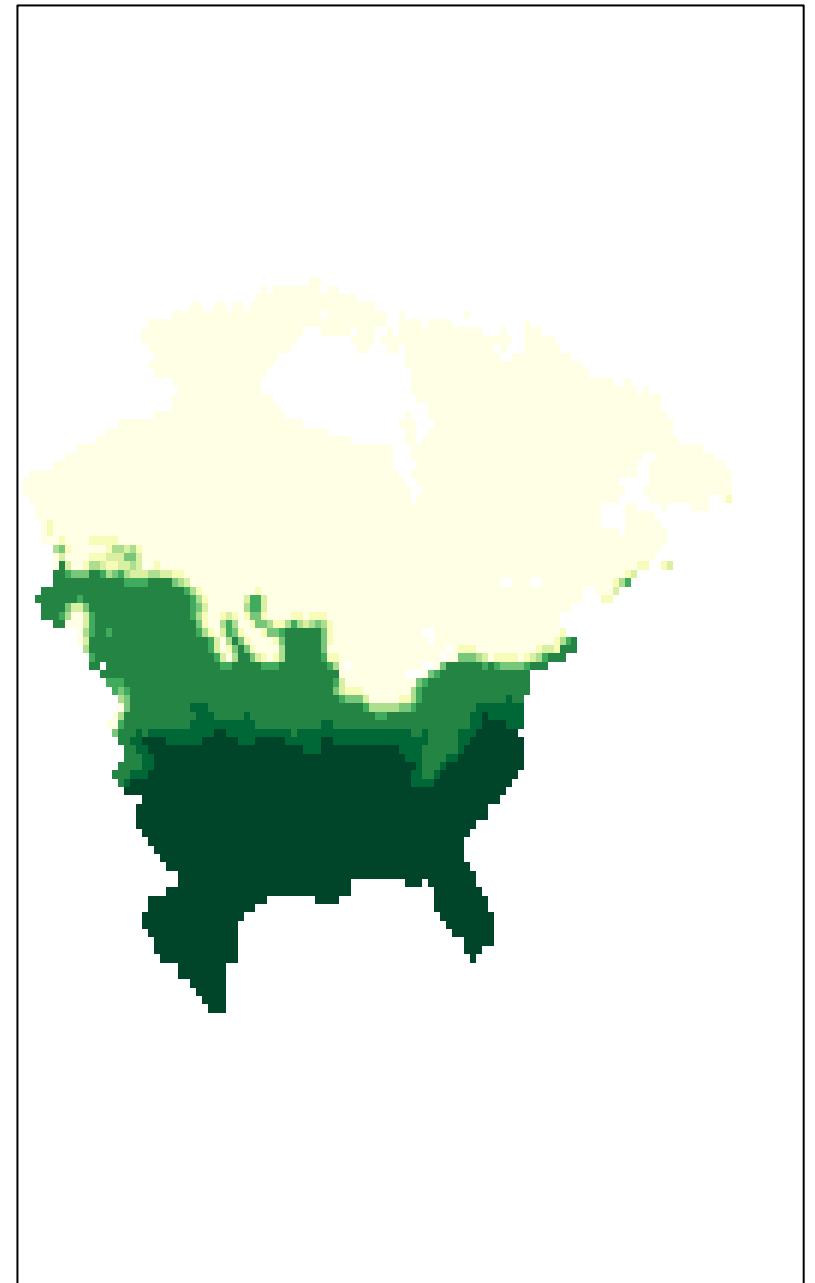


Species skipped = *Fraxinus caroliniana*, GCM = Lorenz_ccsm

MAX, X2000.ybp

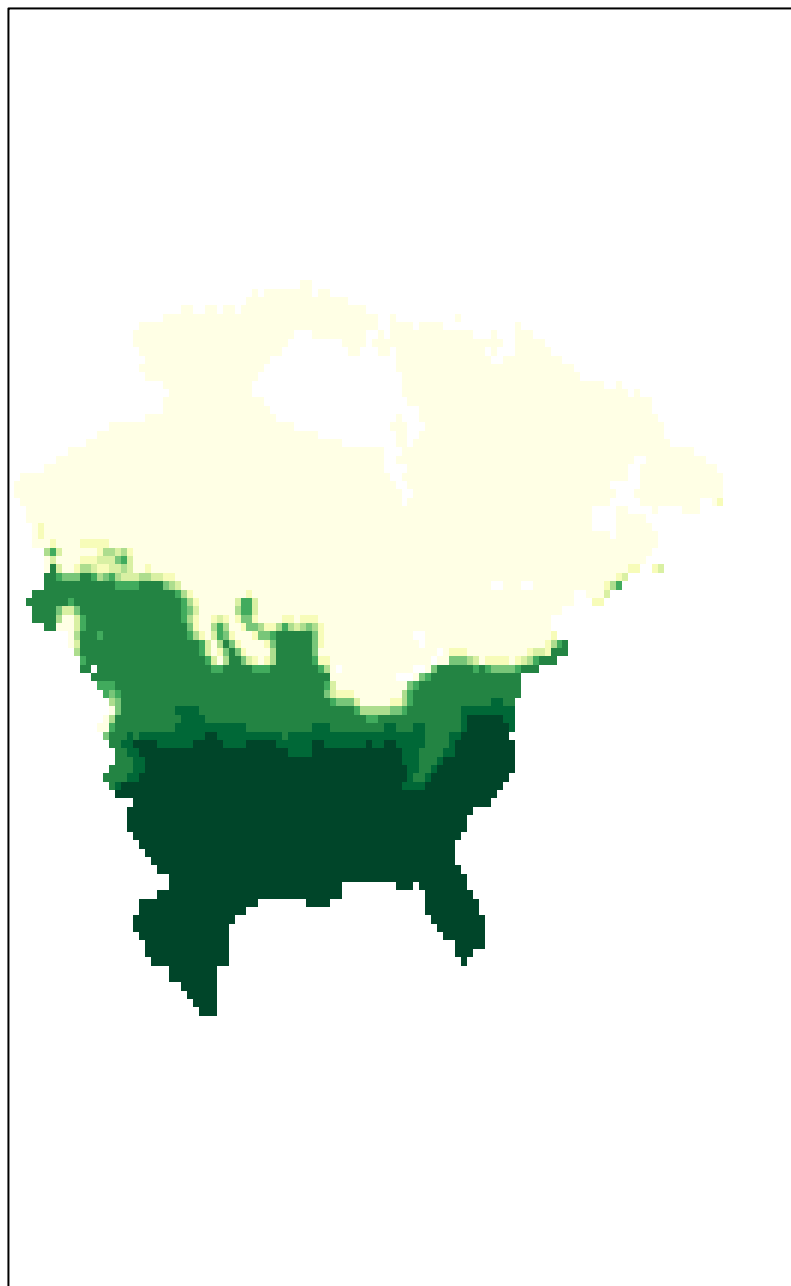


MAX, X2000.ybp

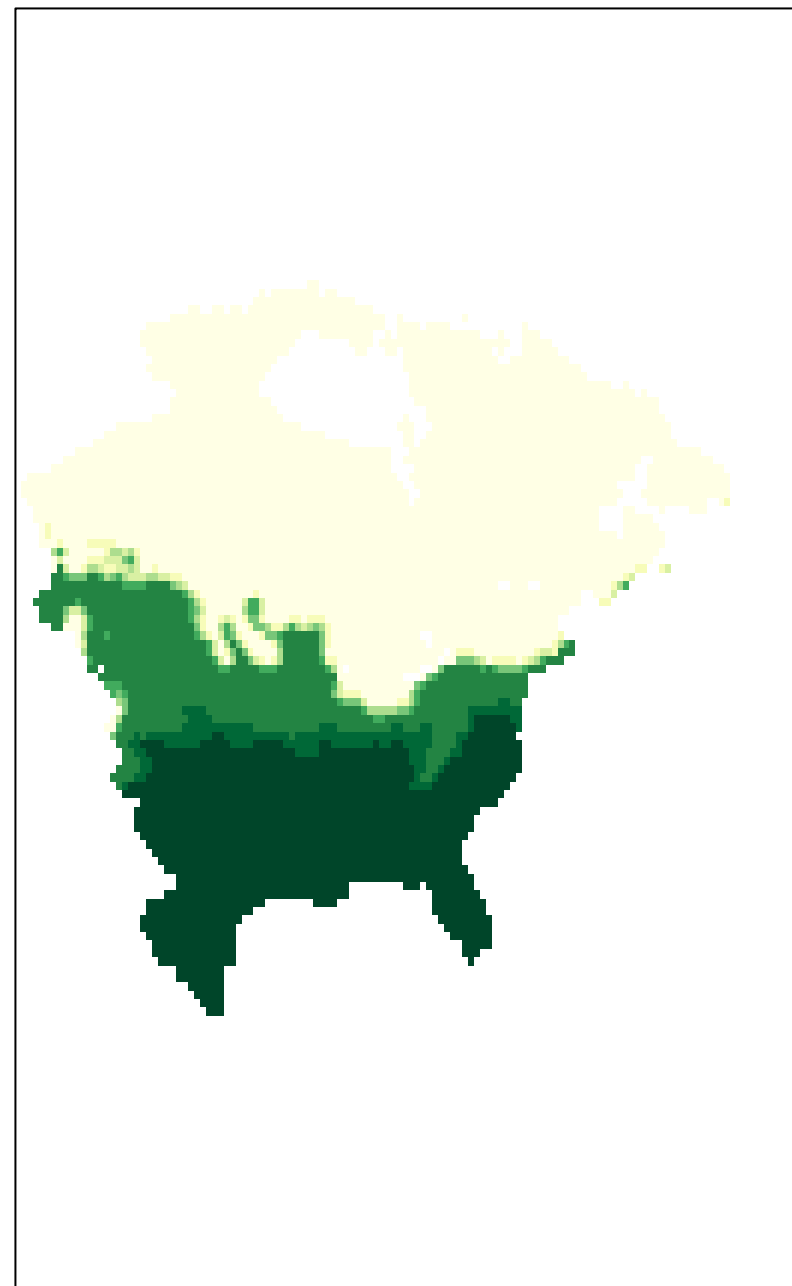


Species skipped = *Fraxinus caroliniana*, GCM = Lorenz_ccsm

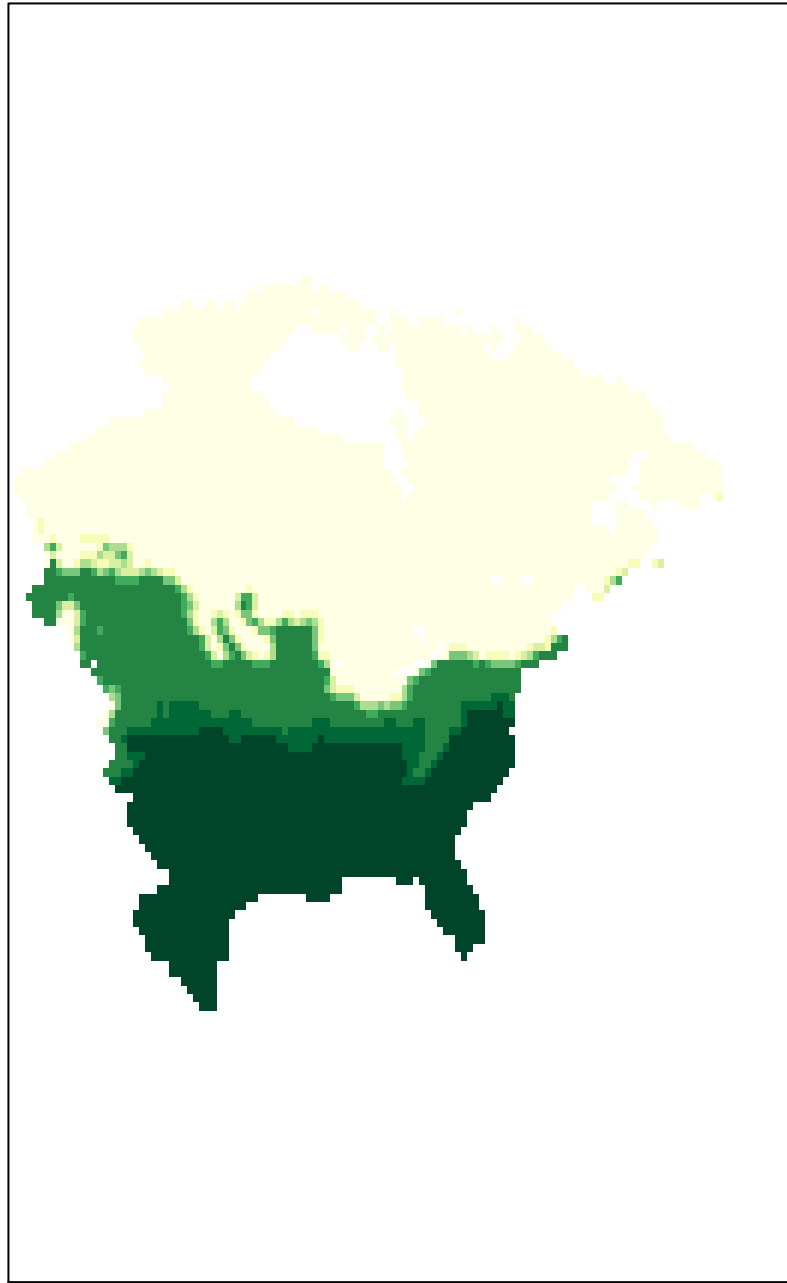
MAX, X1000.ybp



MAX, X1000.ybp



MAX, X0.ybp



MAX, X0.ybp

