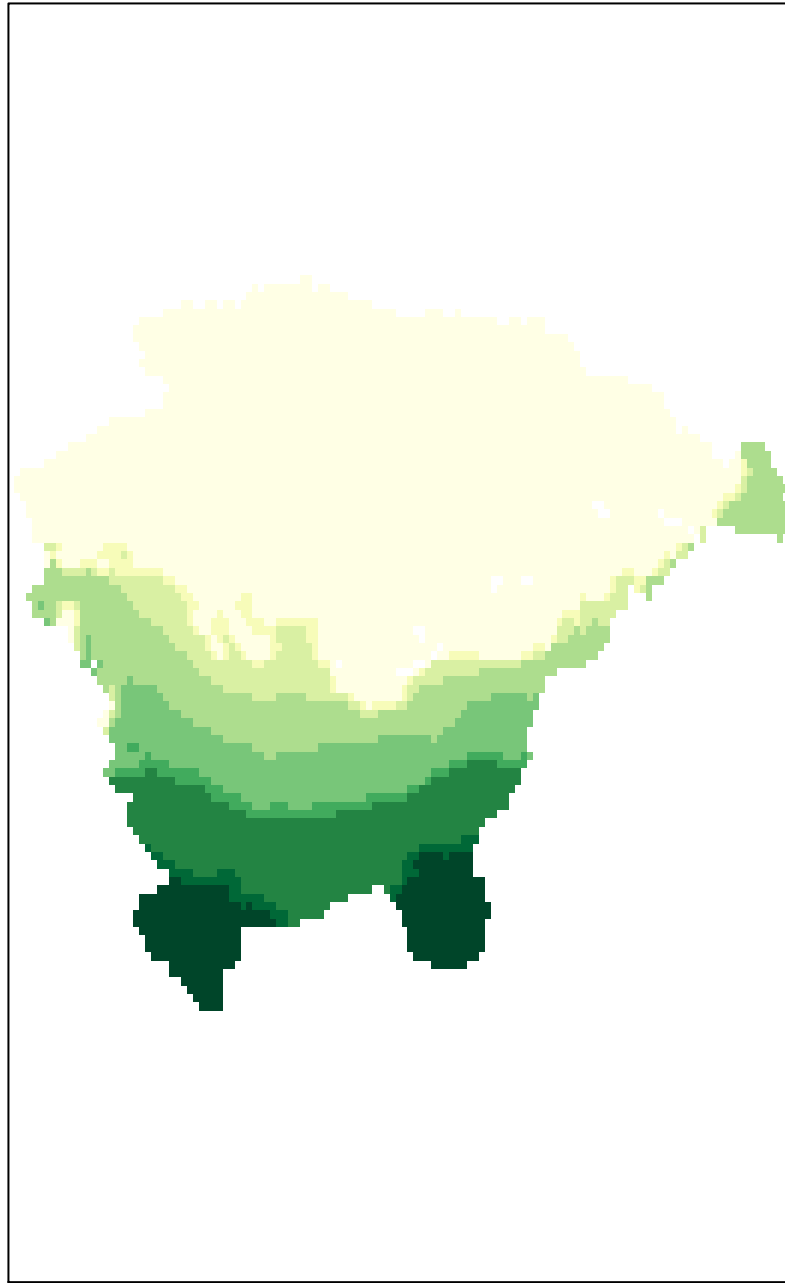
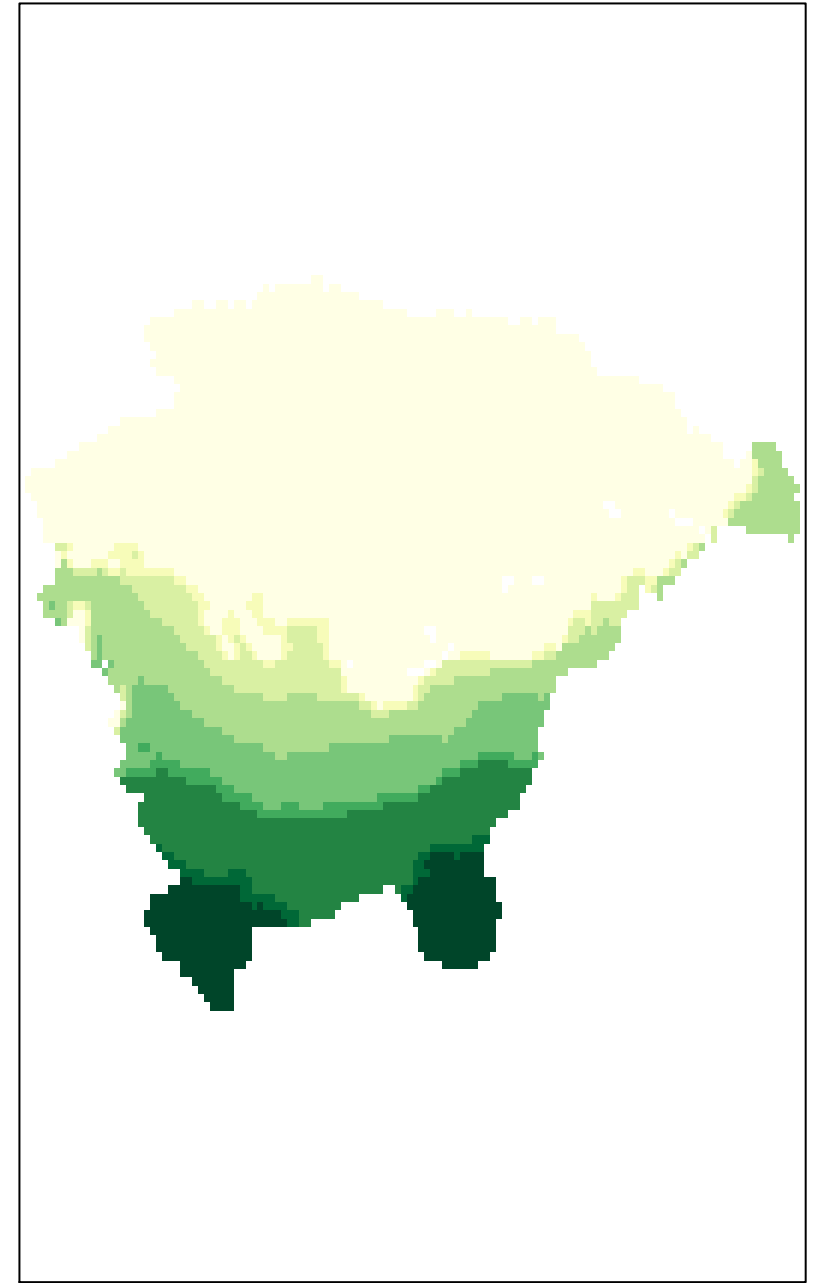


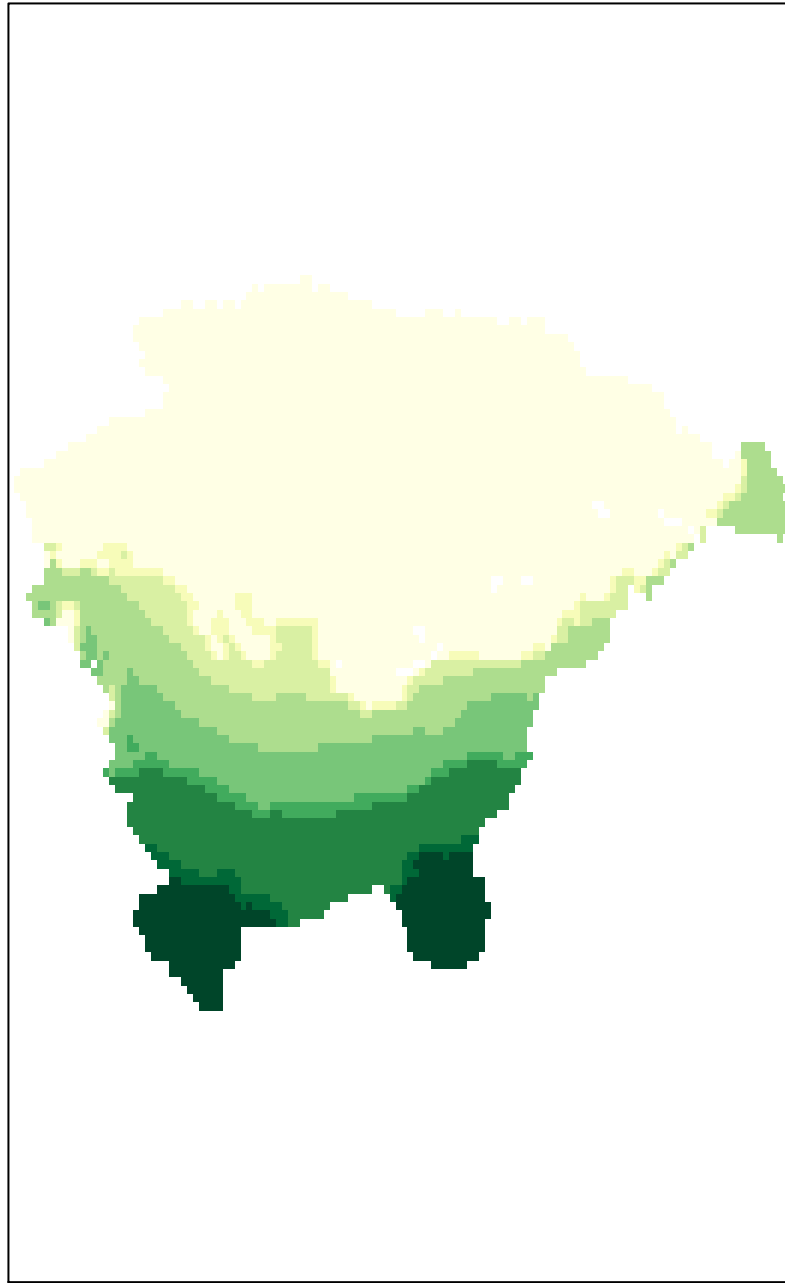
MAX, X21000.ybp



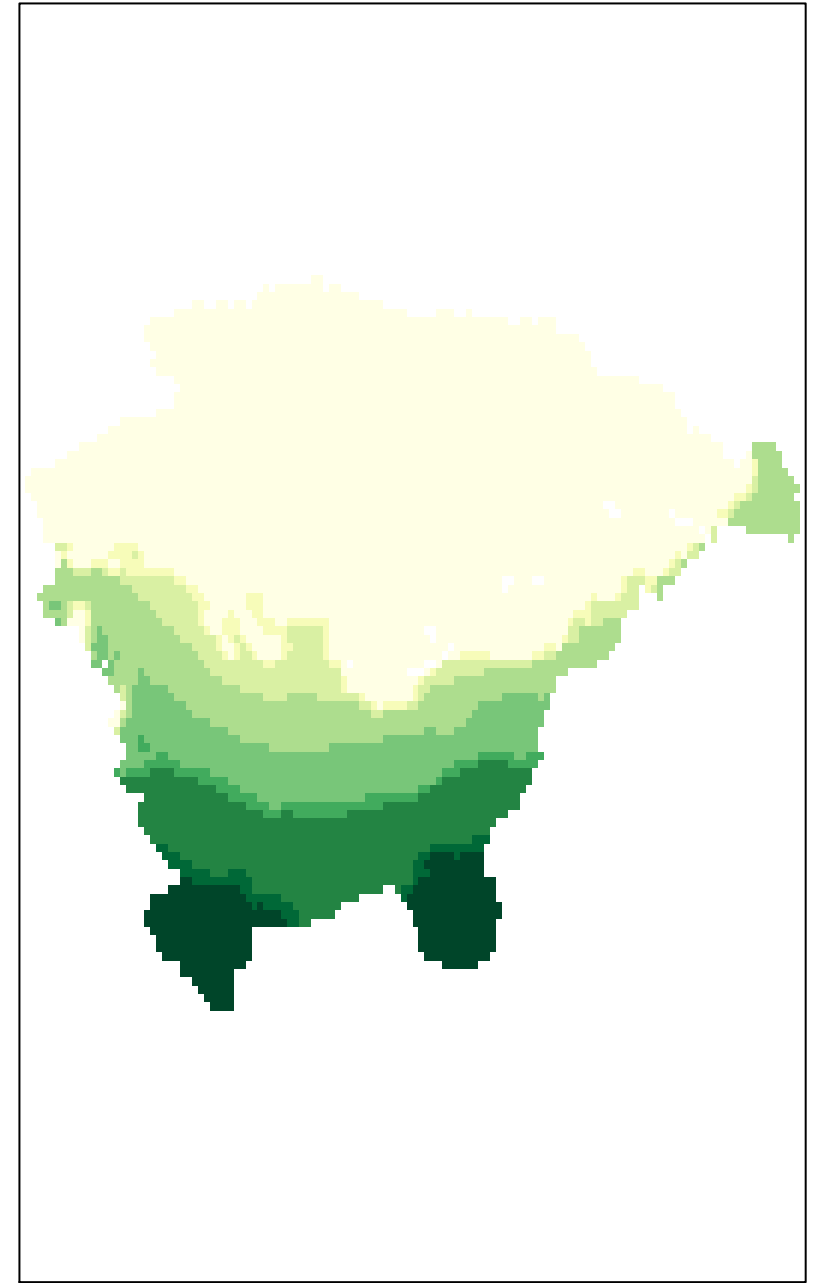
MAX, X21000.ybp



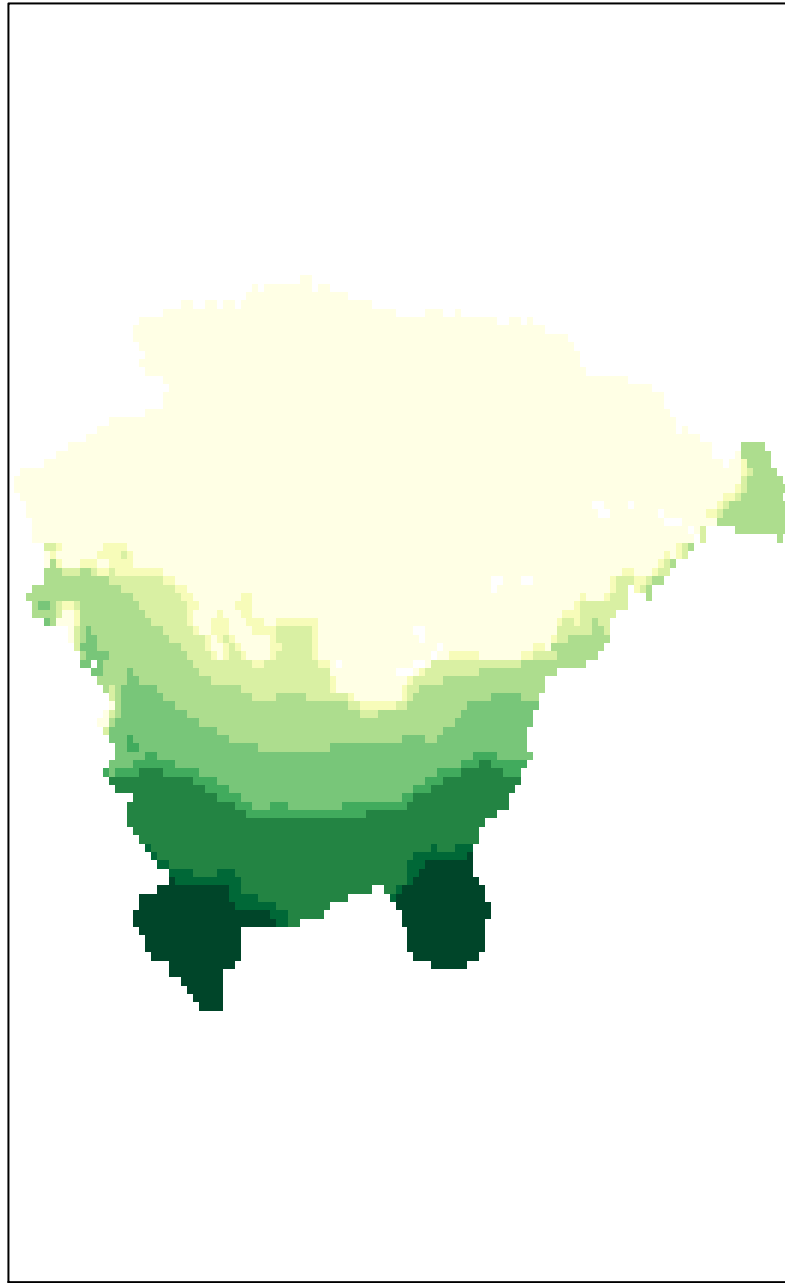
MAX, X20000.ybp



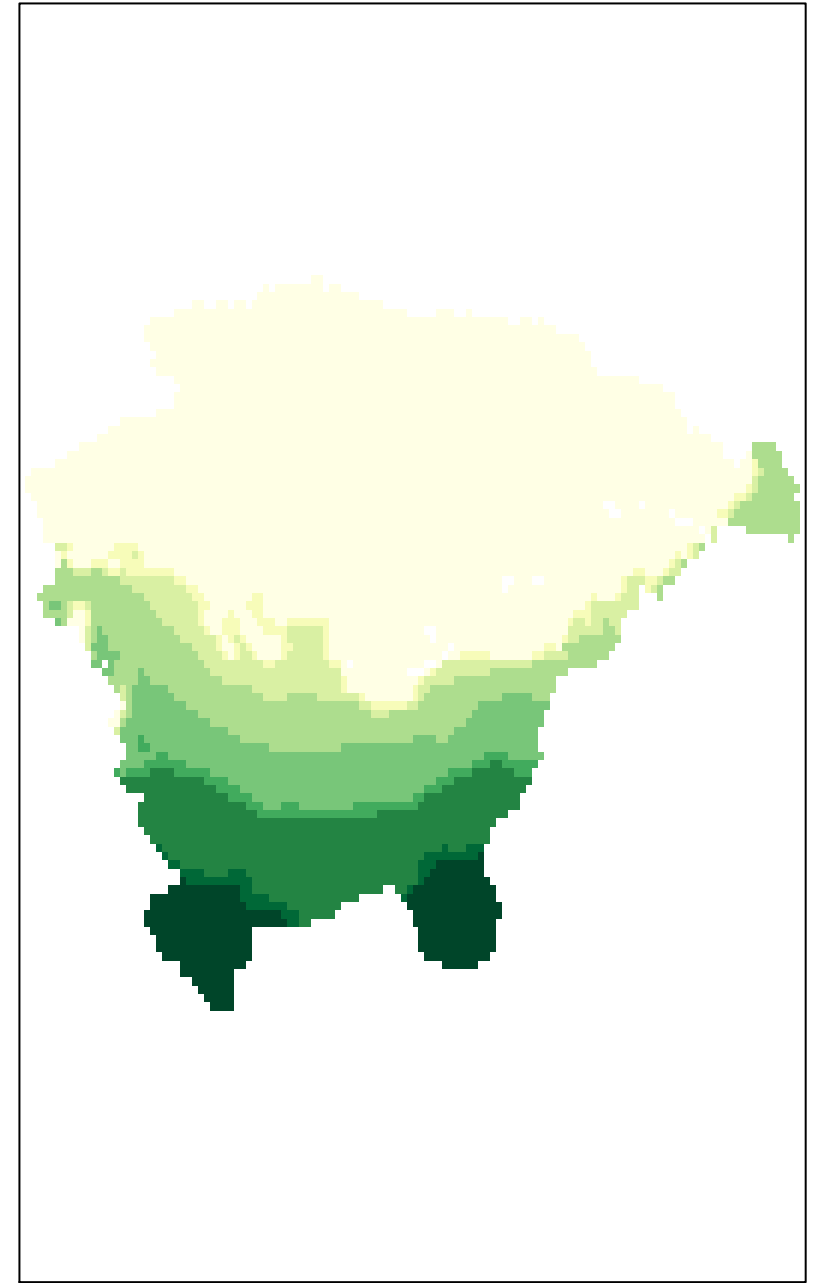
MAX, X20000.ybp



MAX, X19000.ybp

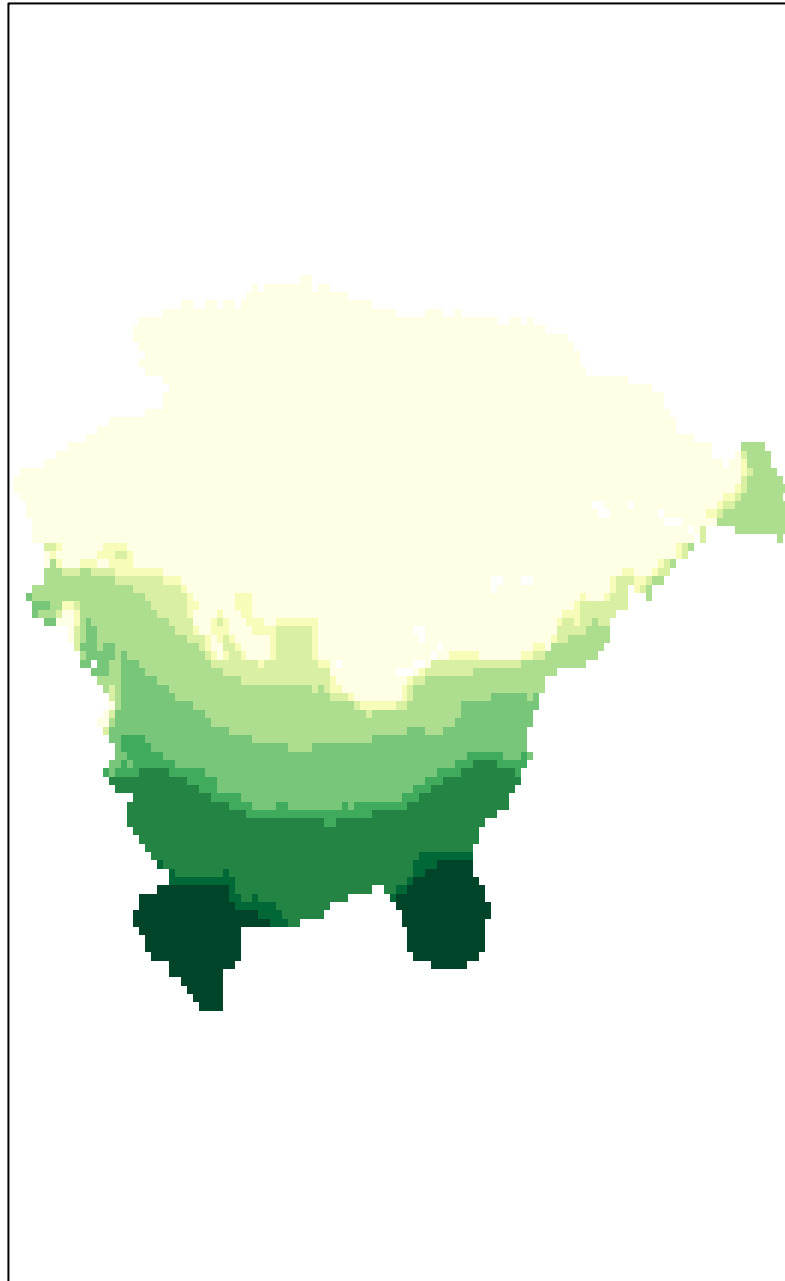


MAX, X19000.ybp

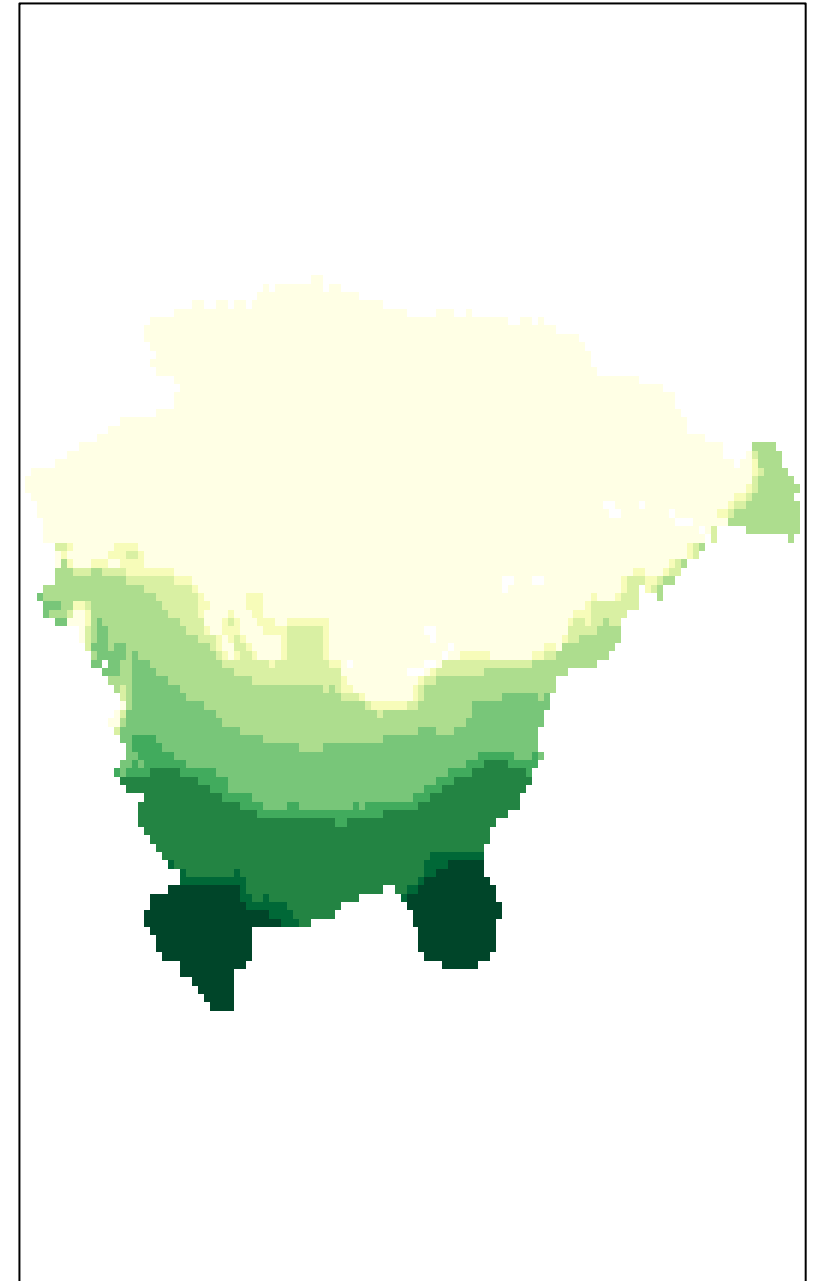


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

MAX, X18000.ybp

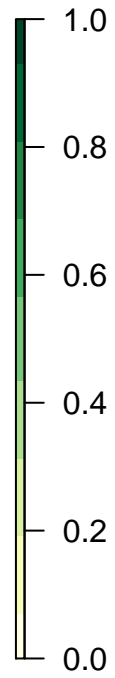
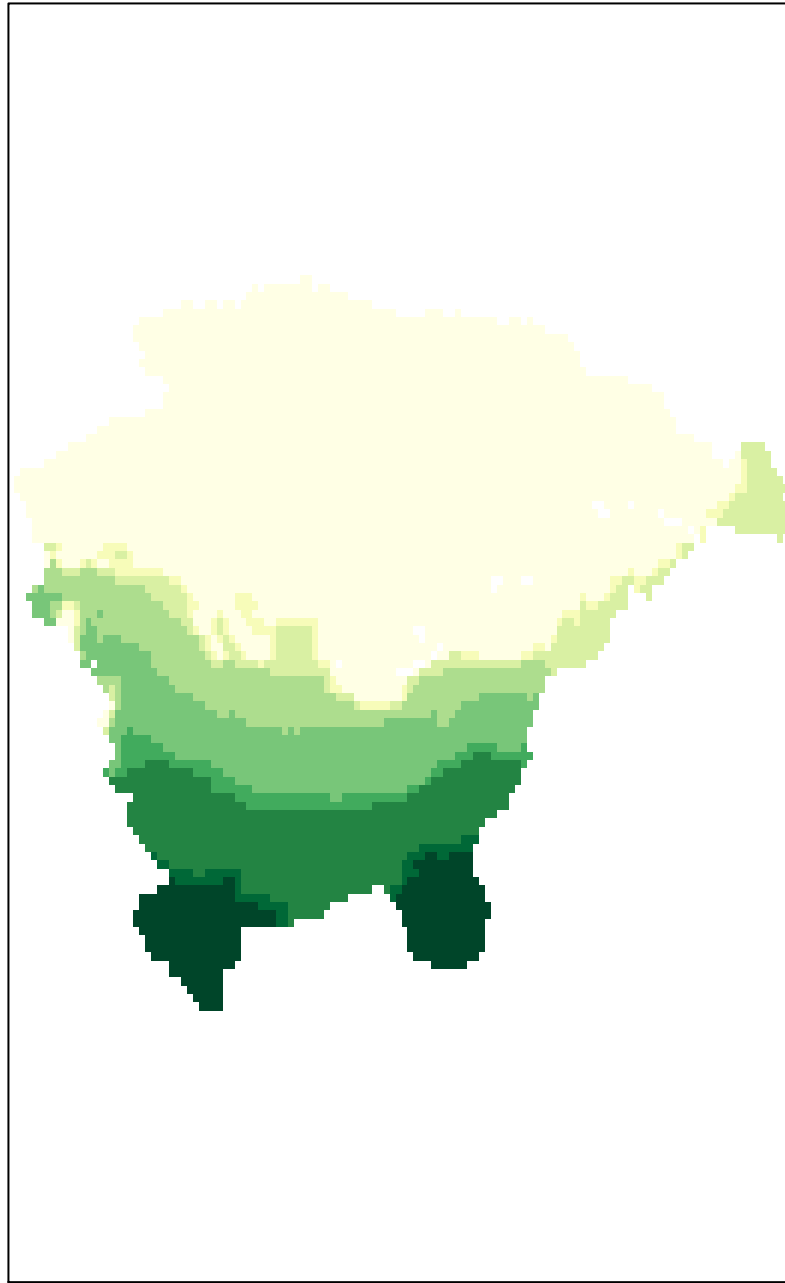


MAX, X18000.ybp

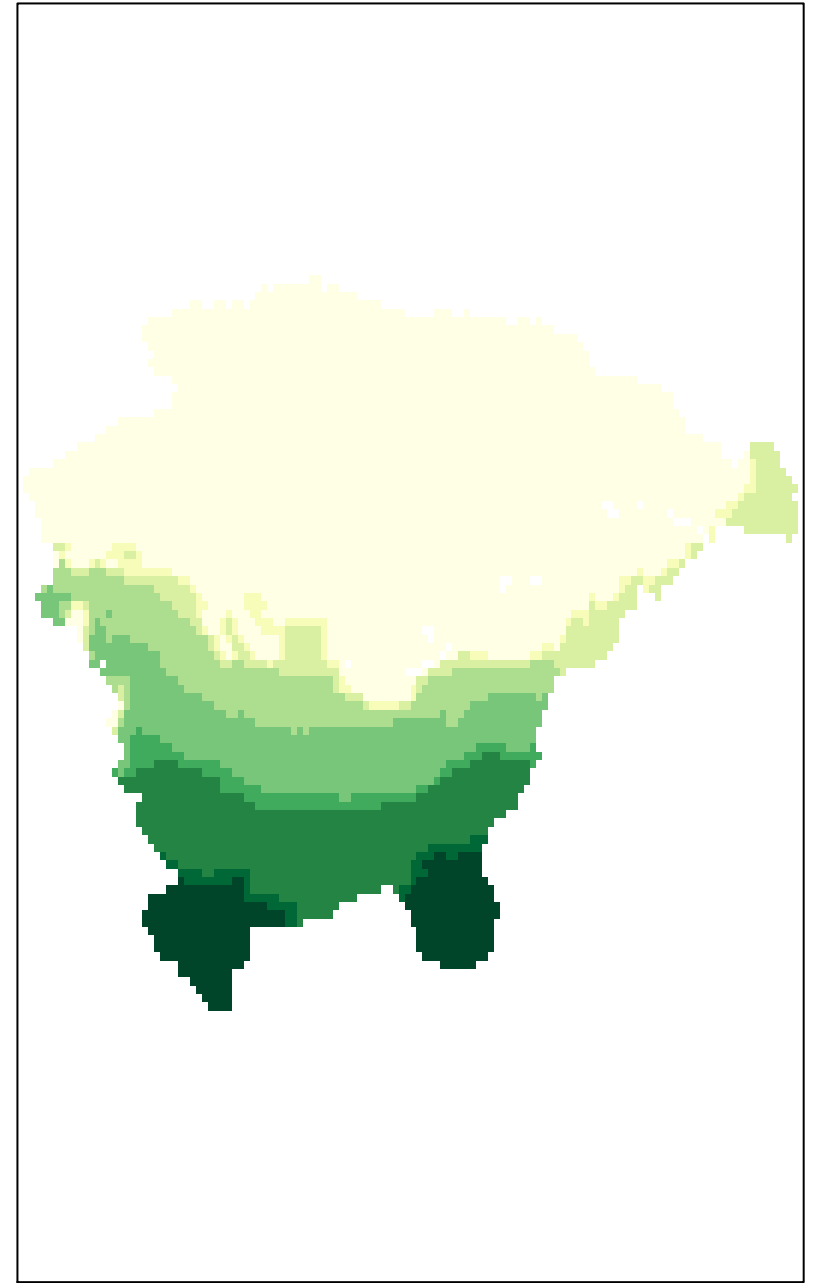


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

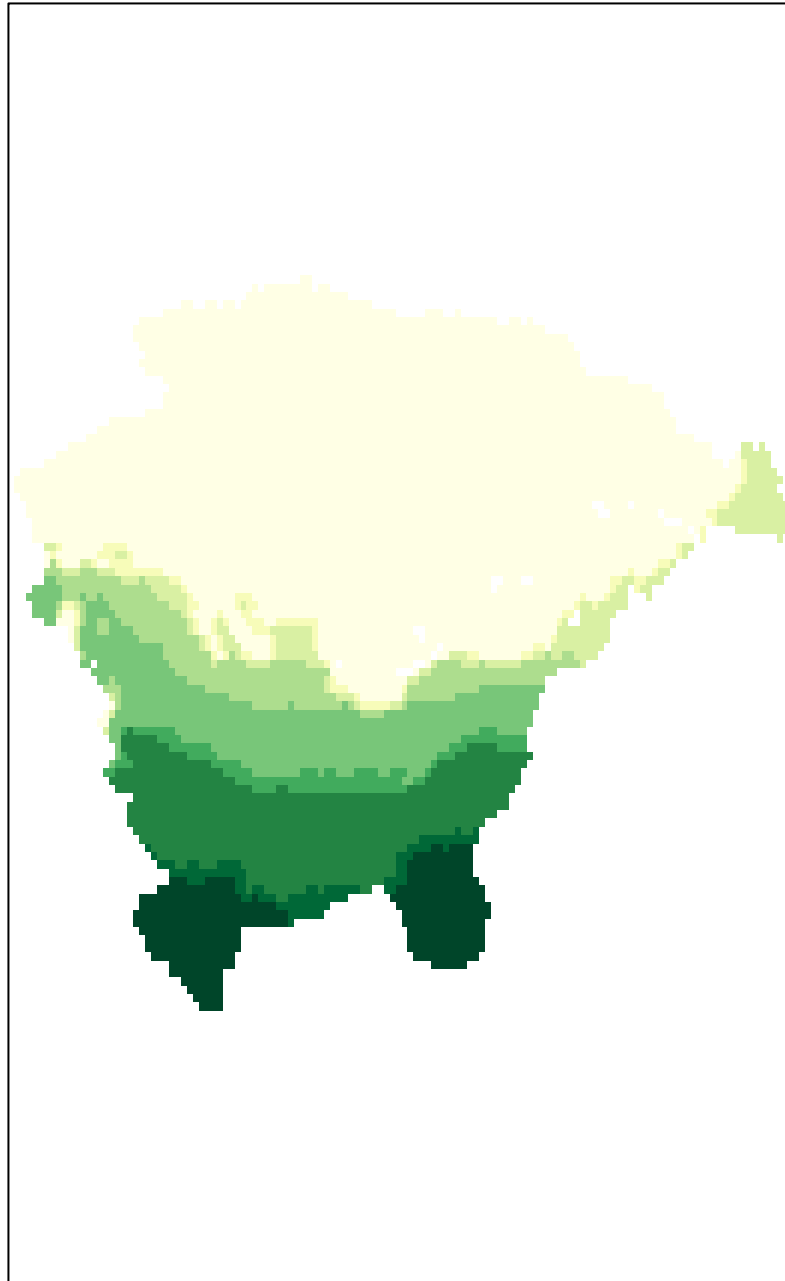
MAX, X17000.ybp



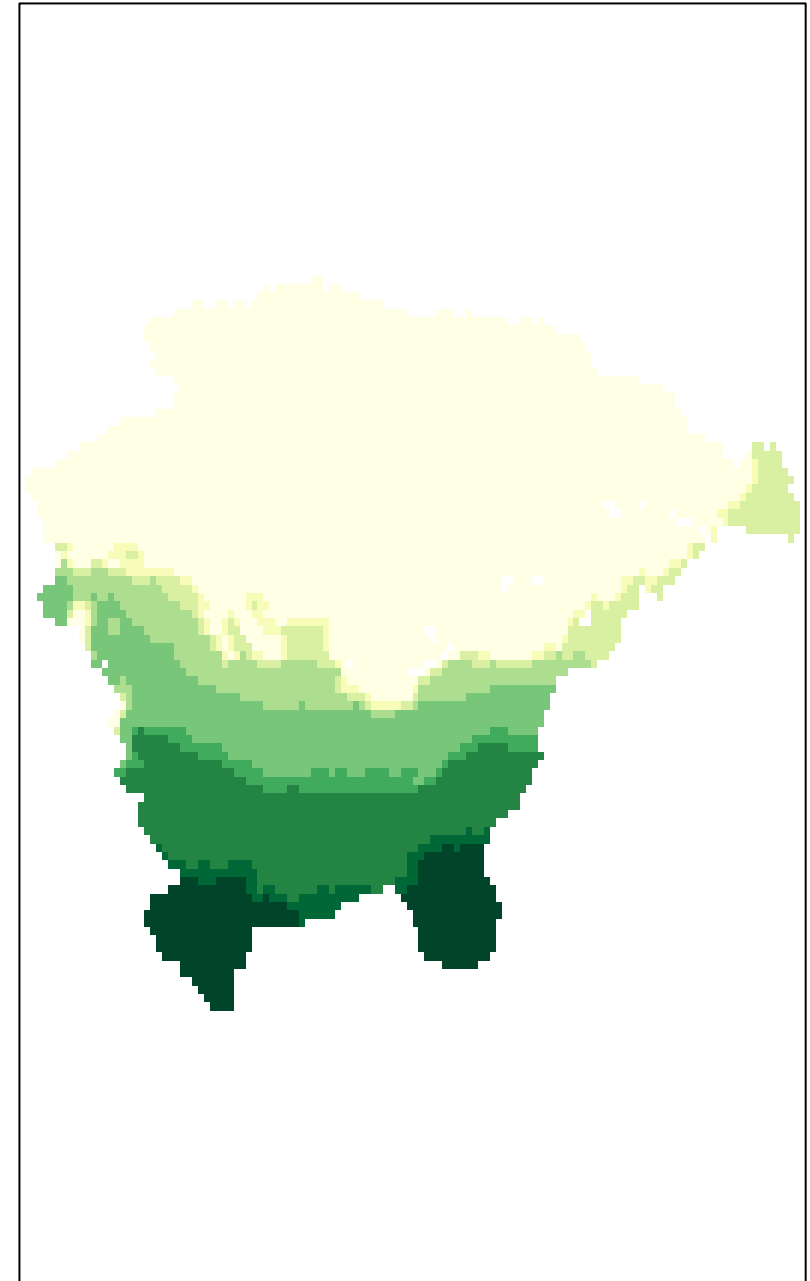
MAX, X17000.ybp



MAX, X16000.ybp

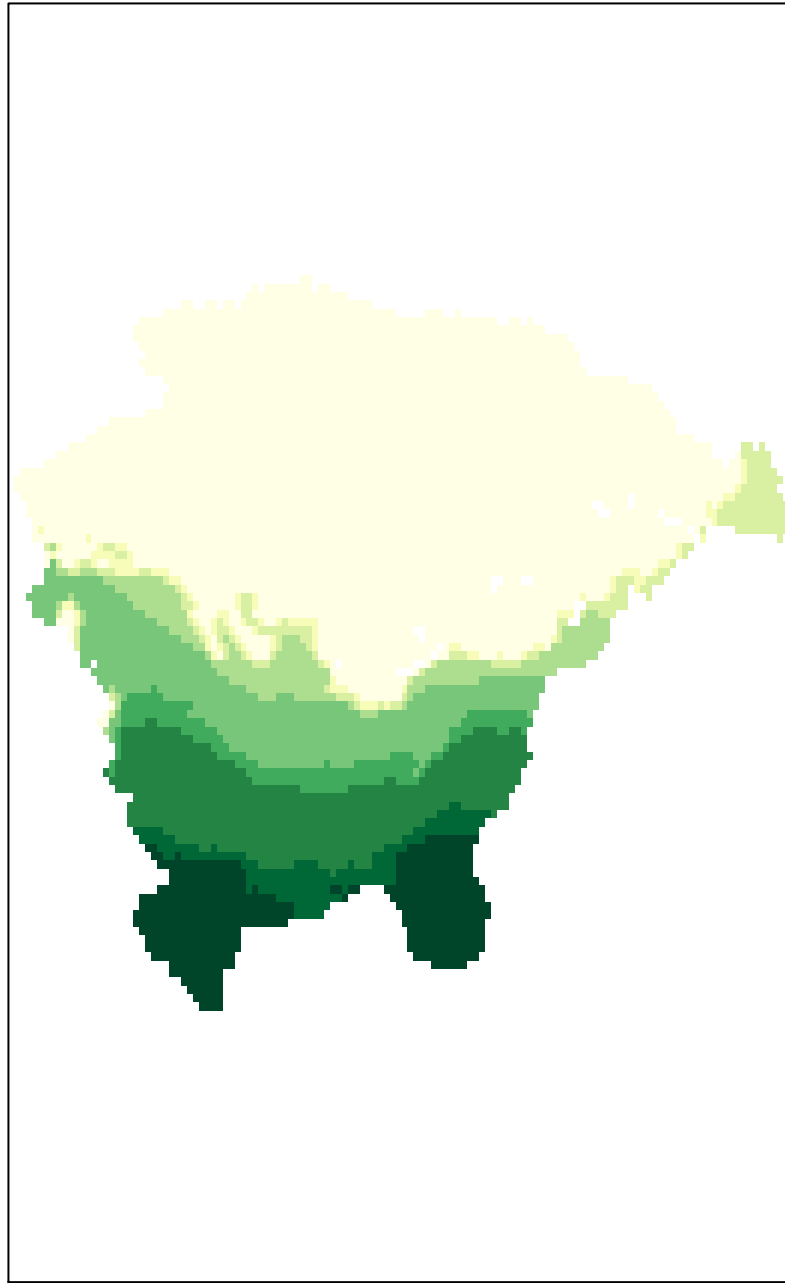


MAX, X16000.ybp

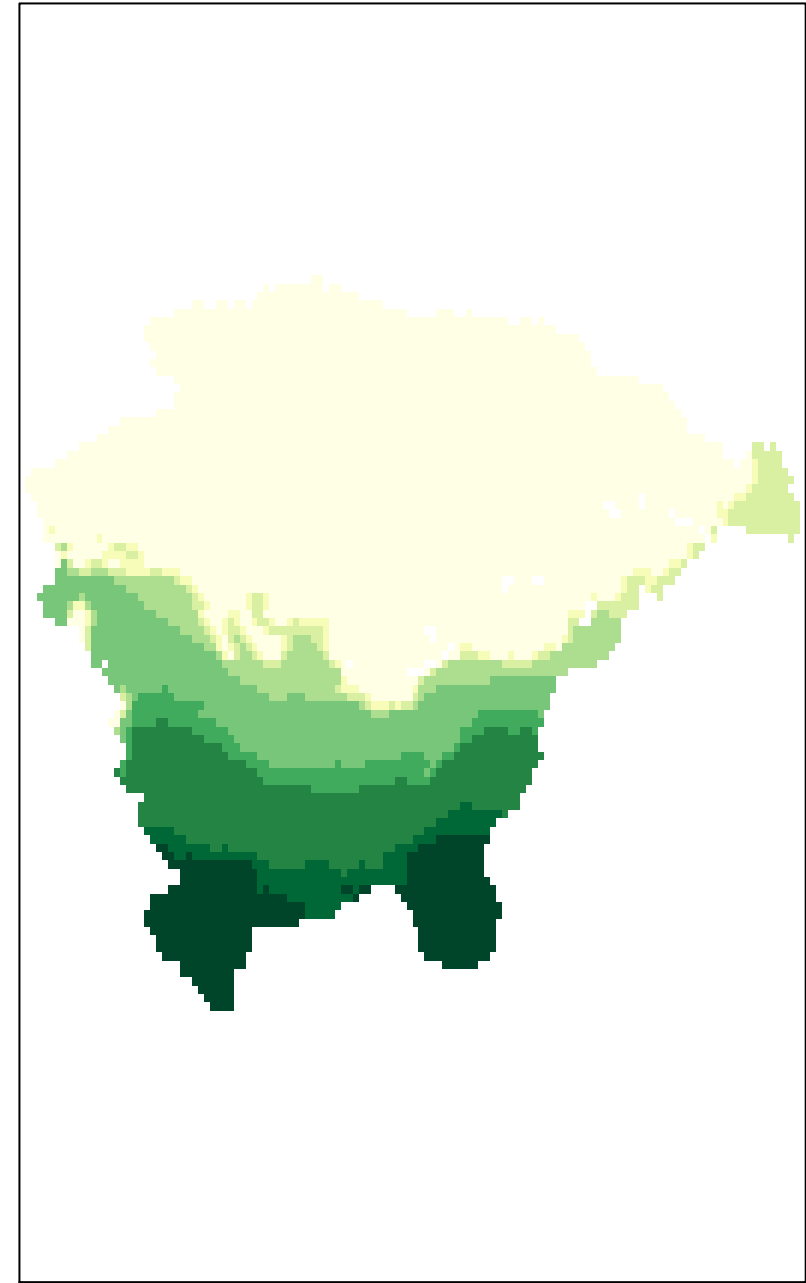


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

MAX, X15000.ybp

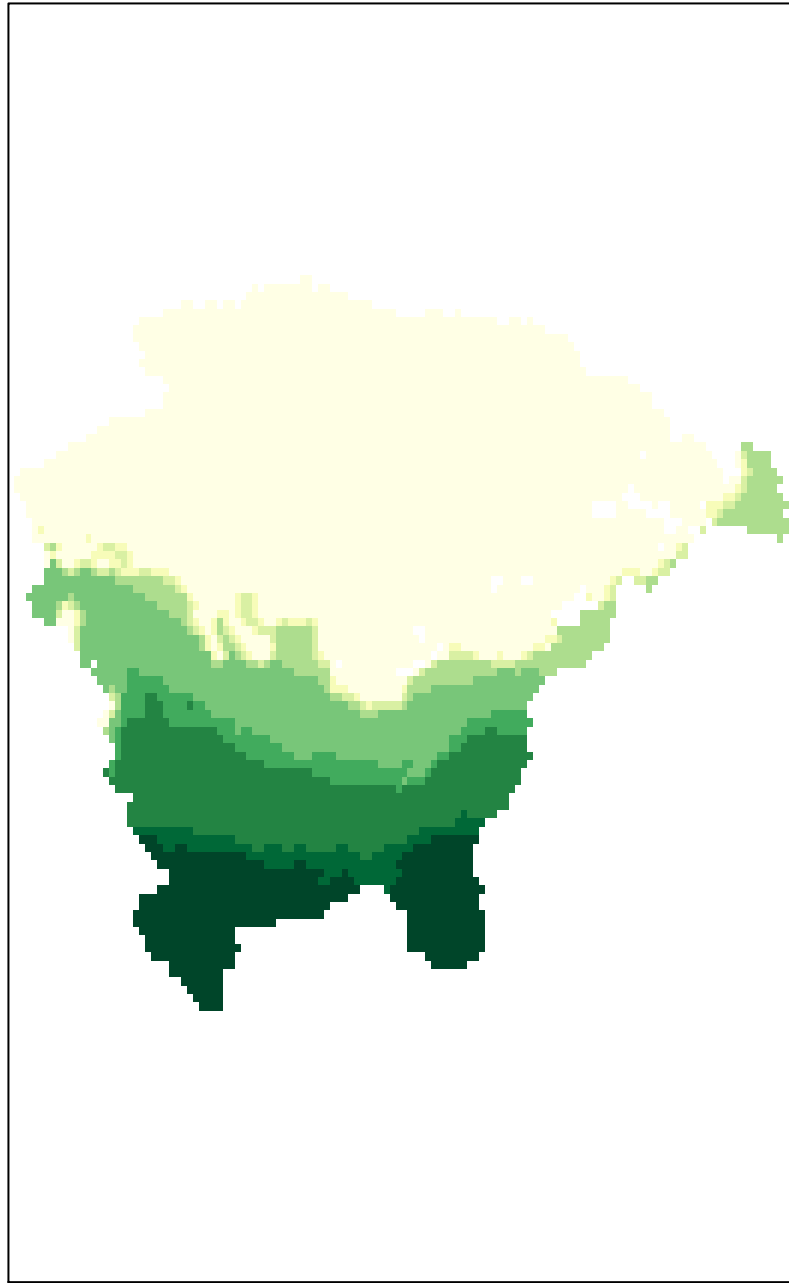


MAX, X15000.ybp

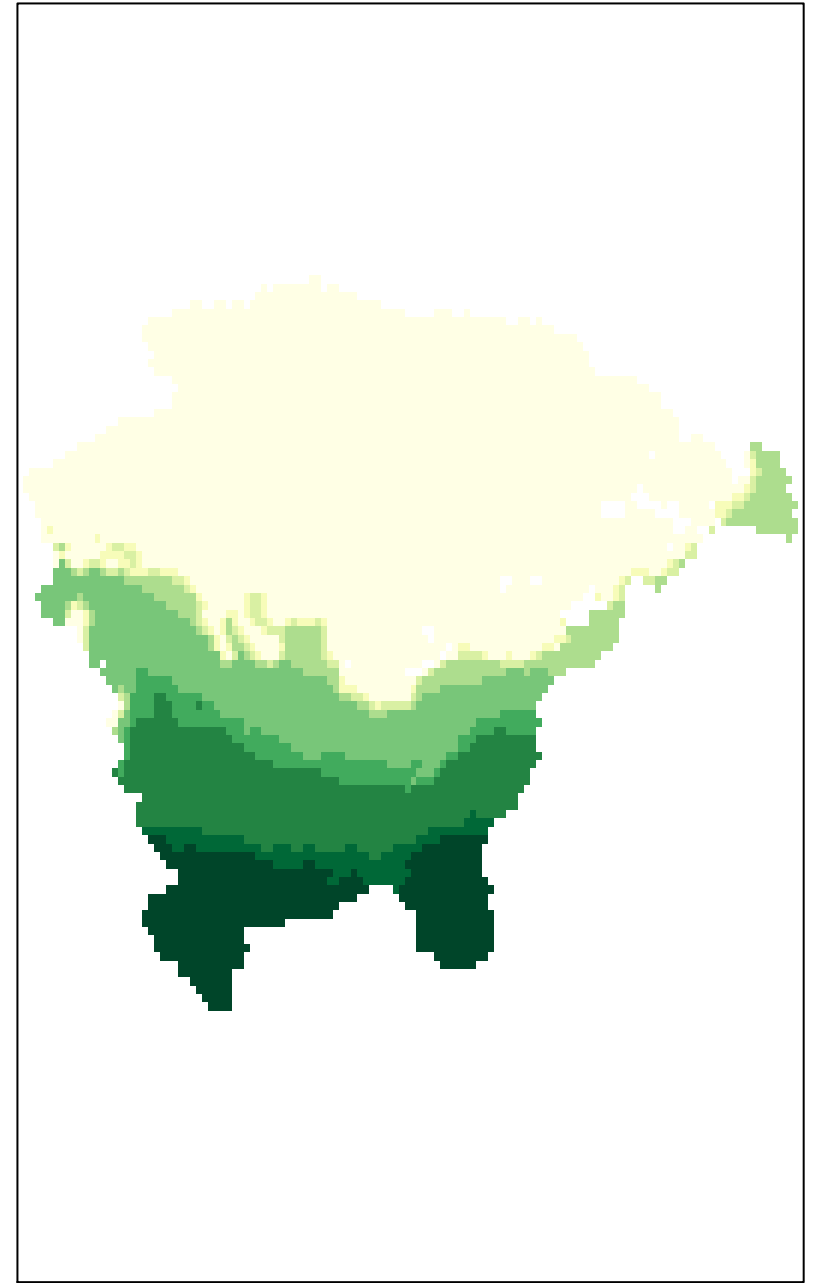


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

MAX, X14000.ybp

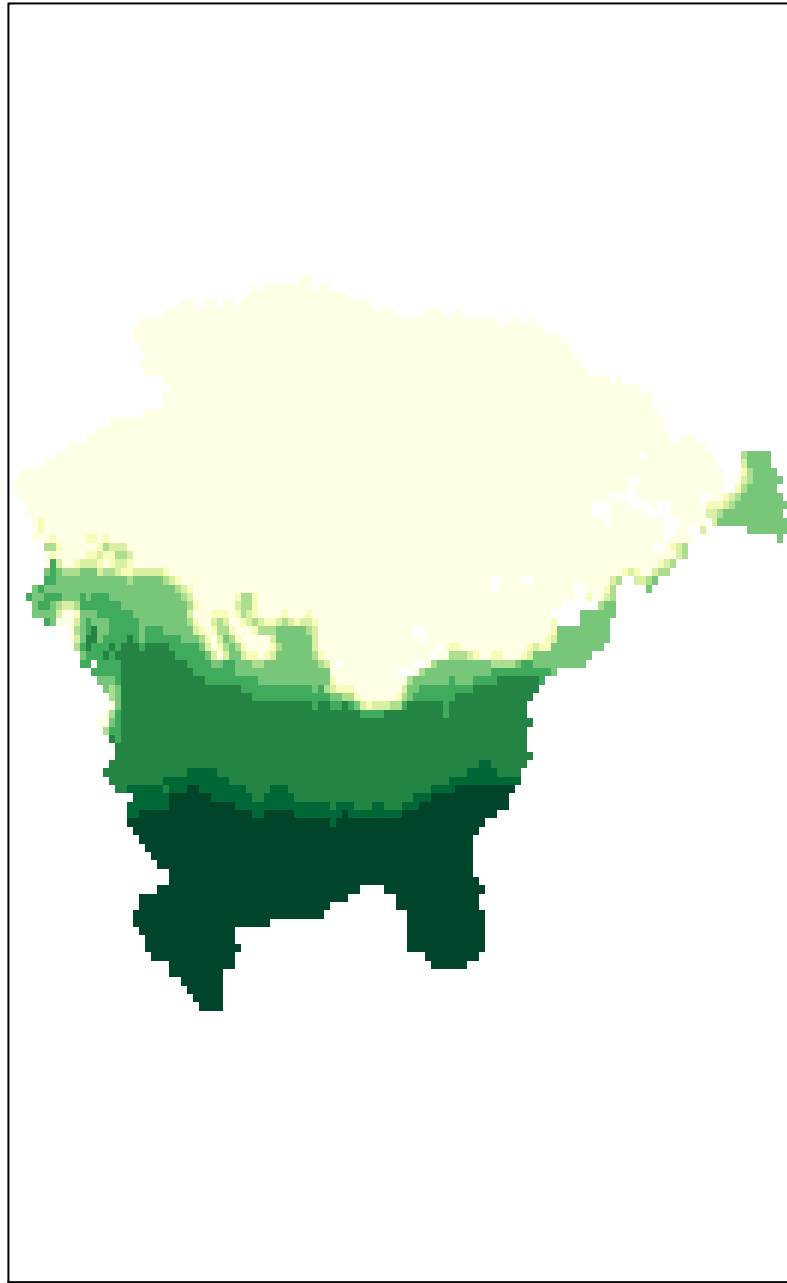


MAX, X14000.ybp

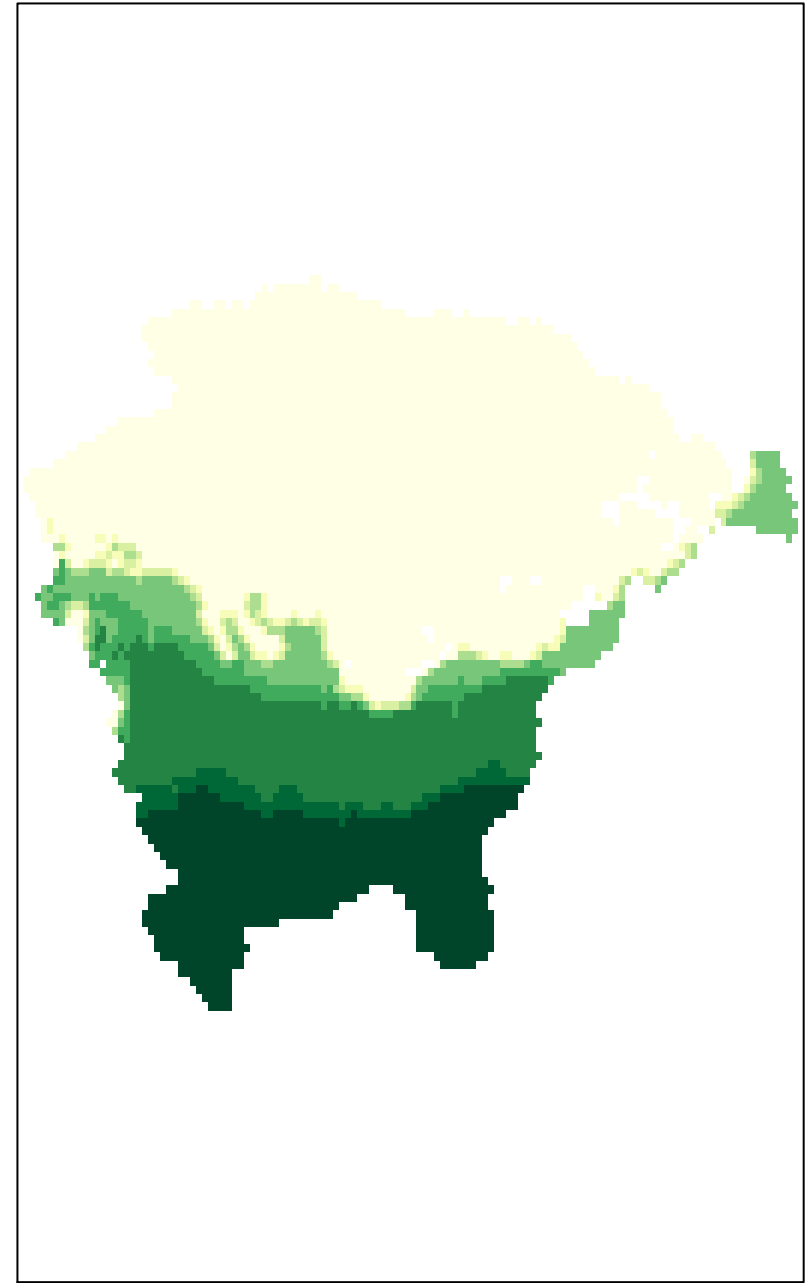


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

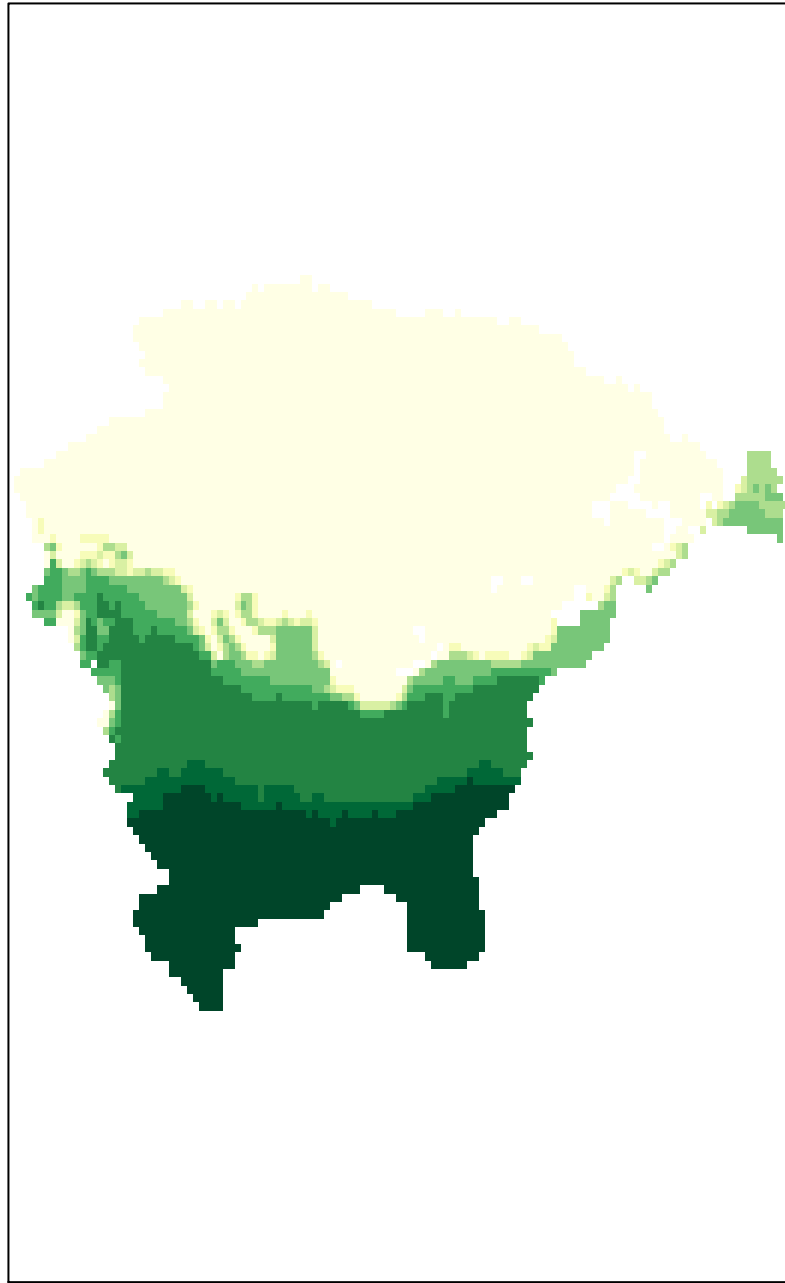
MAX, X13000.ybp



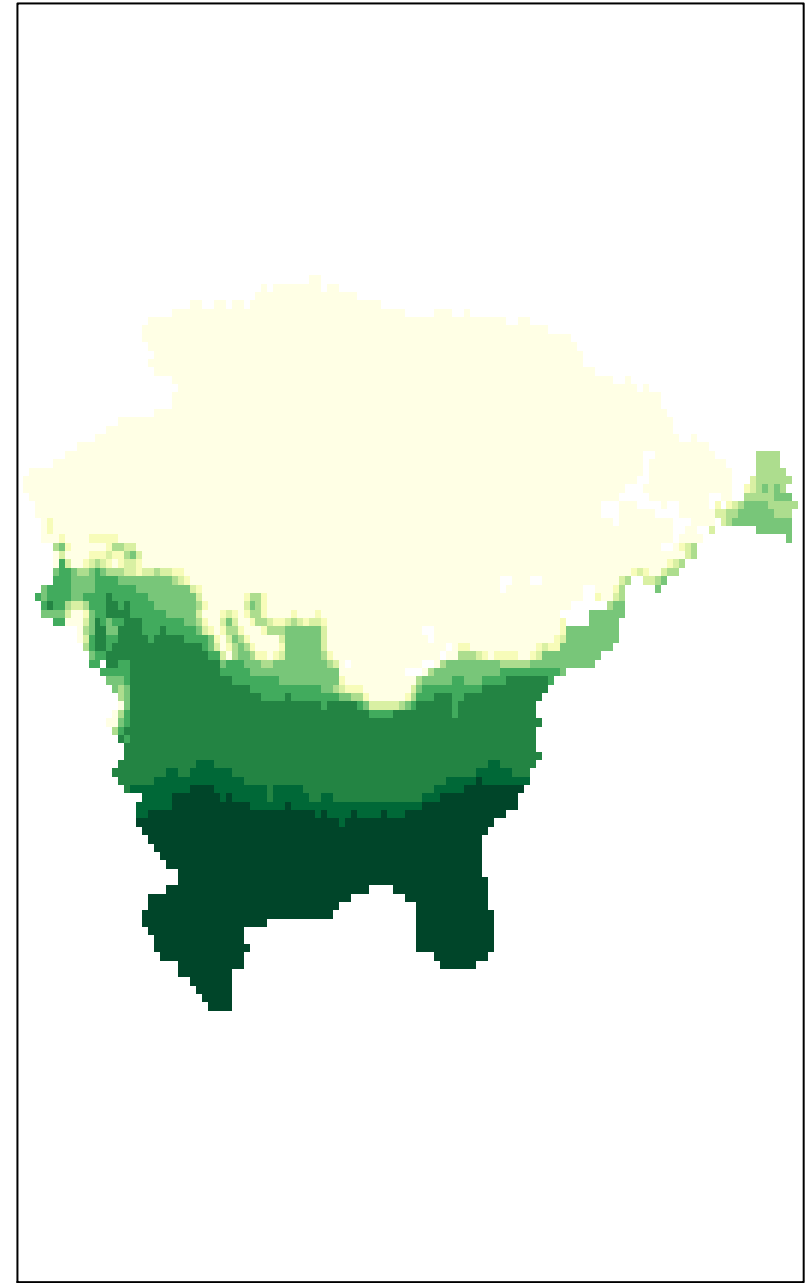
MAX, X13000.ybp



MAX, X12000.ybp

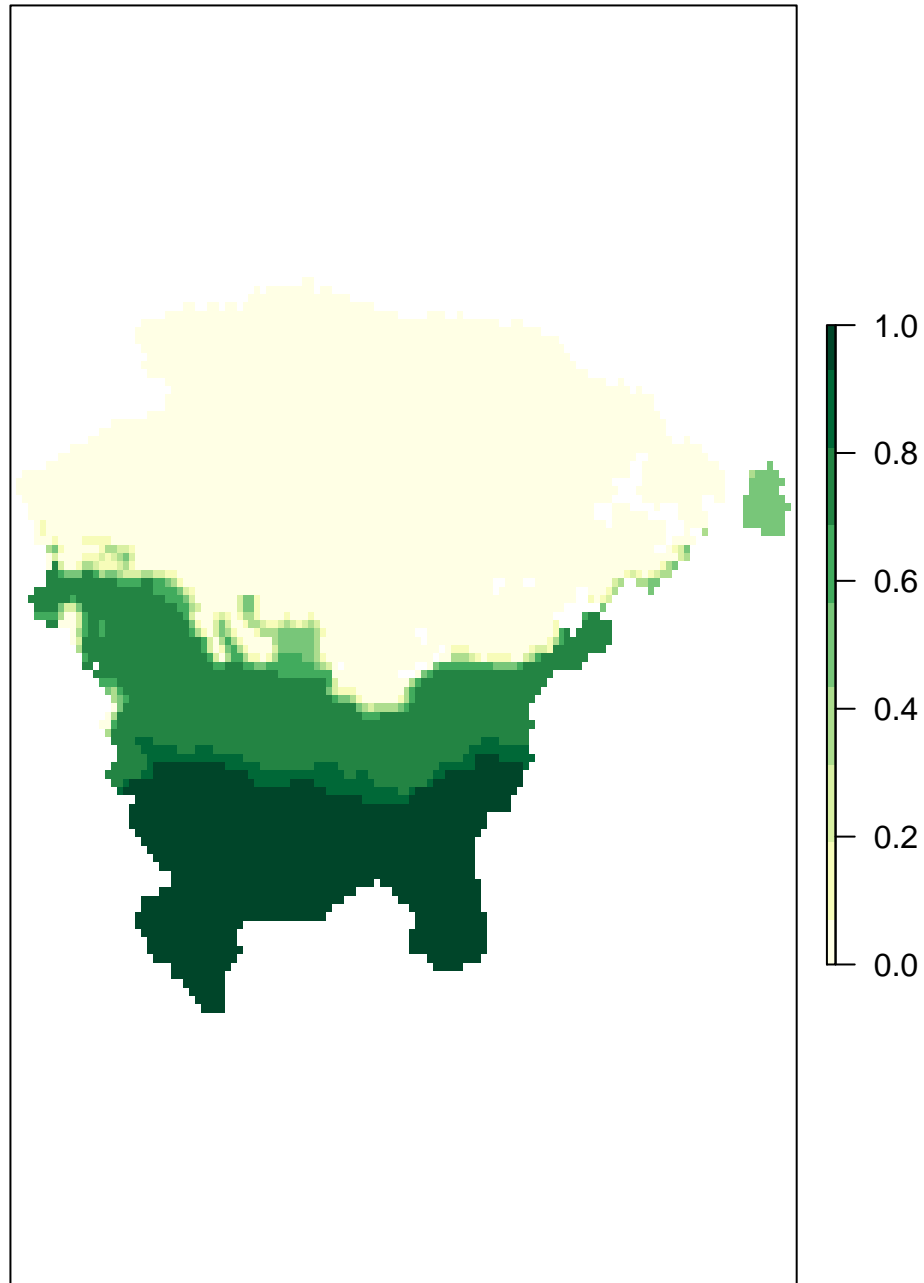


MAX, X12000.ybp

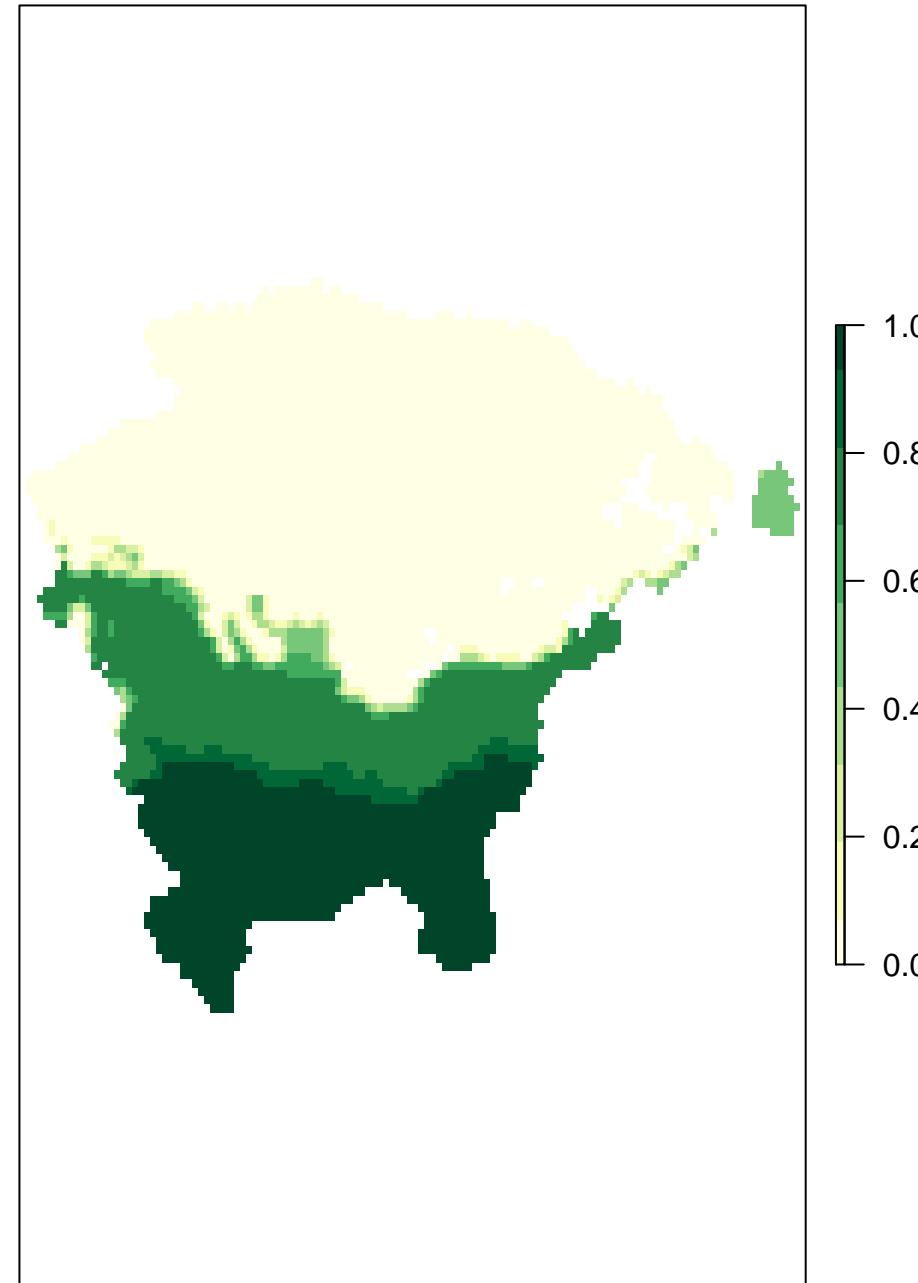


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

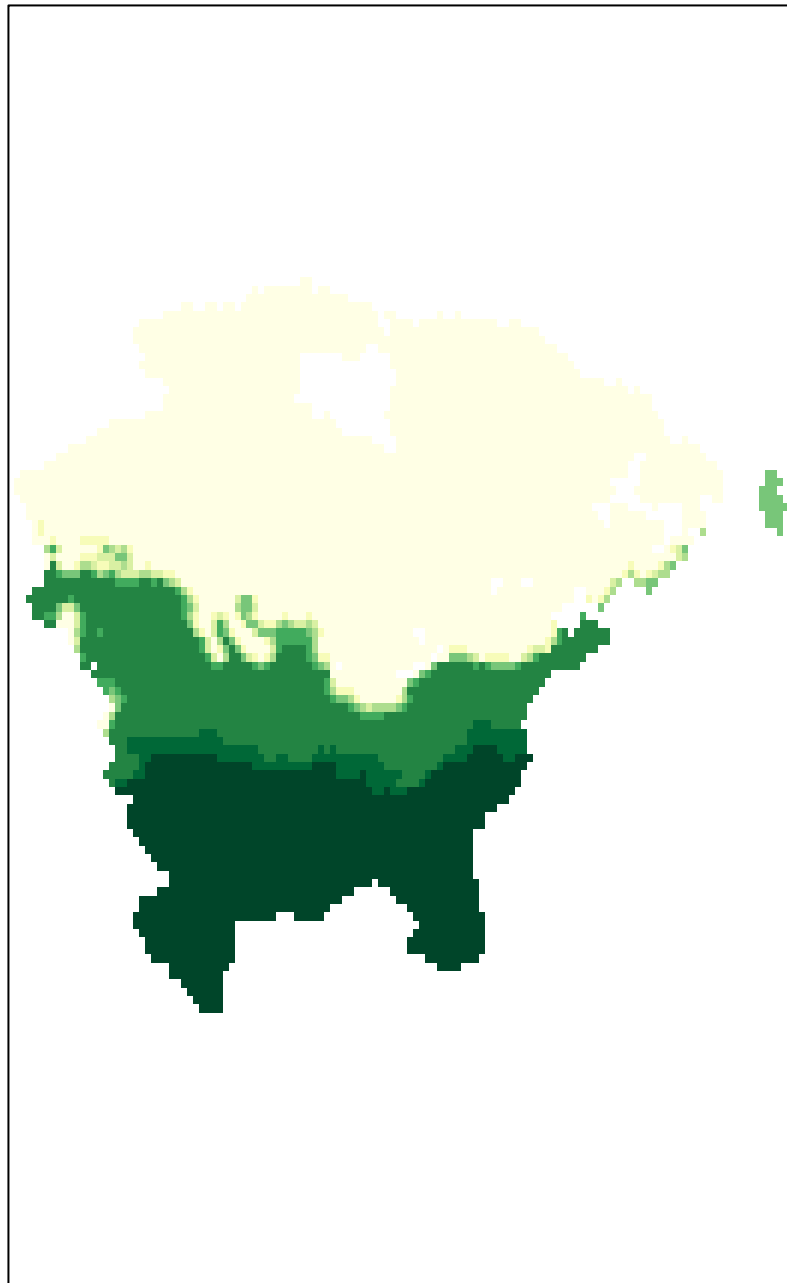
MAX, X11000.ybp



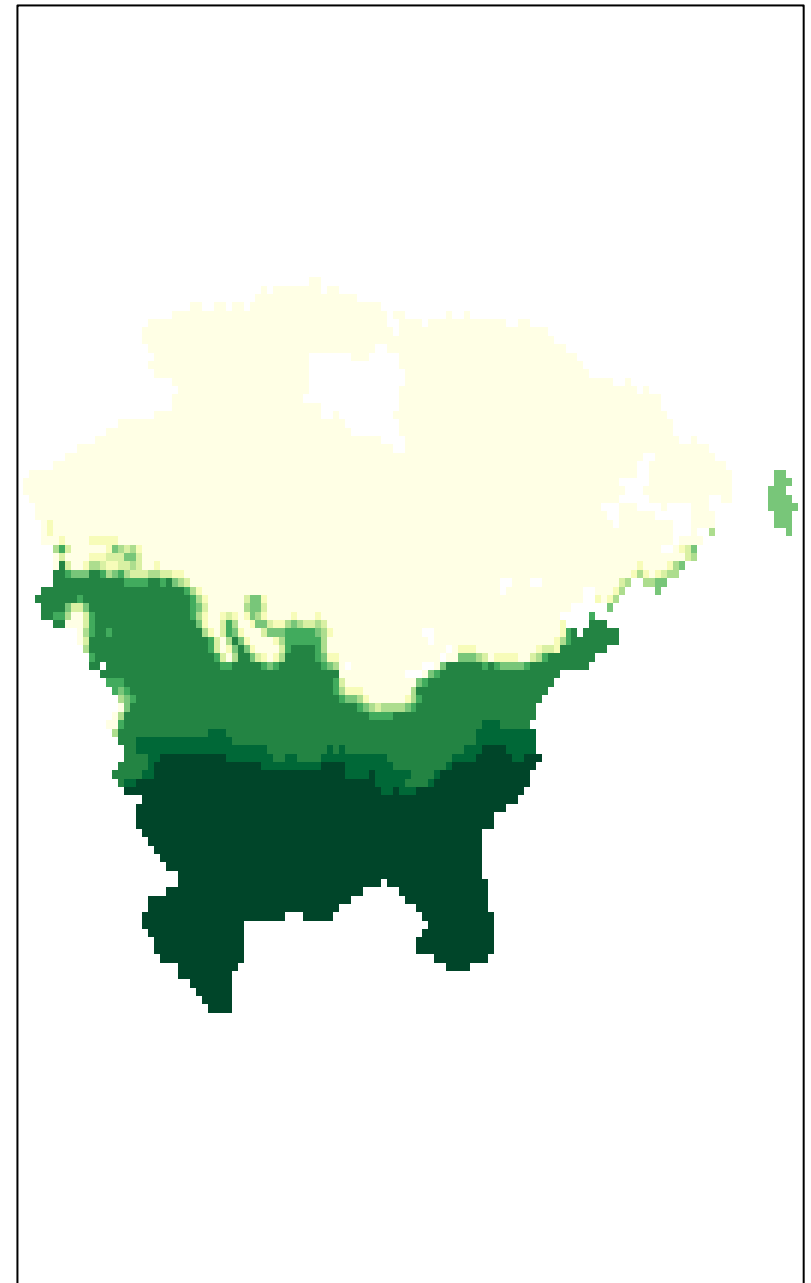
MAX, X11000.ybp



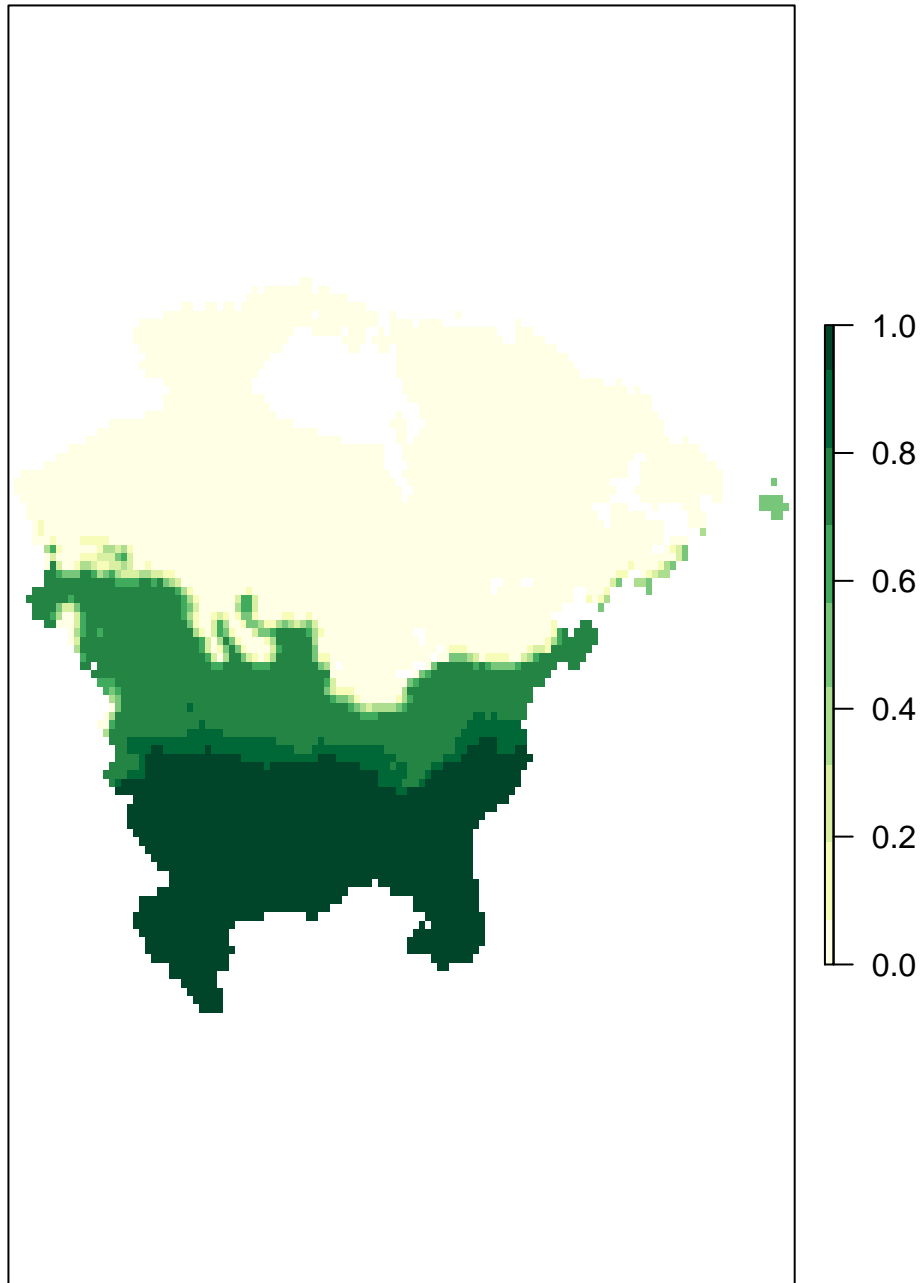
MAX, X10000.ybp



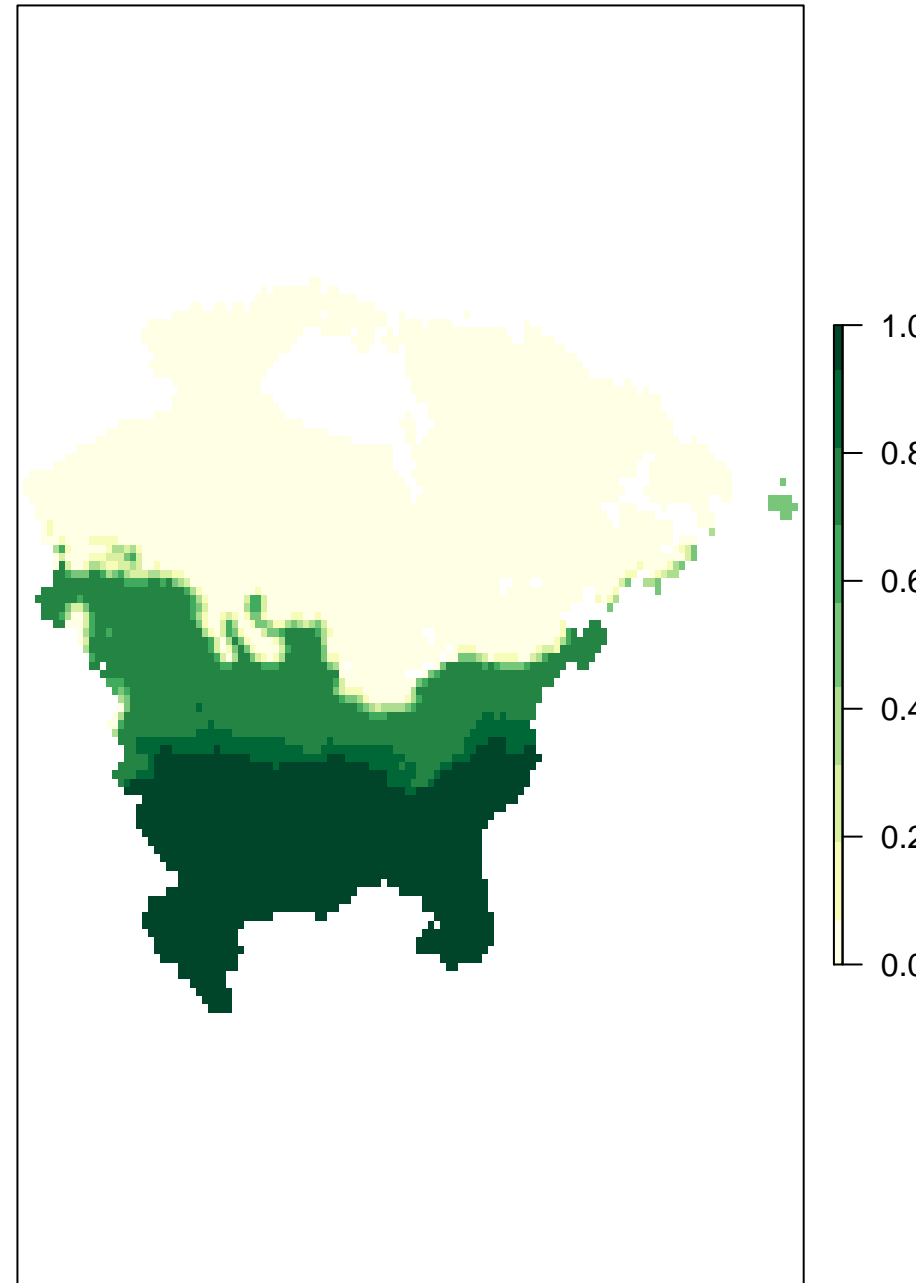
MAX, X10000.ybp



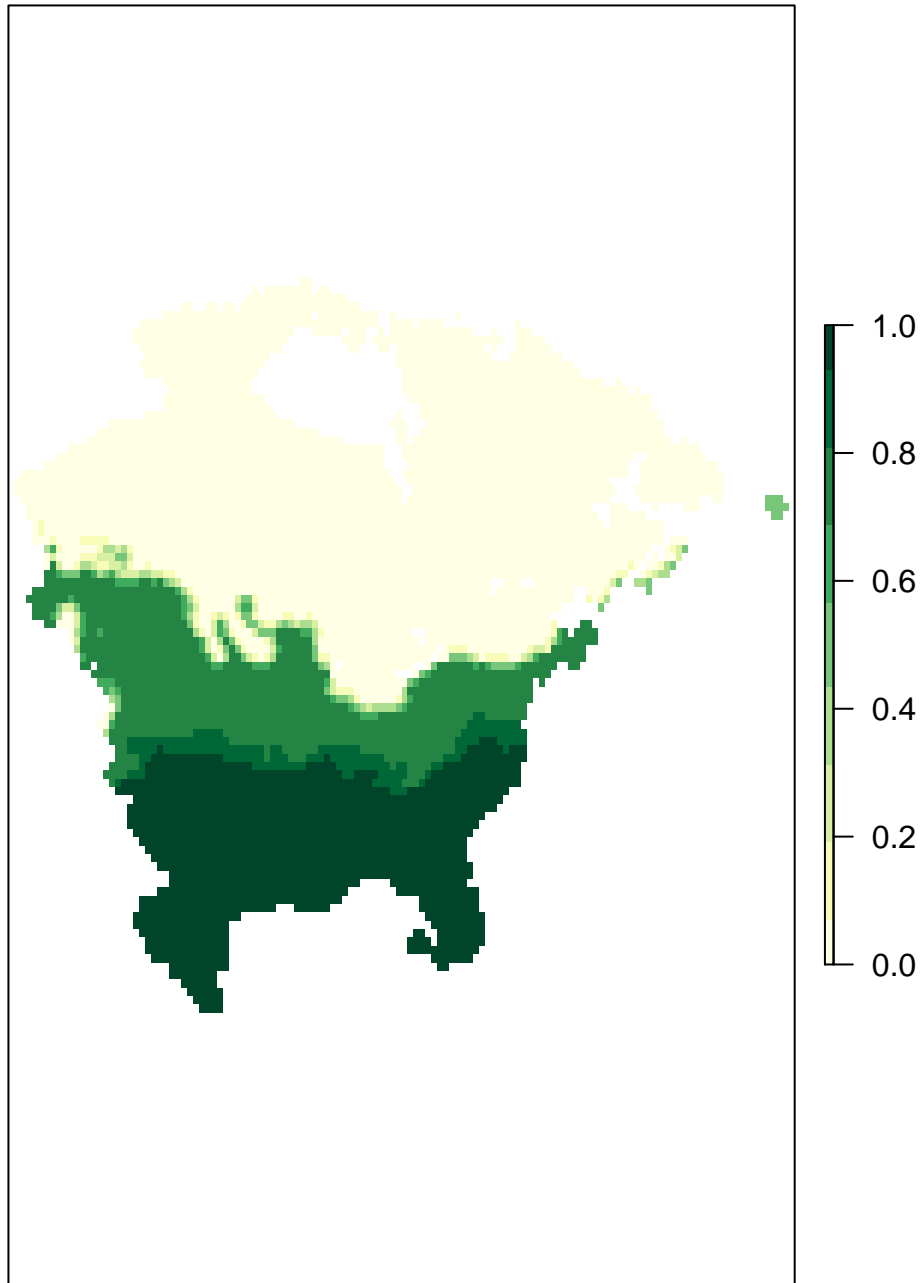
MAX, X9000.ybp



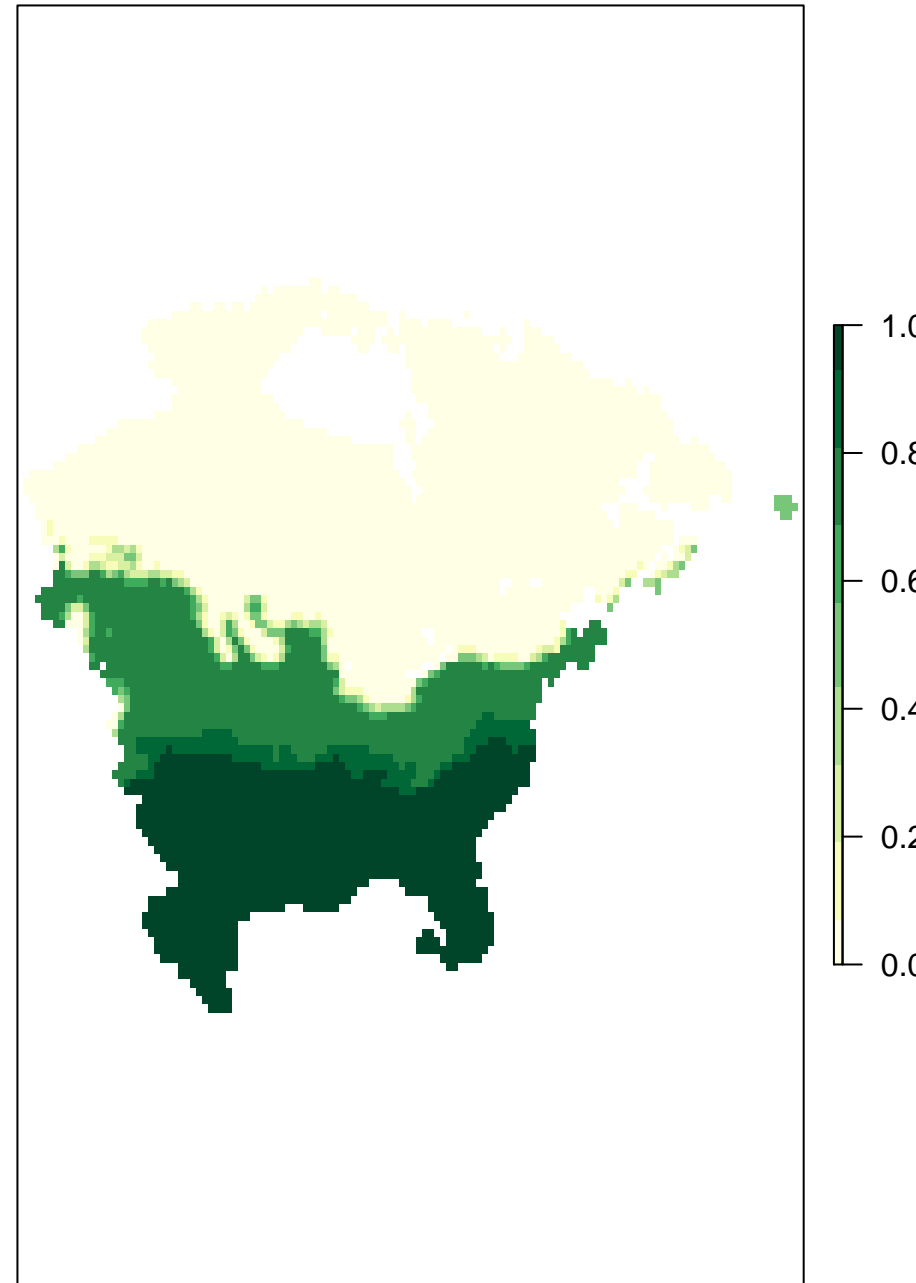
MAX, X9000.ybp



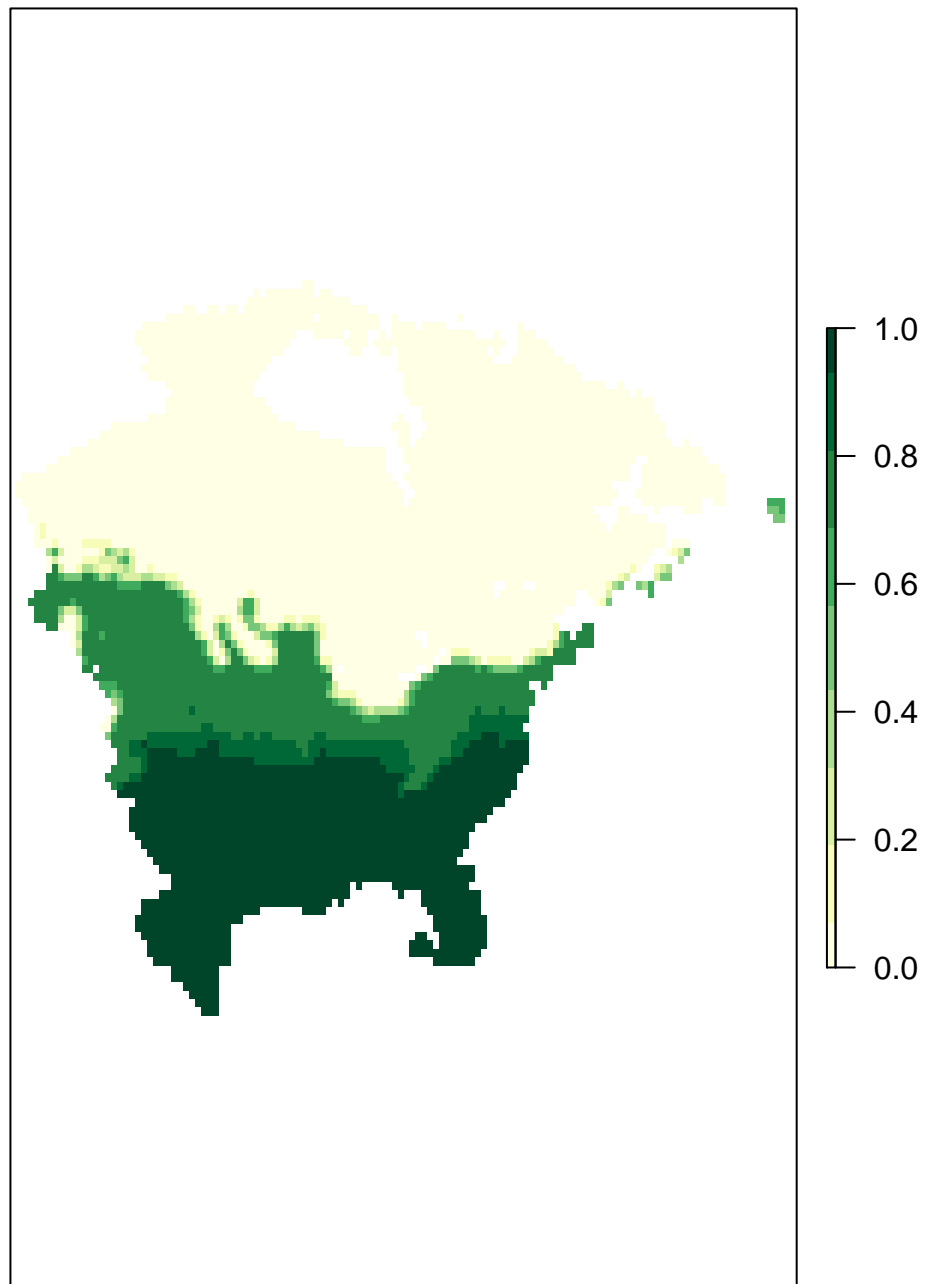
MAX, X8000.ybp



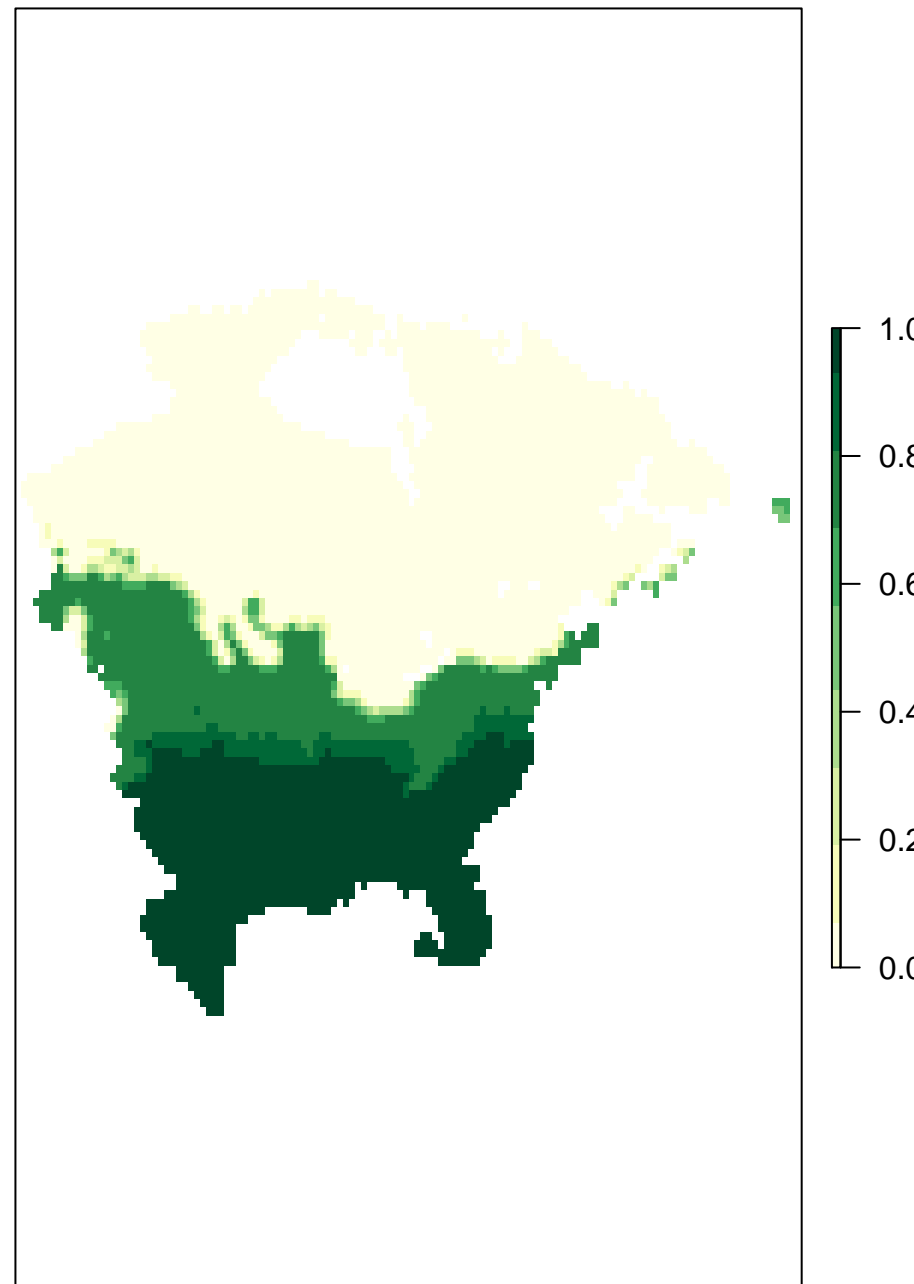
MAX, X8000.ybp



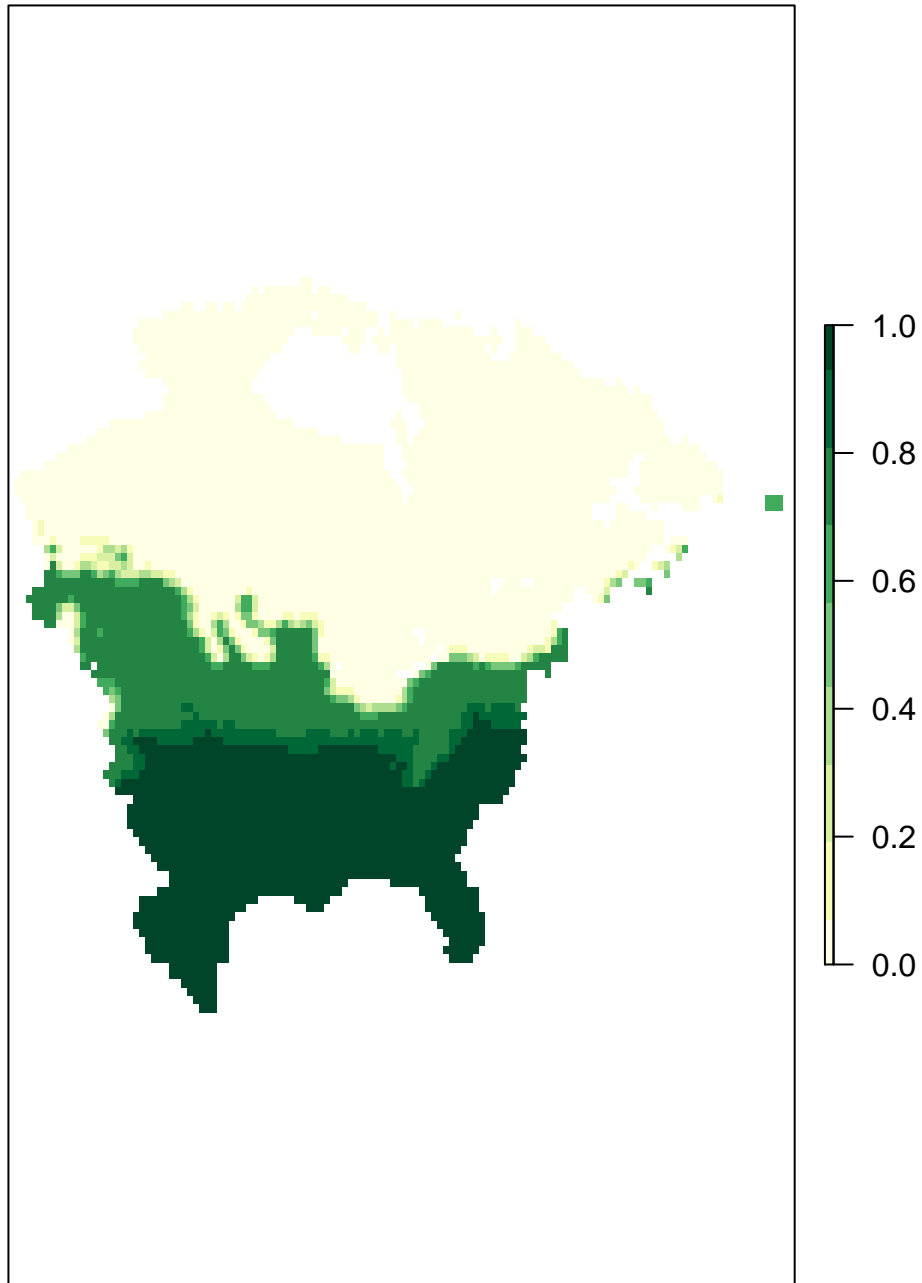
MAX, X7000.ybp



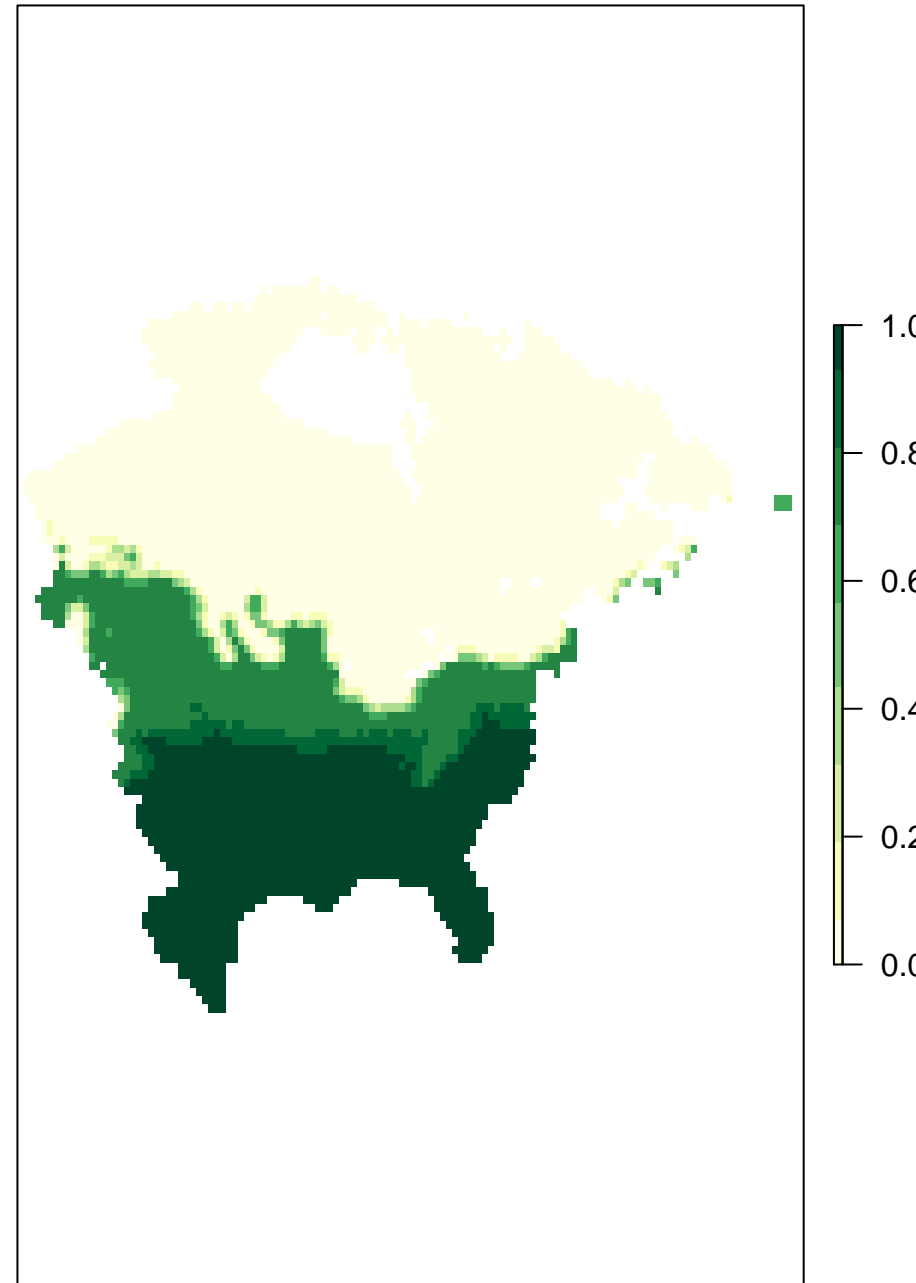
MAX, X7000.ybp



MAX, X6000.ybp

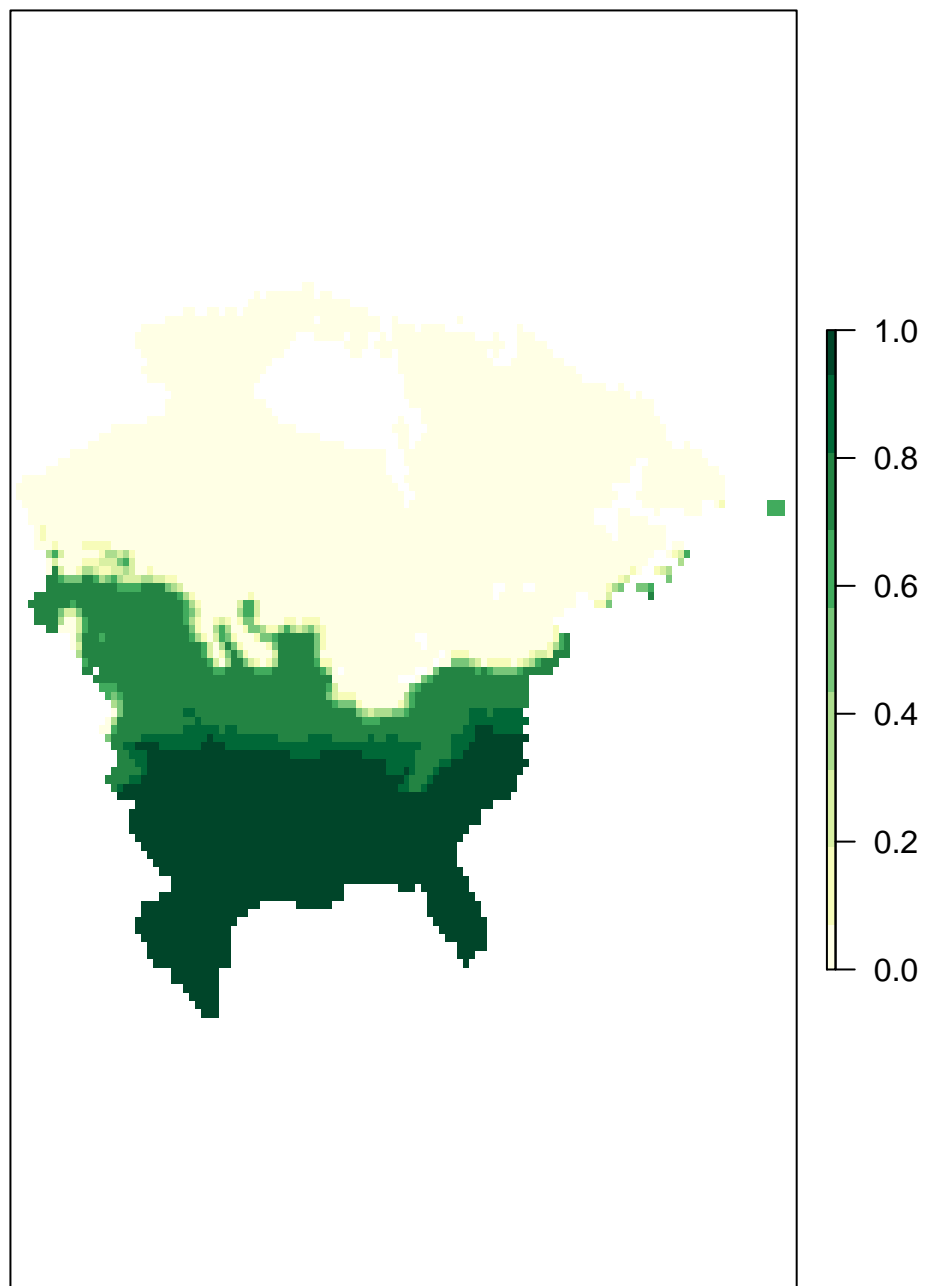


MAX, X6000.ybp

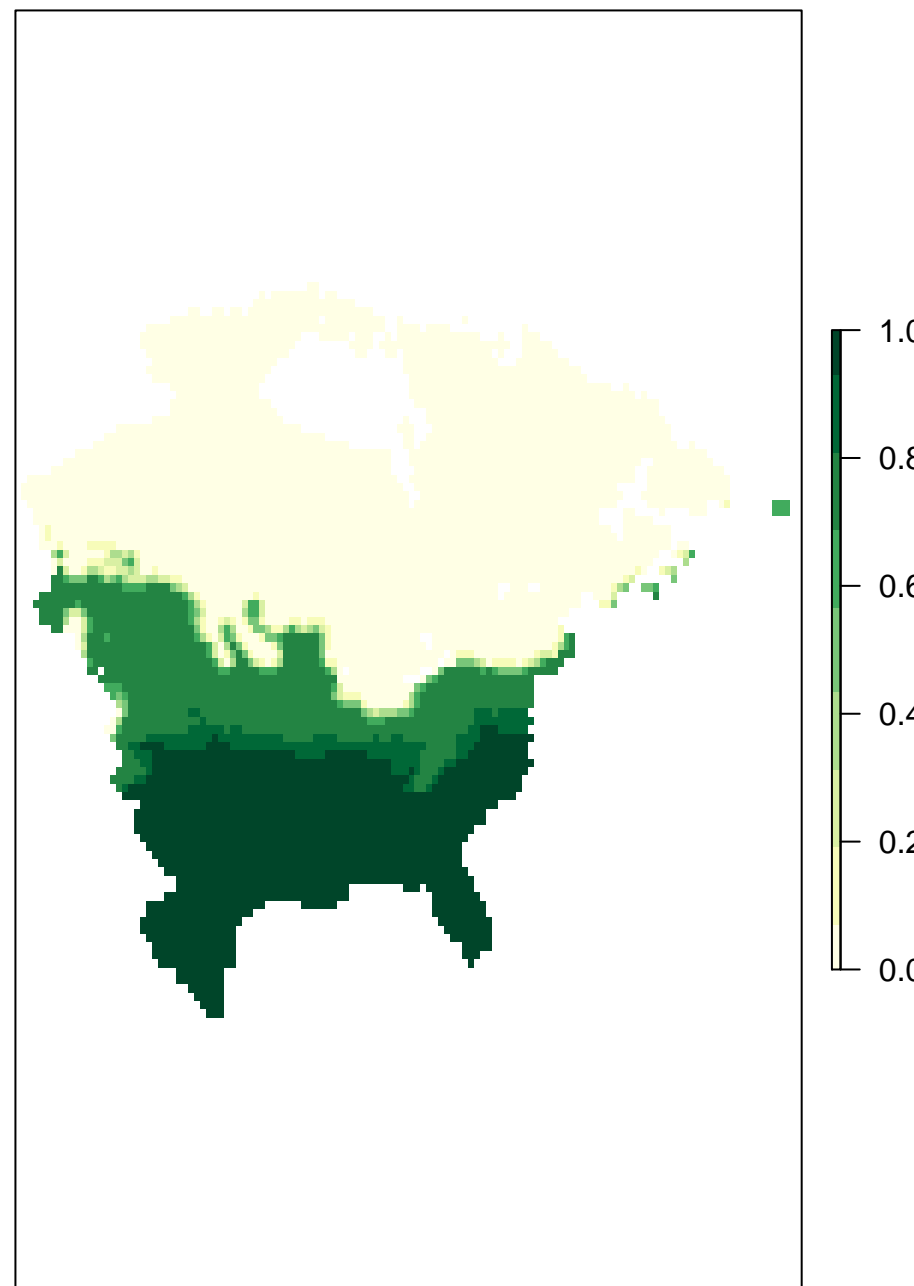


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

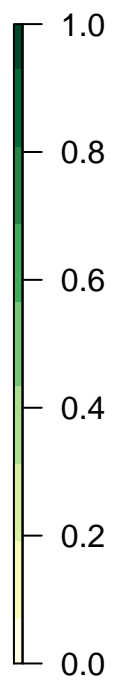
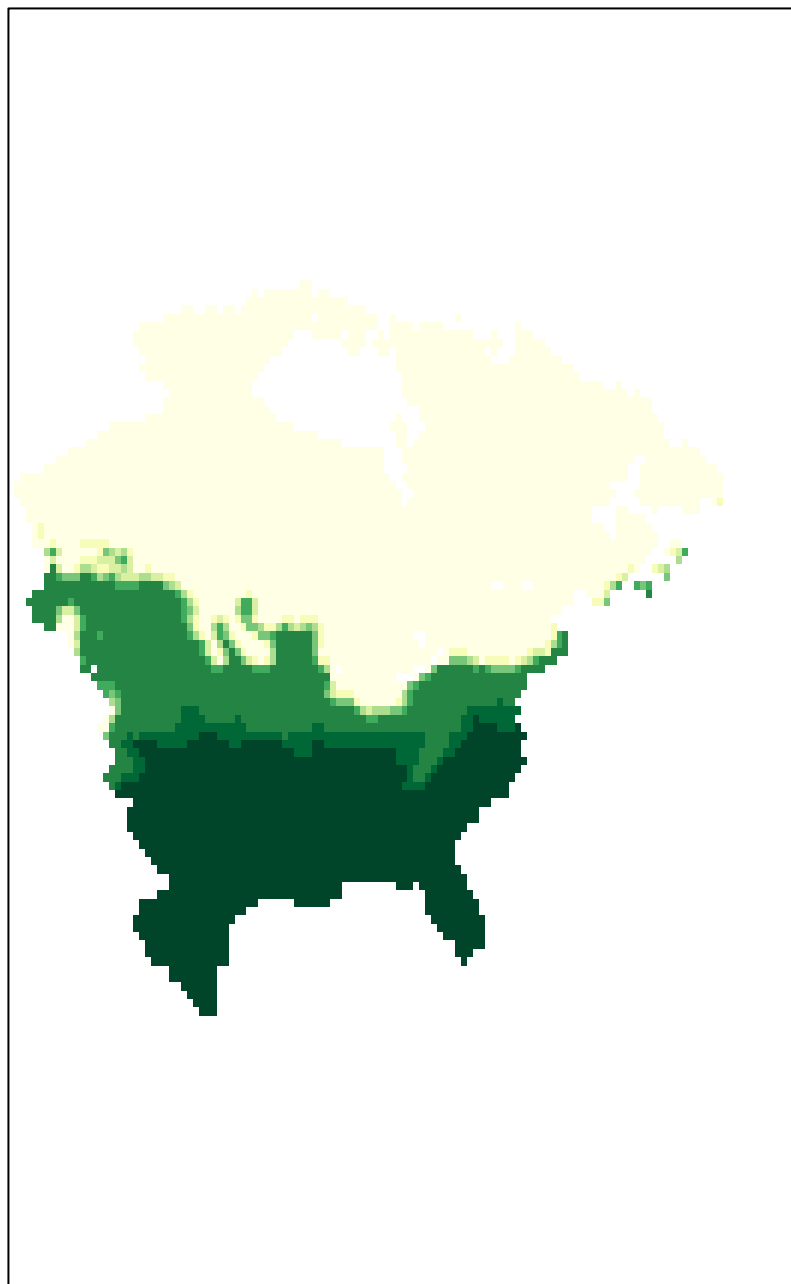
MAX, X5000.ybp



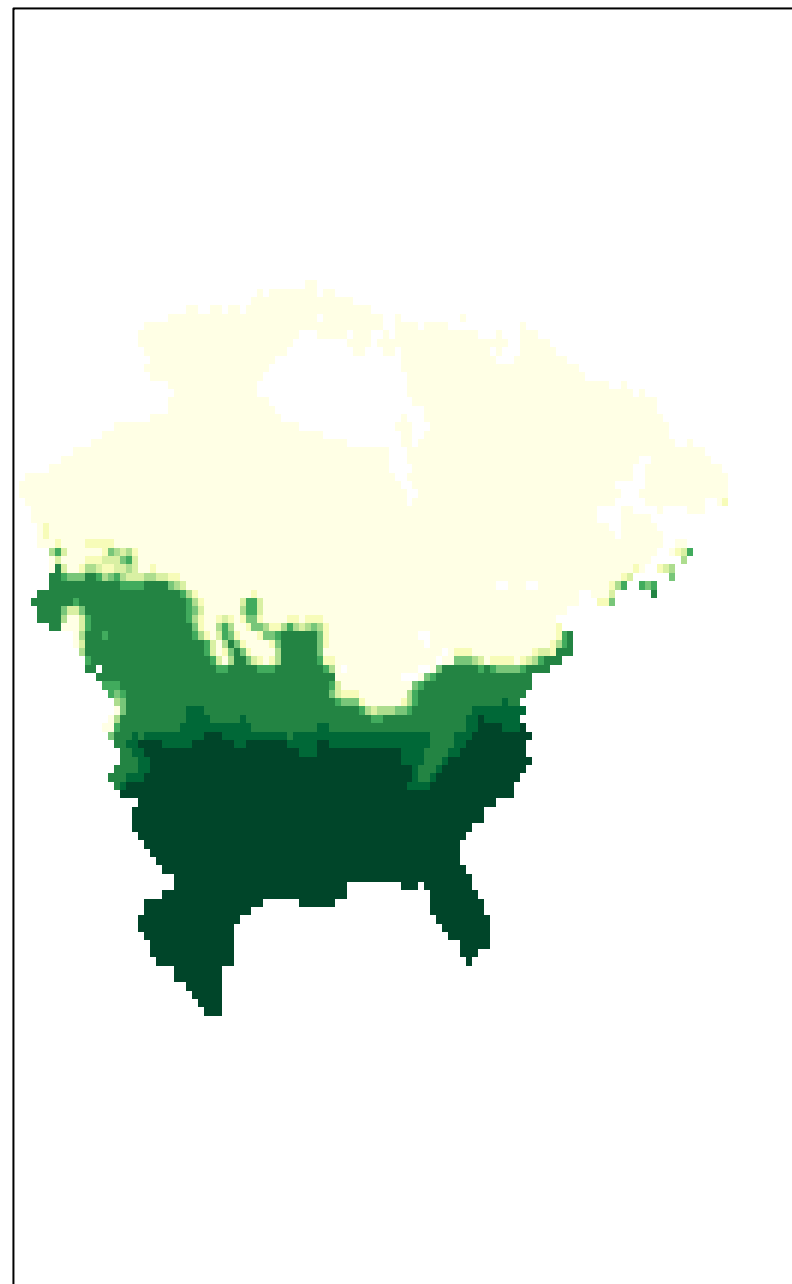
MAX, X5000.ybp



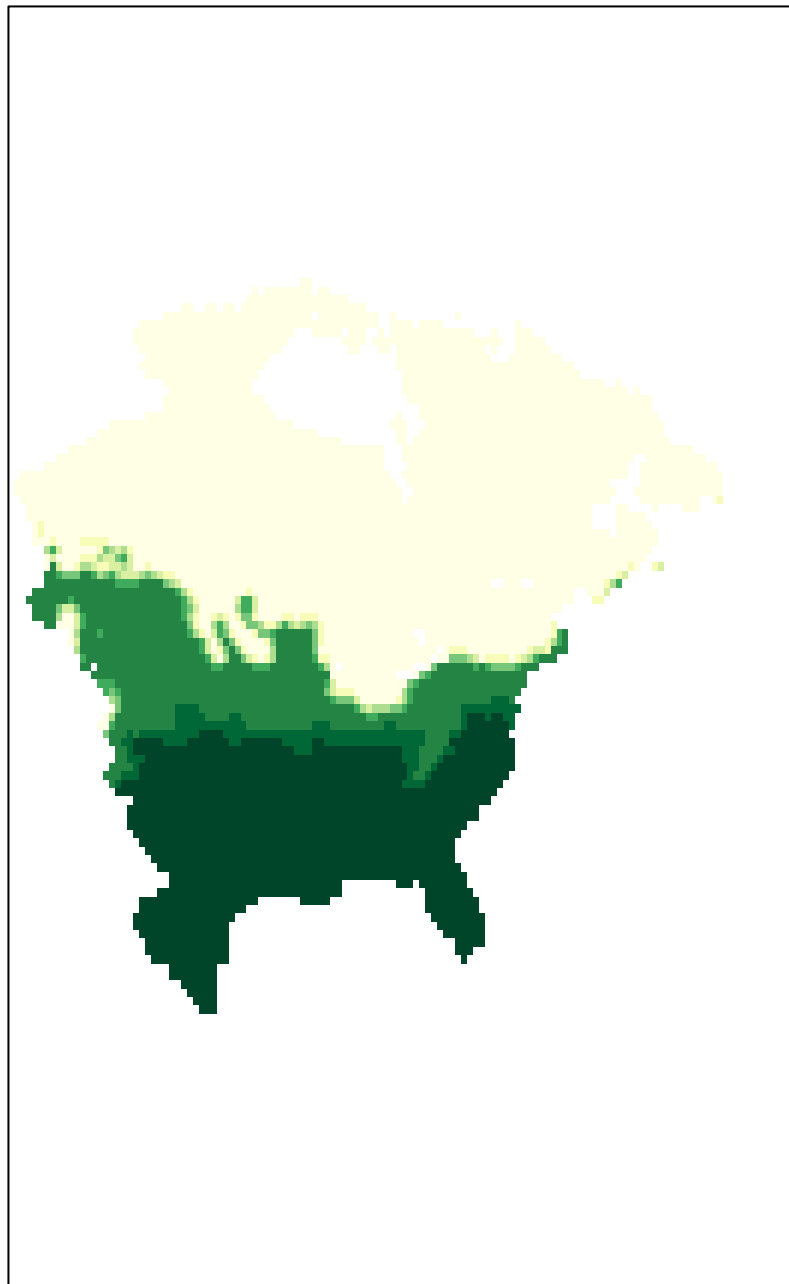
MAX, X4000.ybp



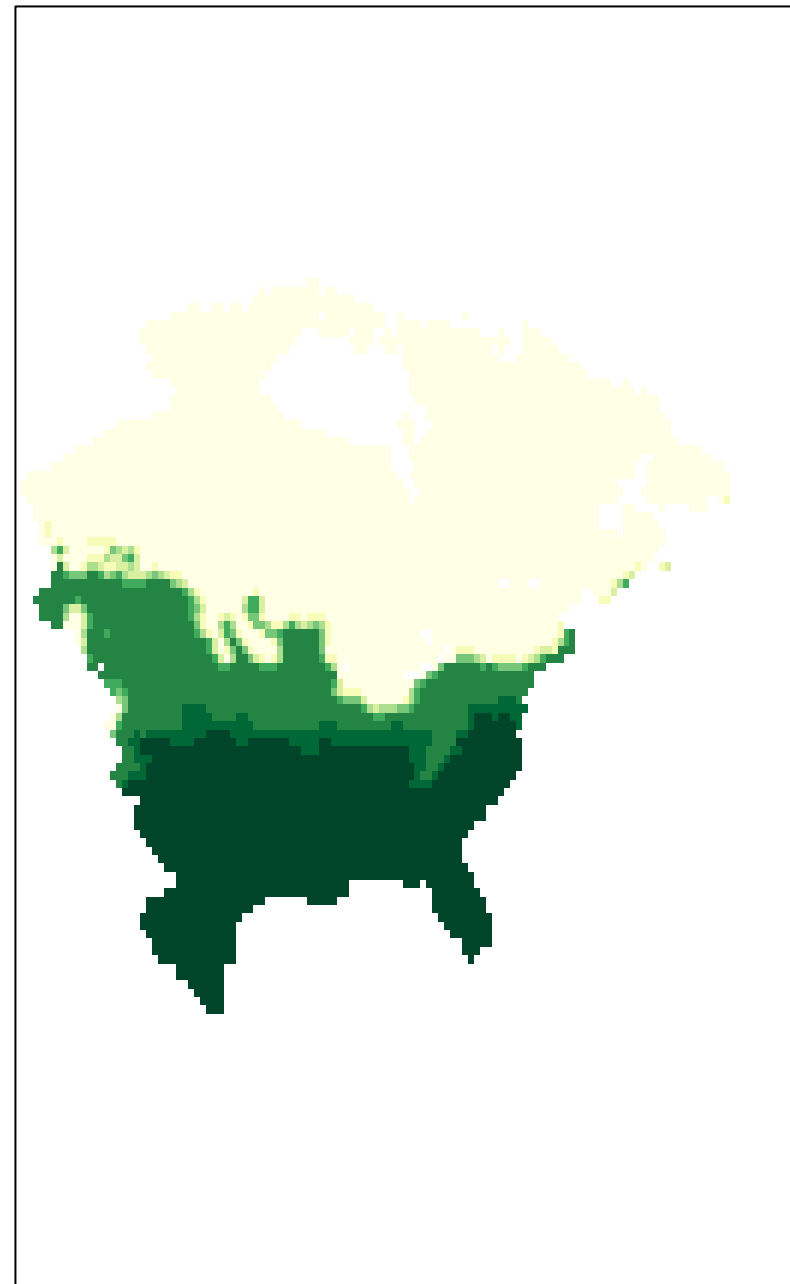
MAX, X4000.ybp



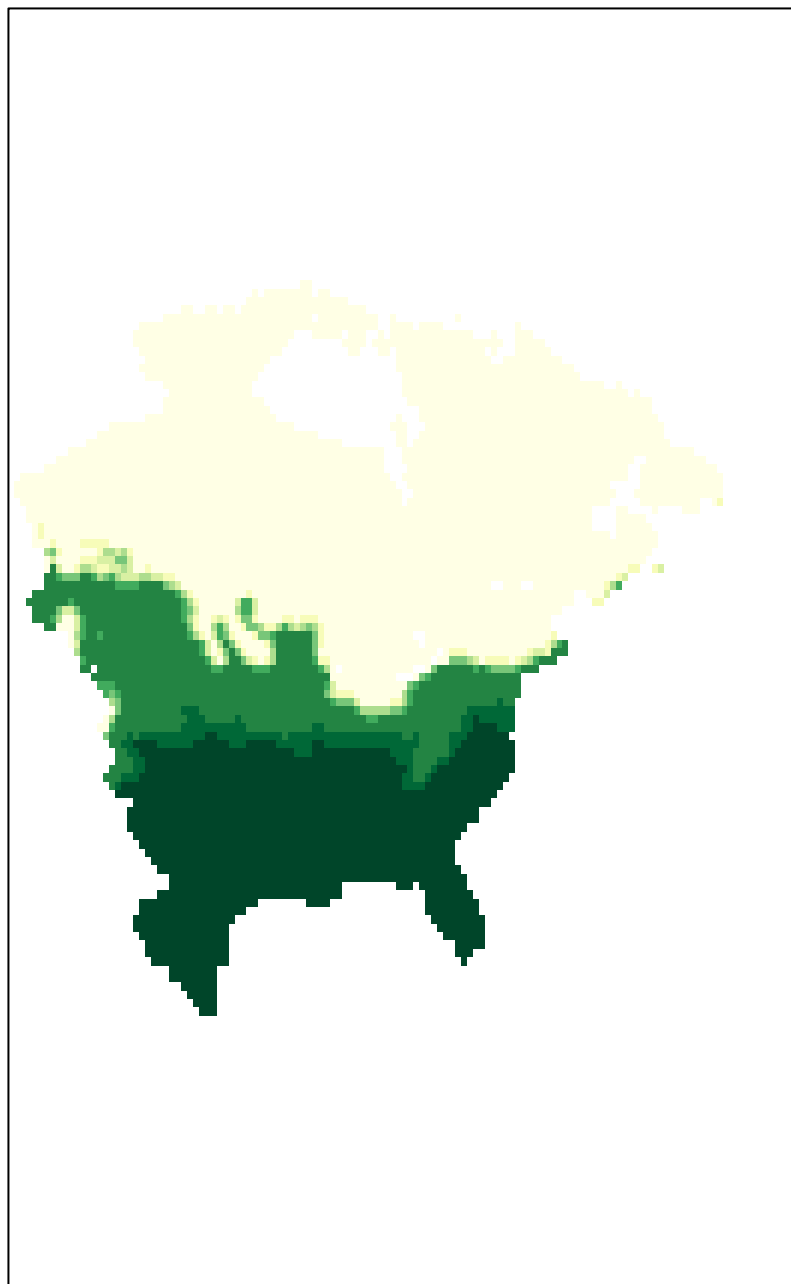
MAX, X3000.ybp



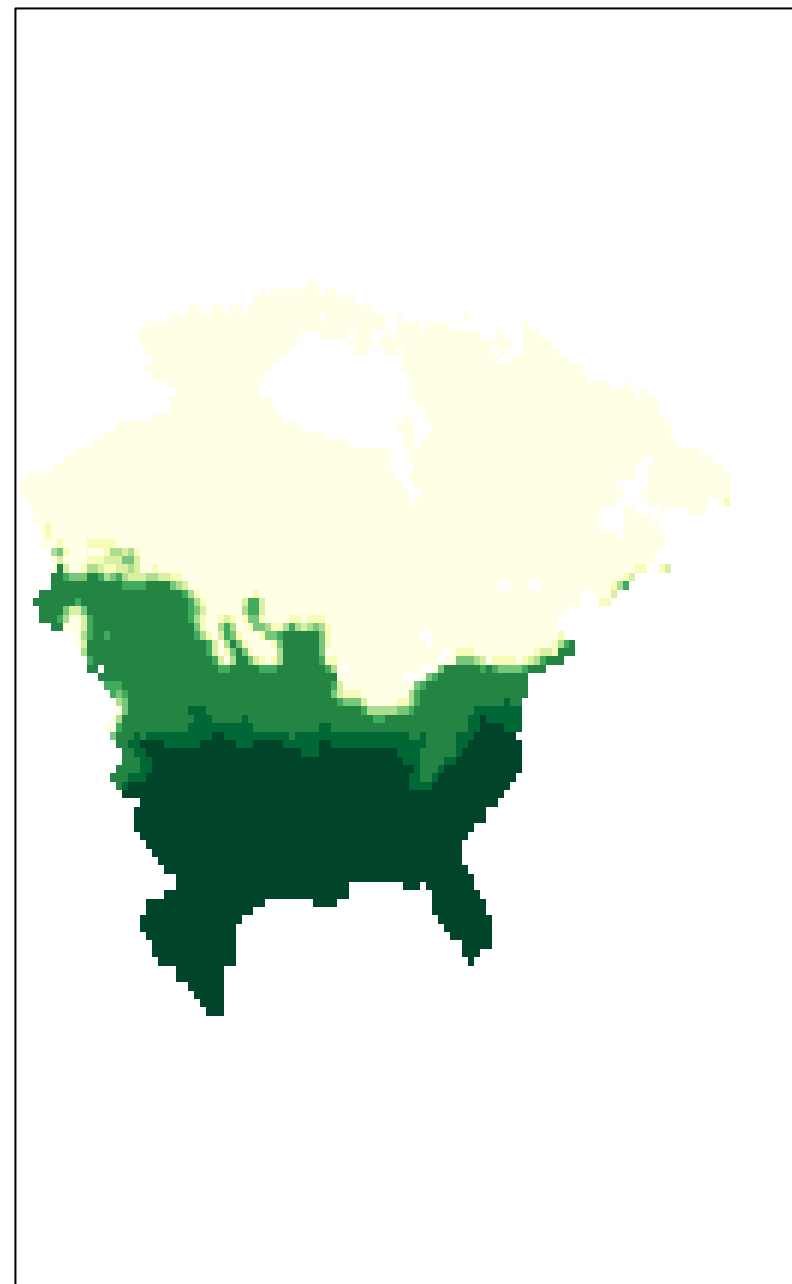
MAX, X3000.ybp



MAX, X2000.ybp

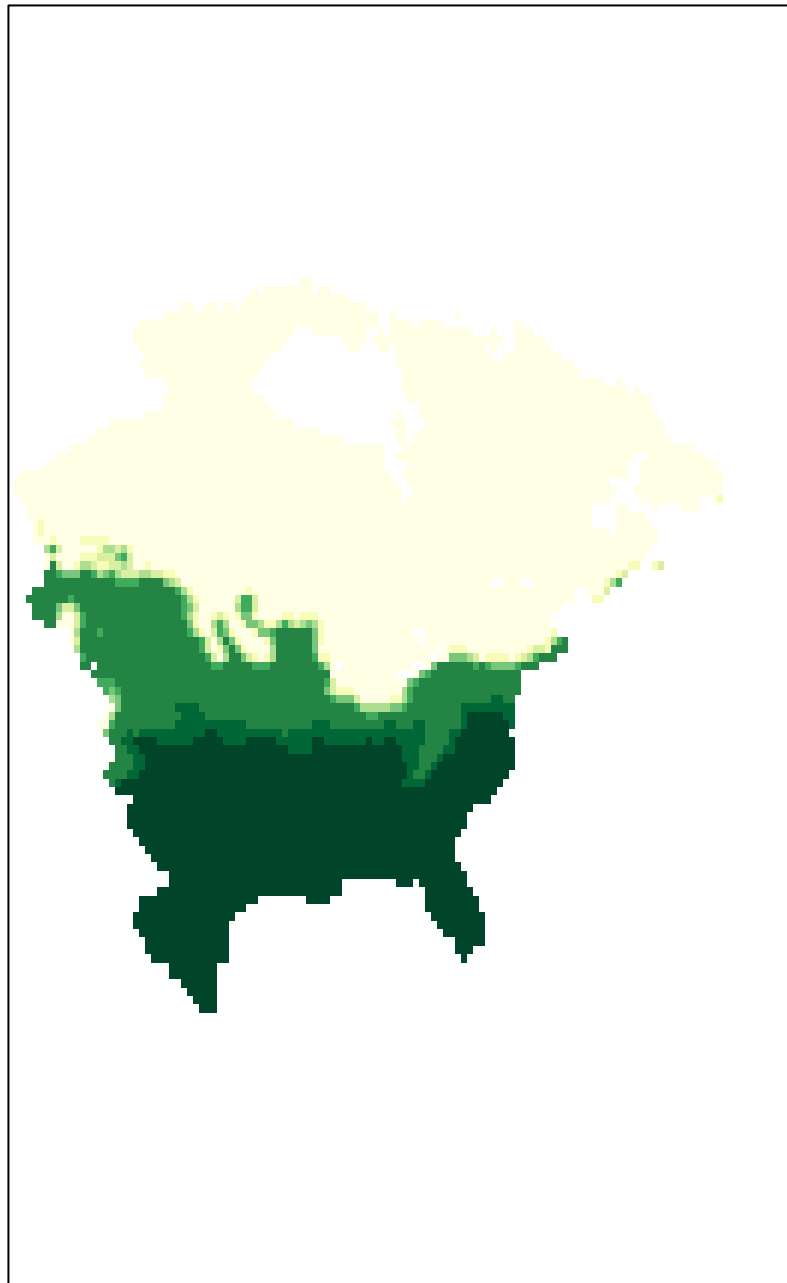


MAX, X2000.ybp

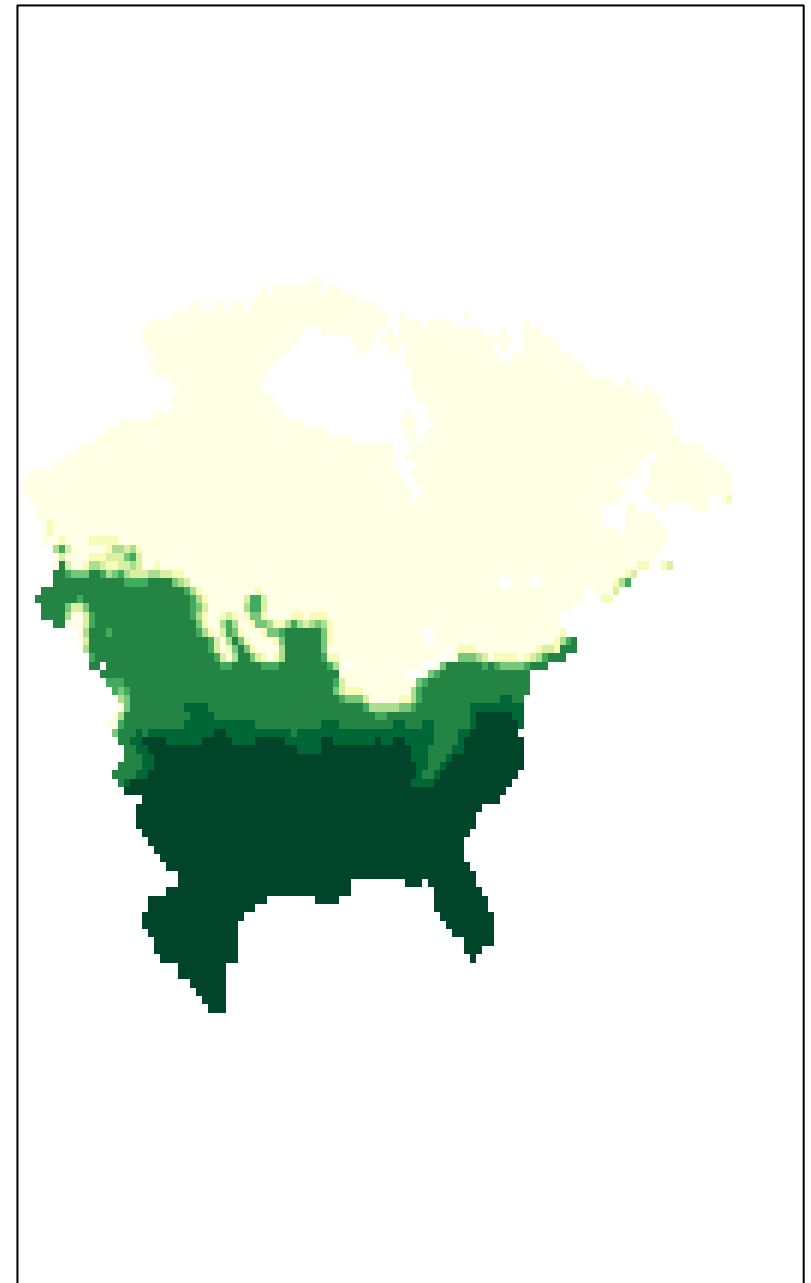


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

MAX, X1000.ybp

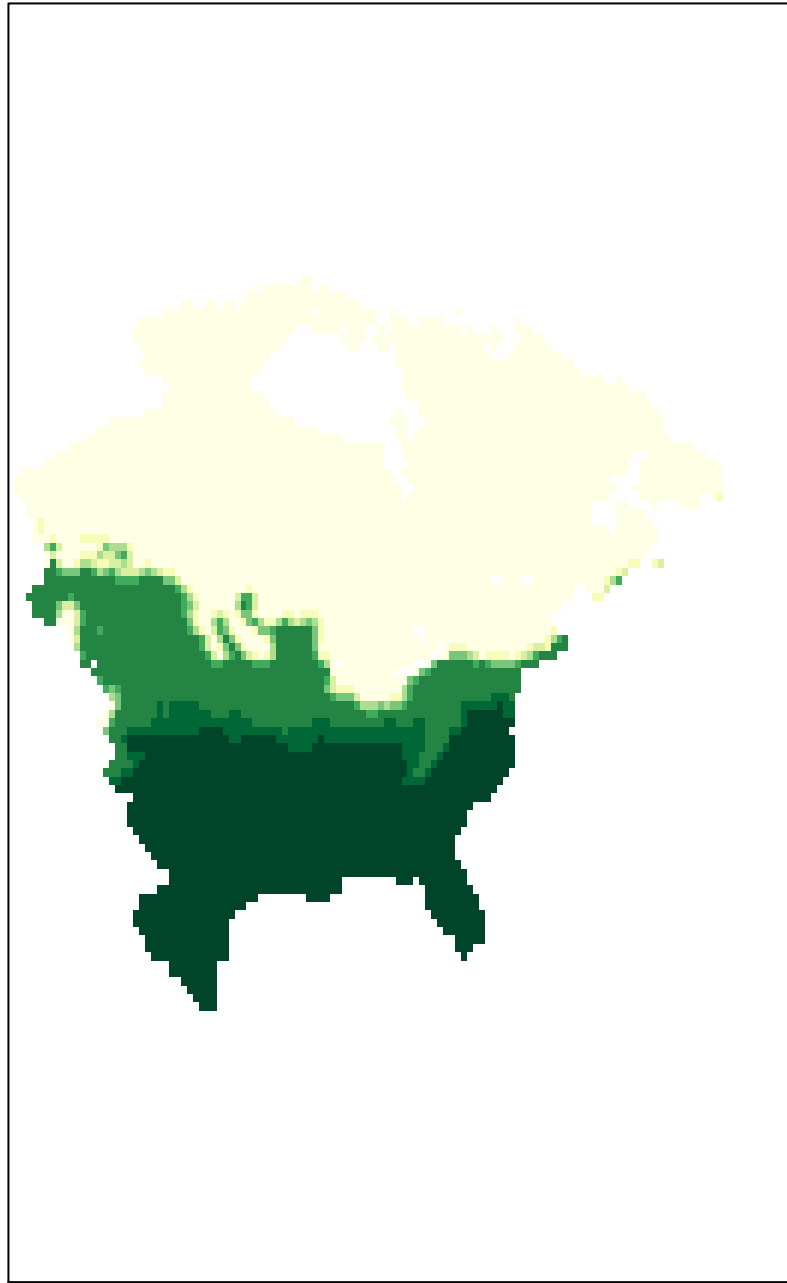


MAX, X1000.ybp

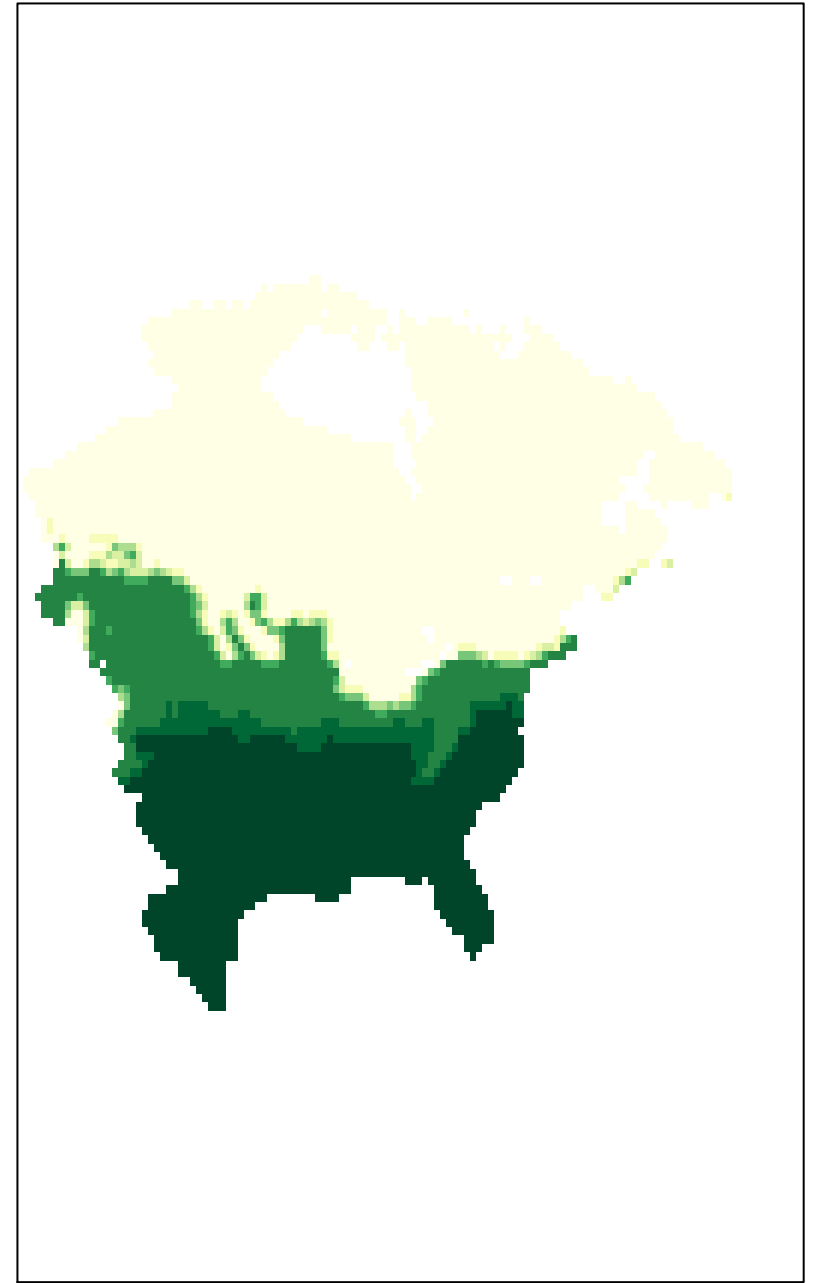


Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm

MAX, X0.ybp



MAX, X0.ybp



Species skipped = *Fraxinus greggii*, GCM = Lorenz_ccsm