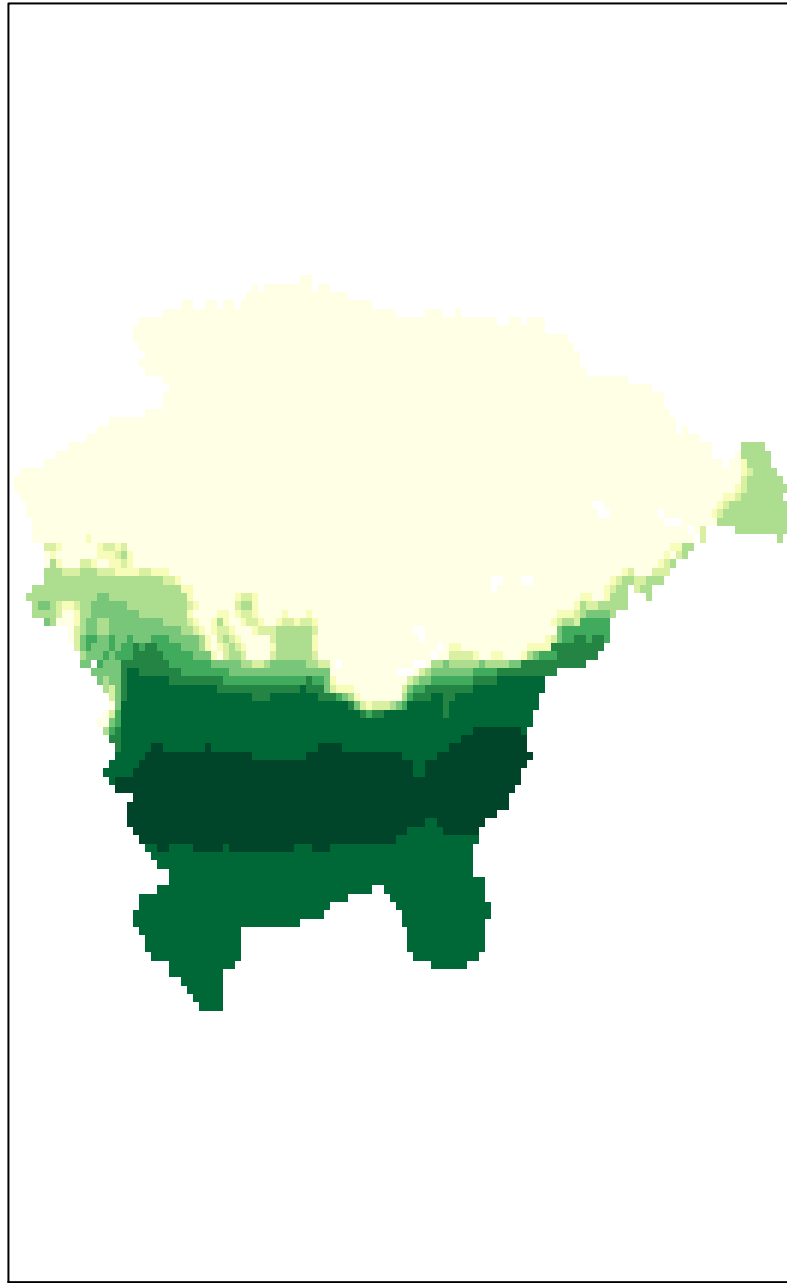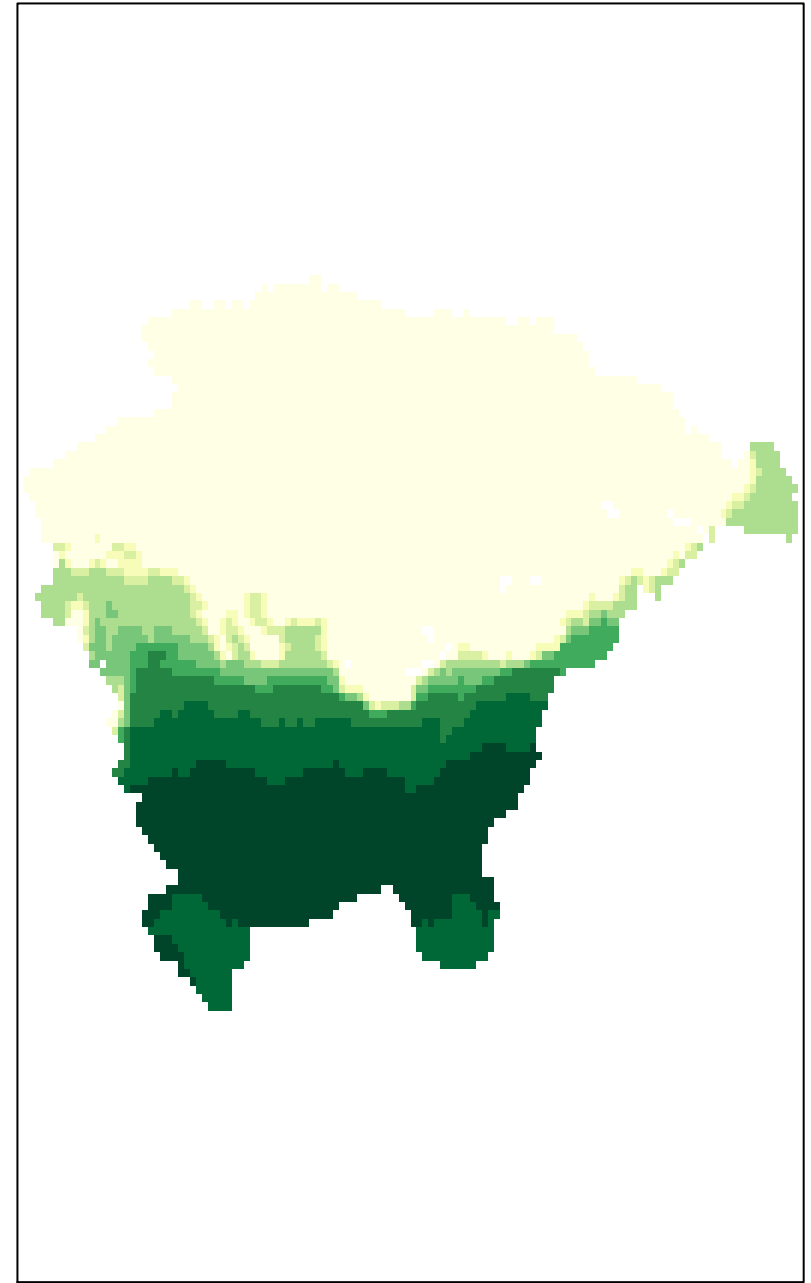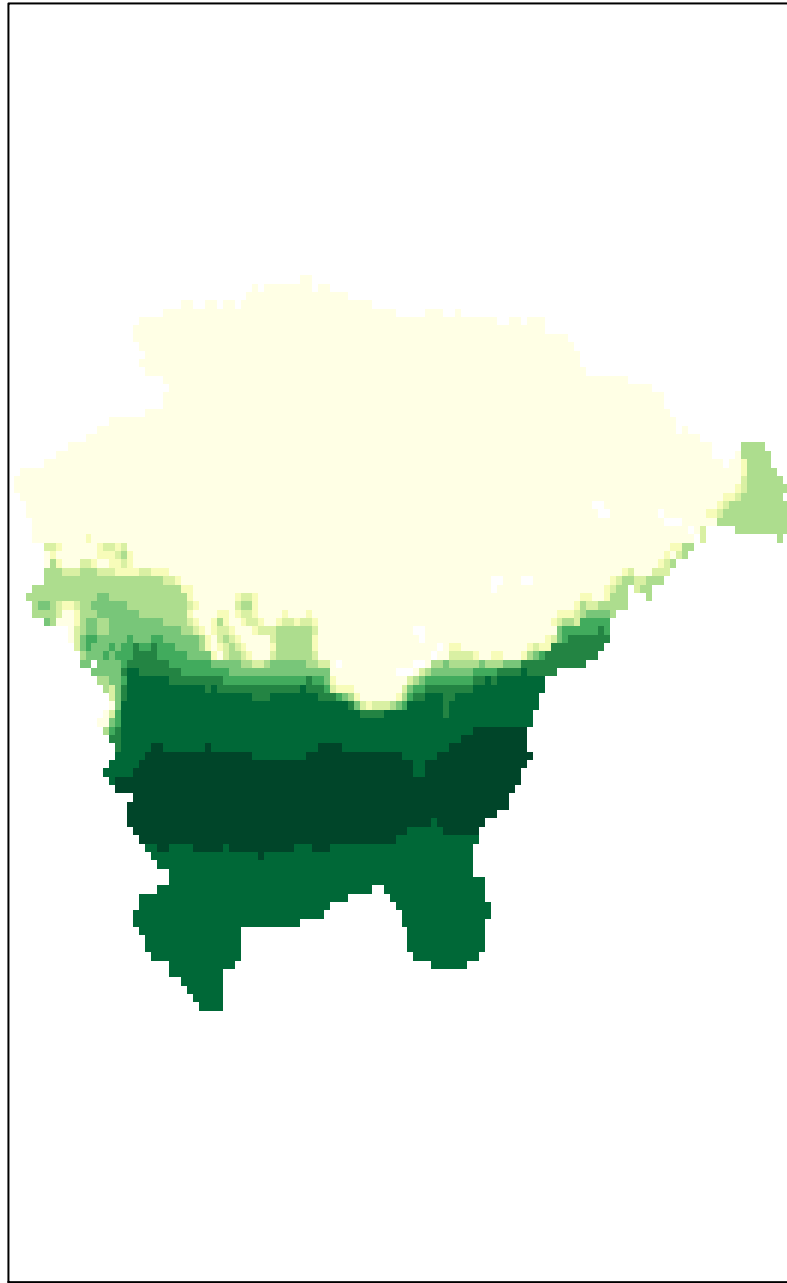**SUM, X21000.ybp**

SUM, X21000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X20000.ybp

SUM, X20000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X19000.ybp
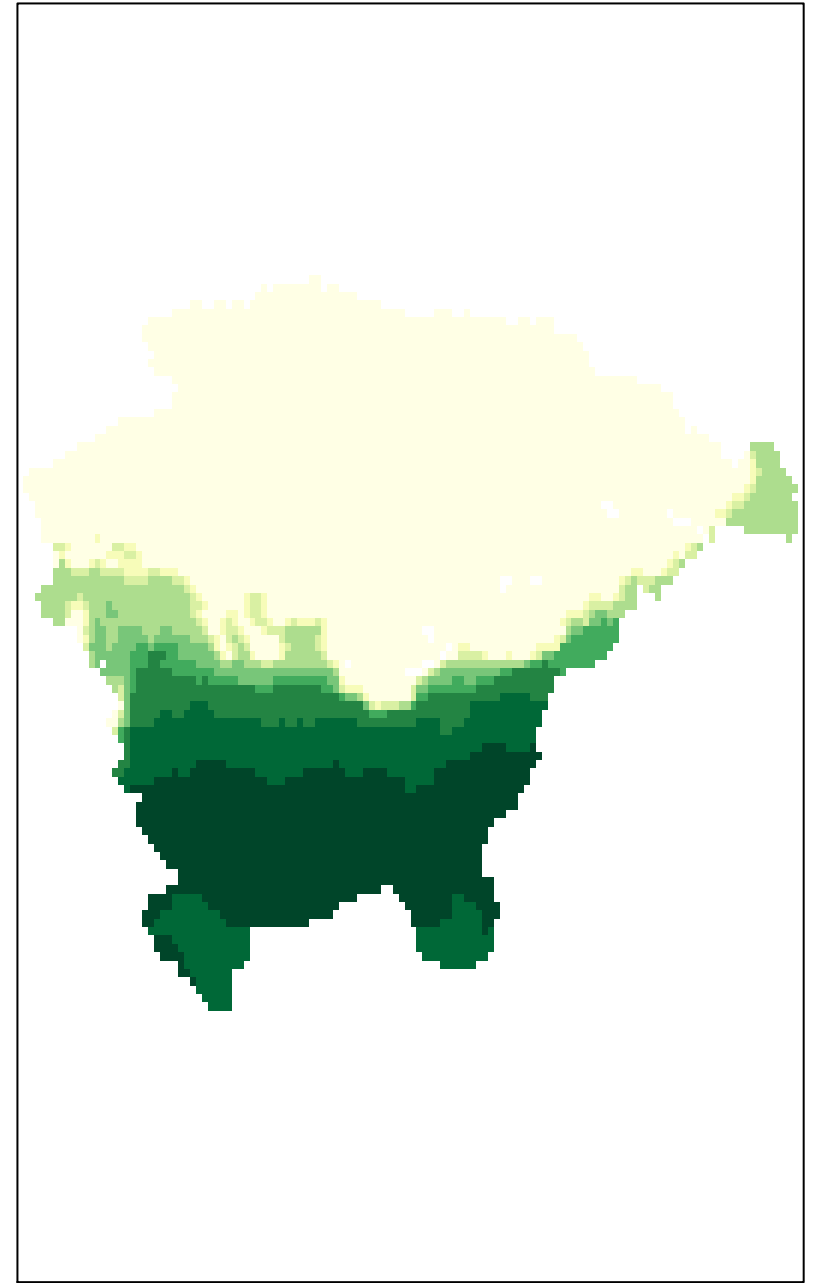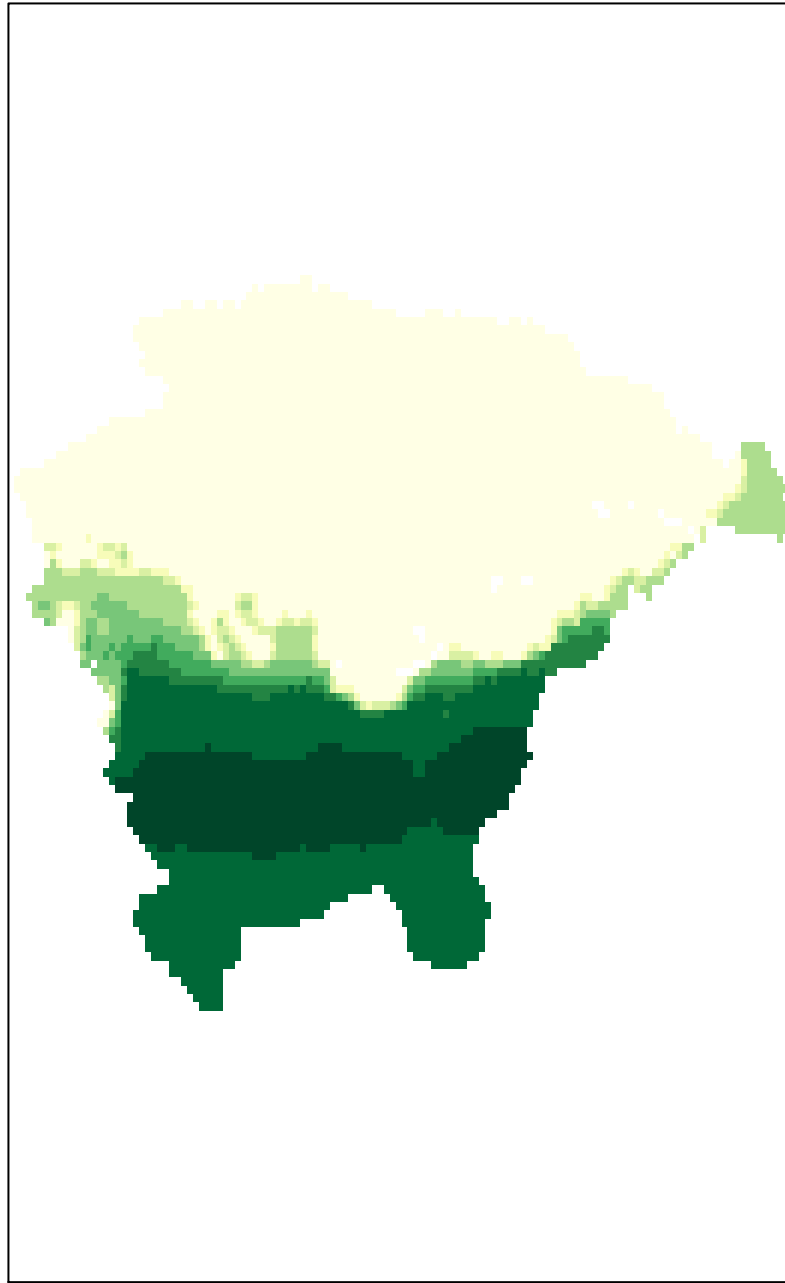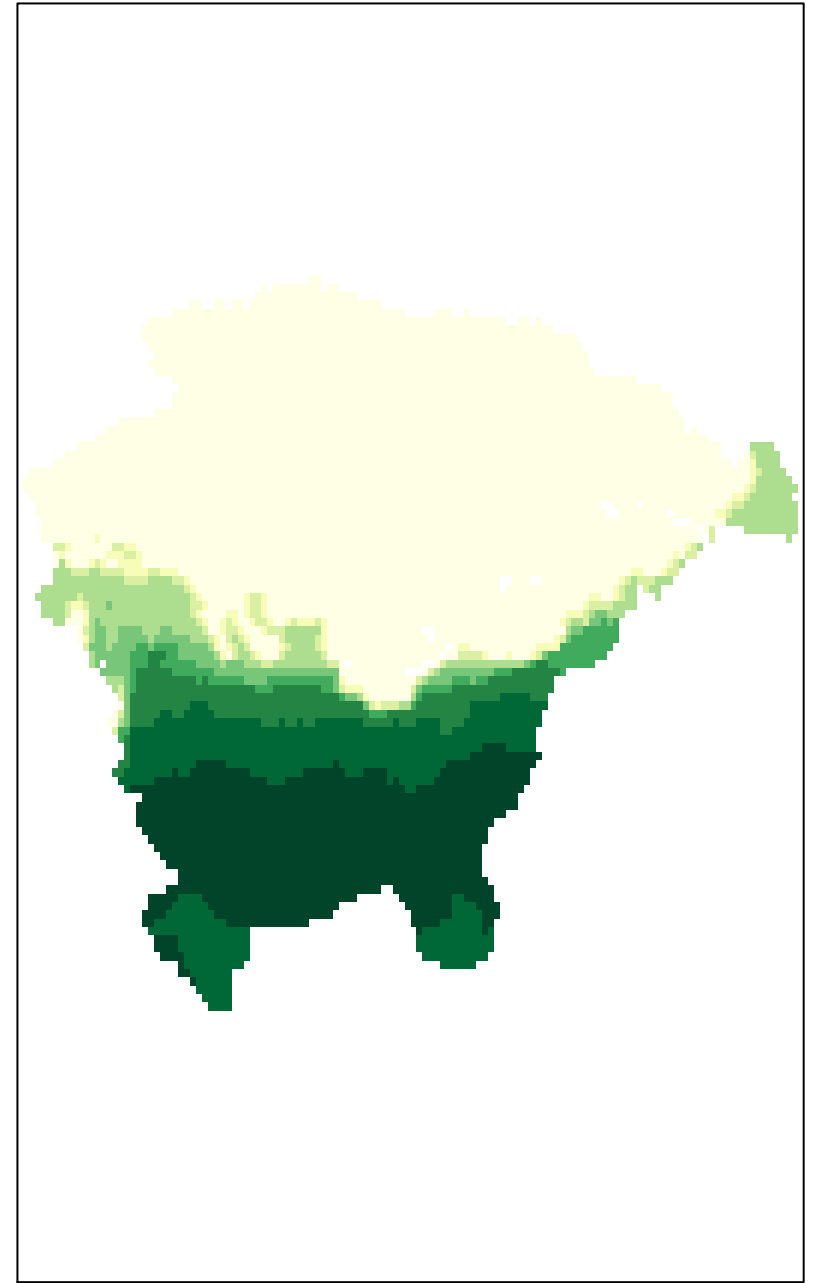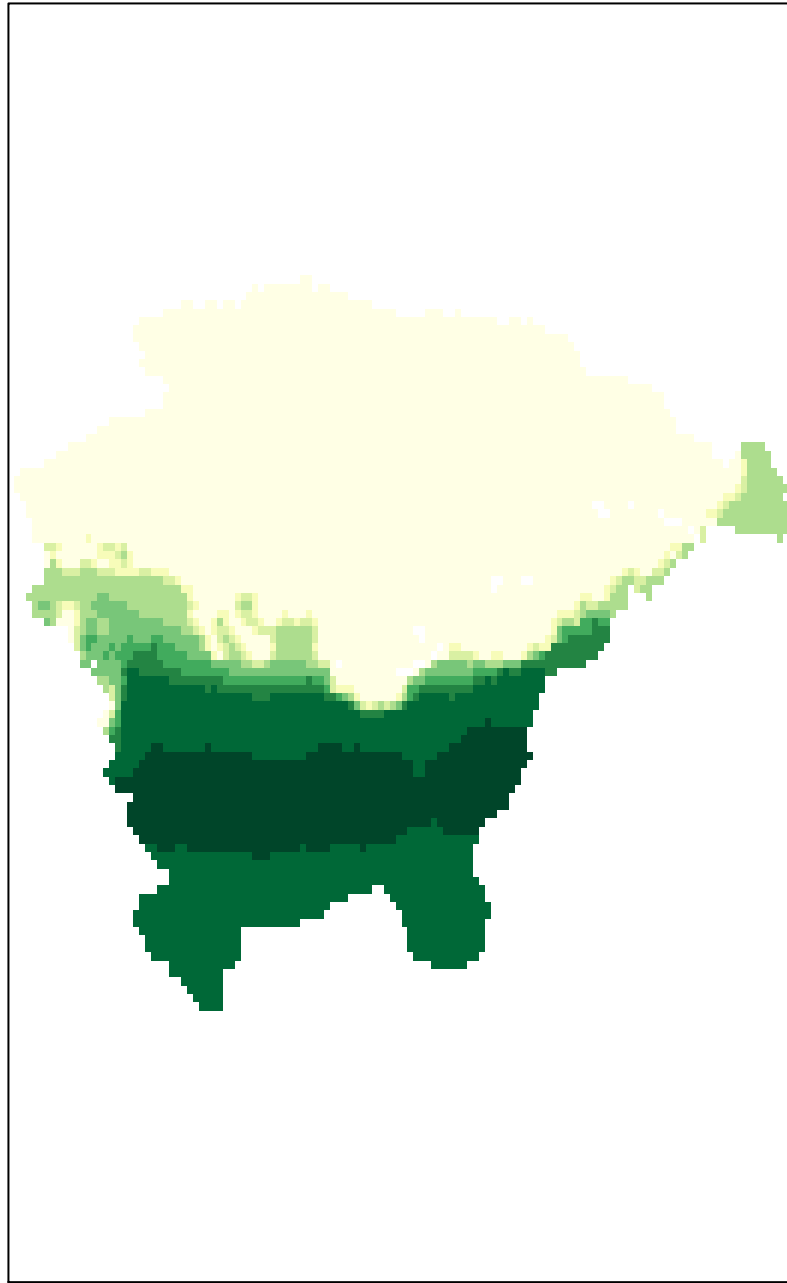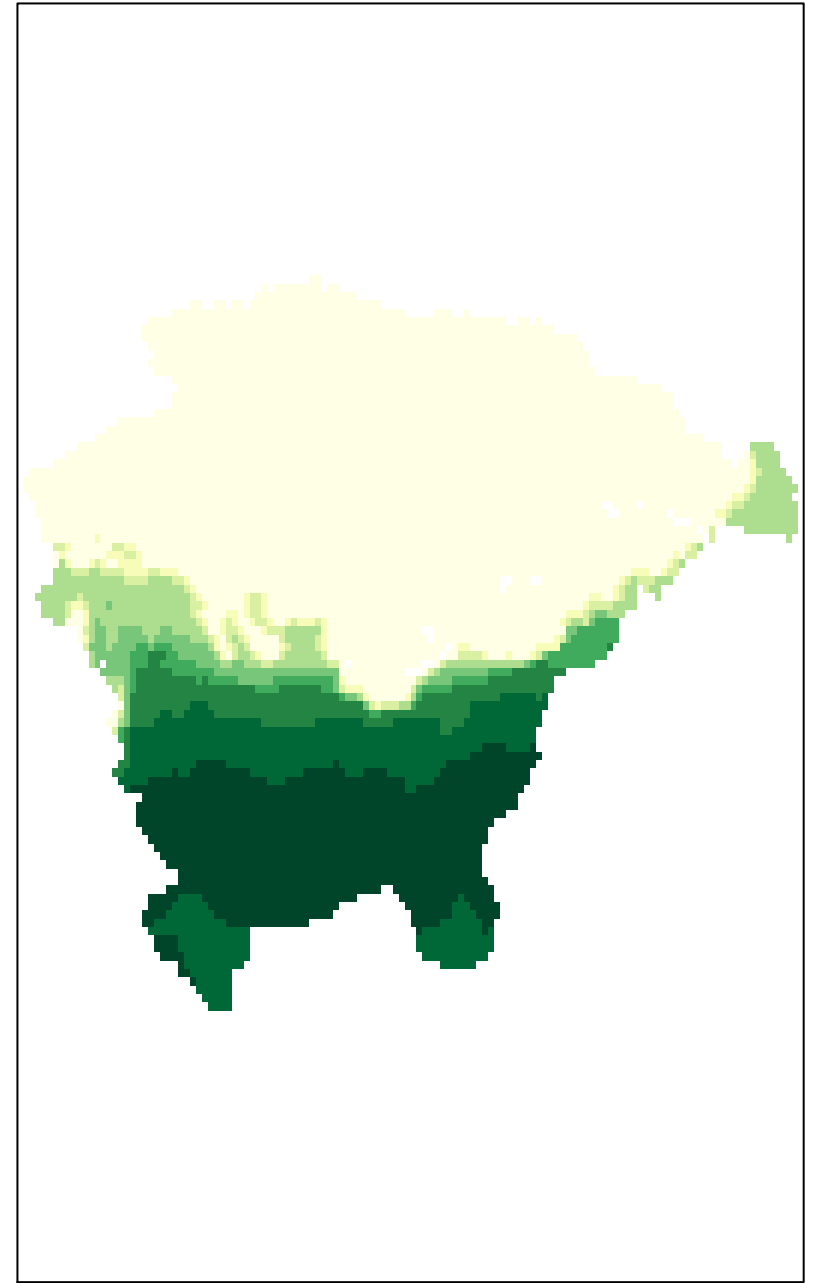
SUM, X19000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X18000.ybp

SUM, X18000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X17000.ybp

SUM, X17000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X16000.ybp

SUM, X16000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X15000.ybp

SUM, X15000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt
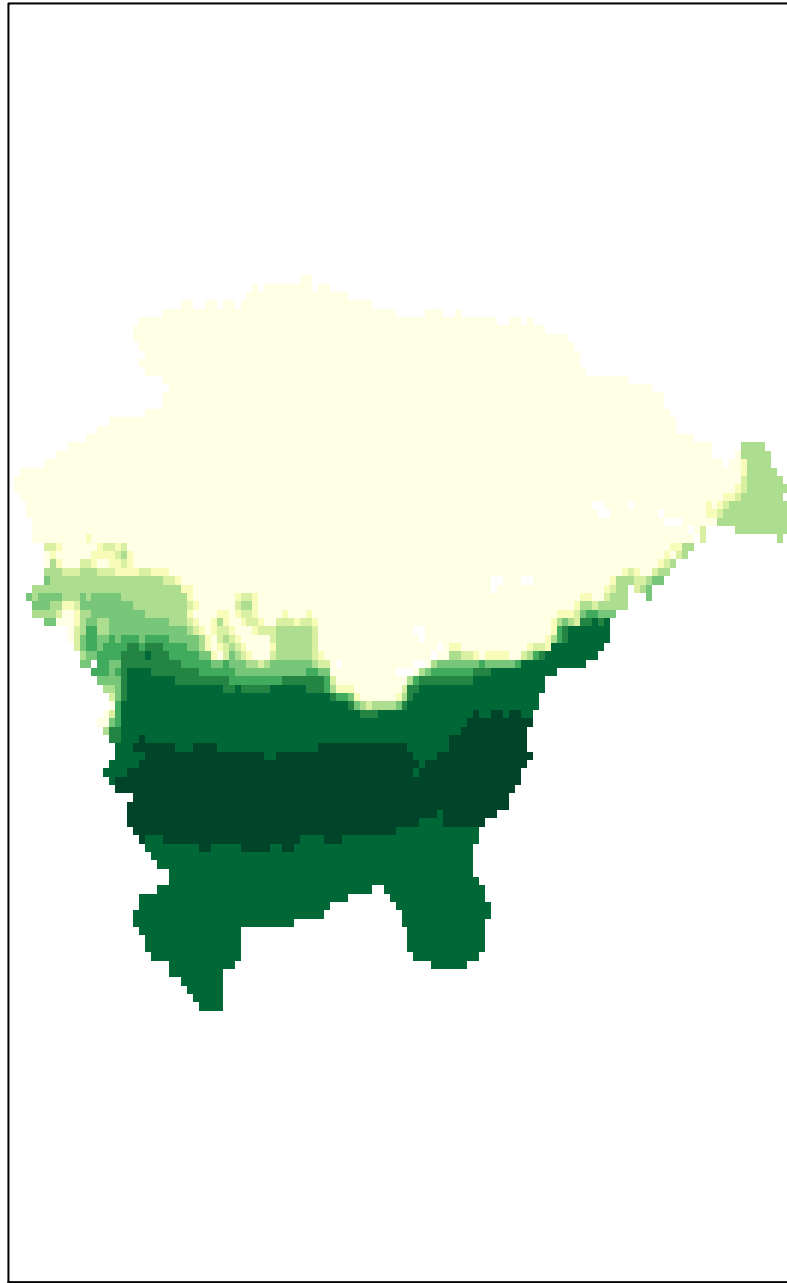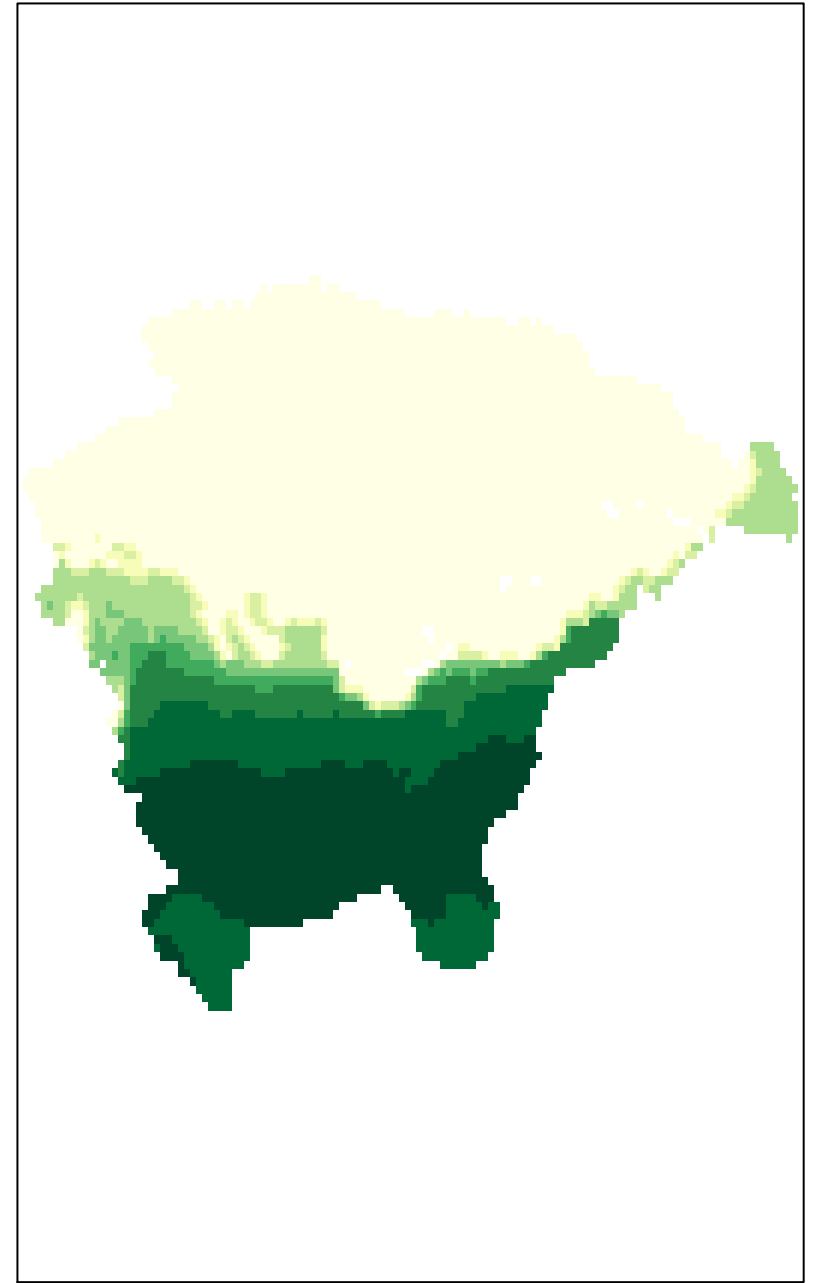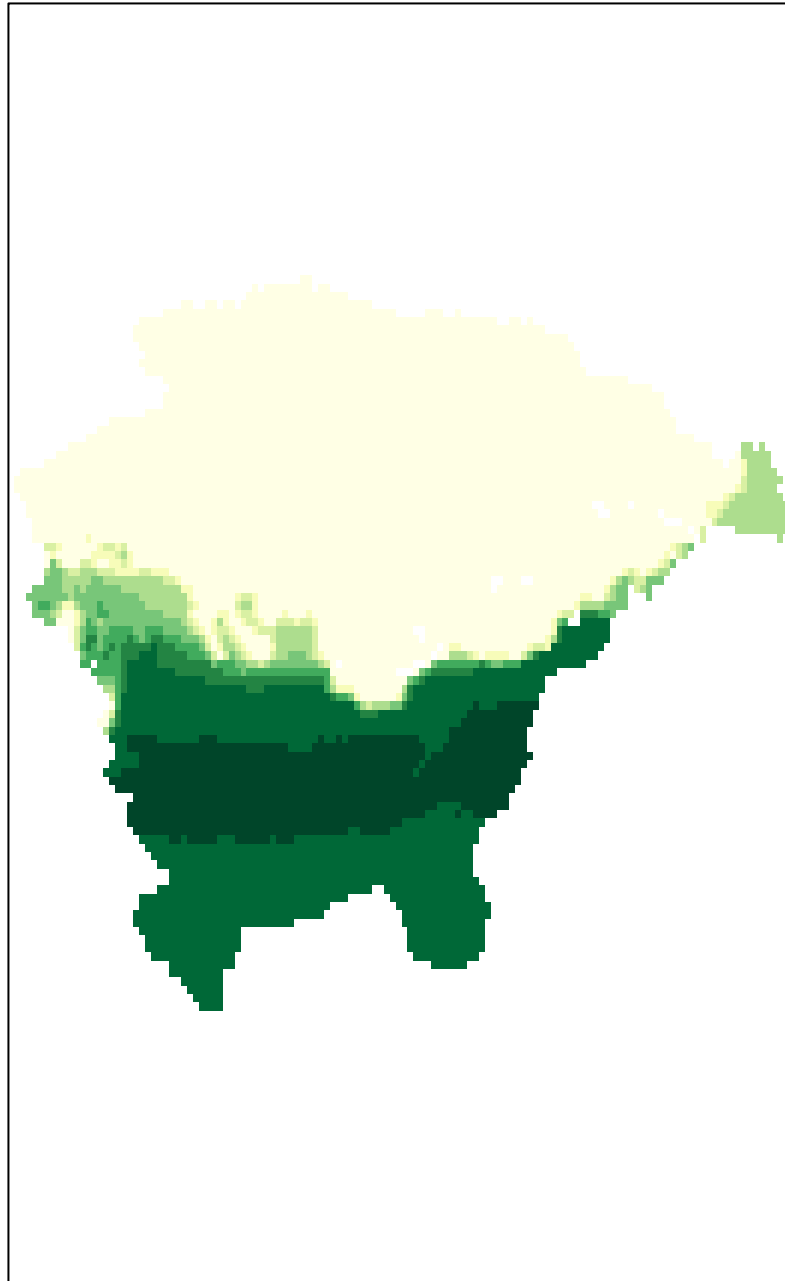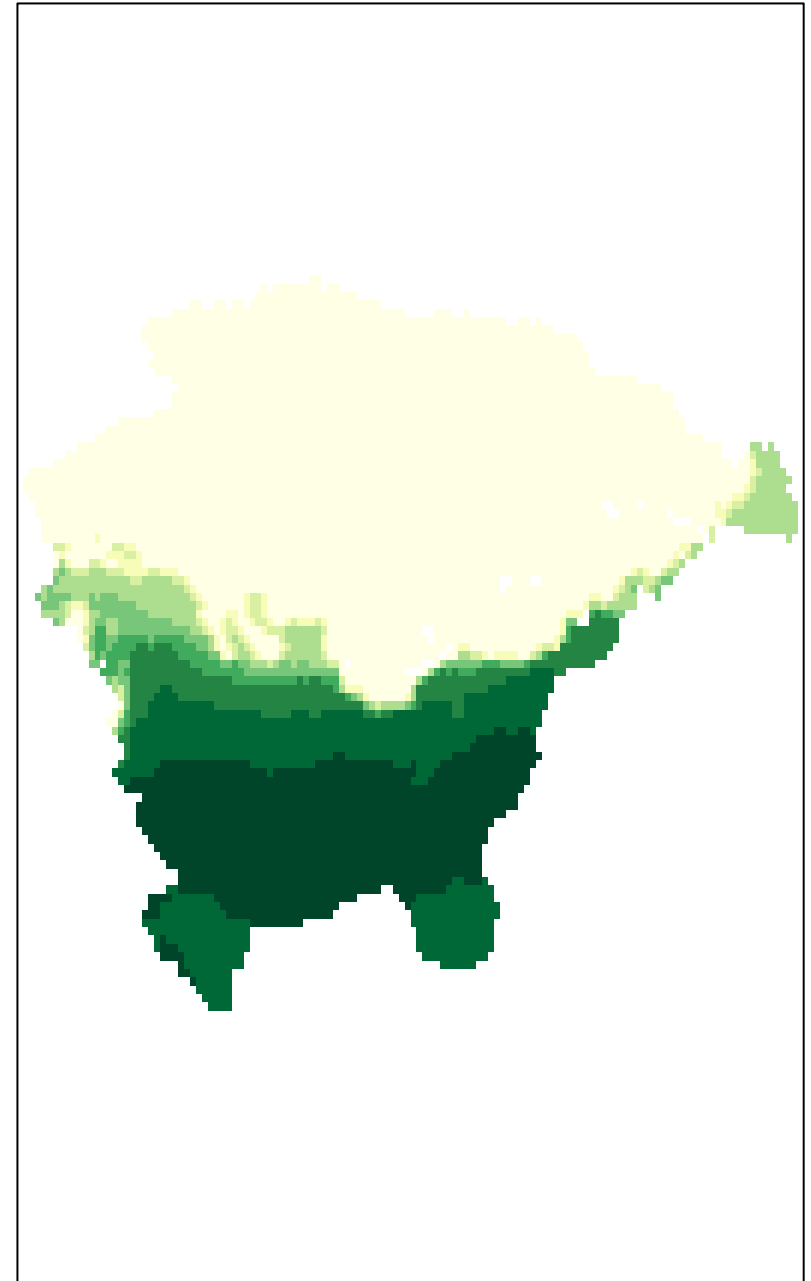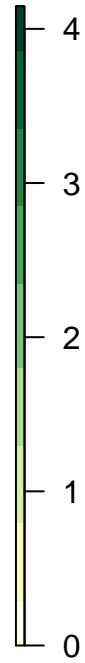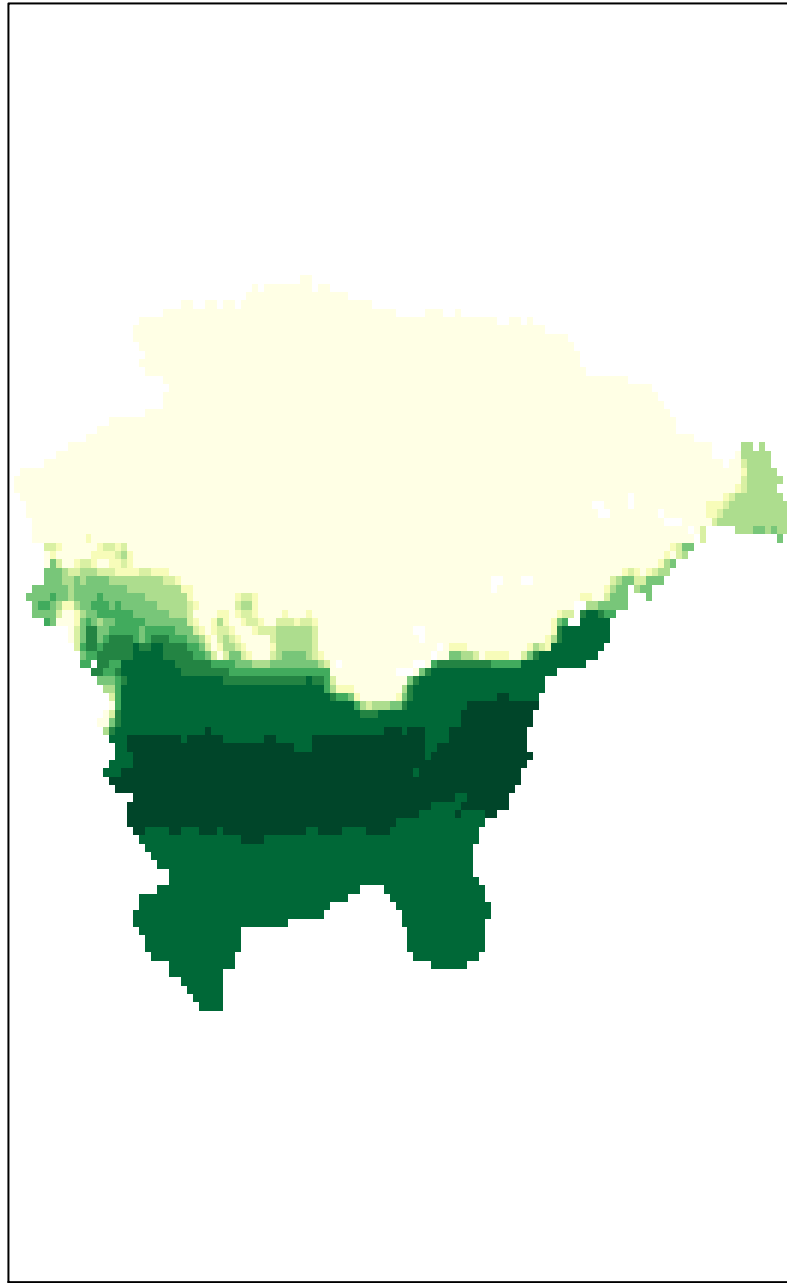
SUM, X14000.ybp
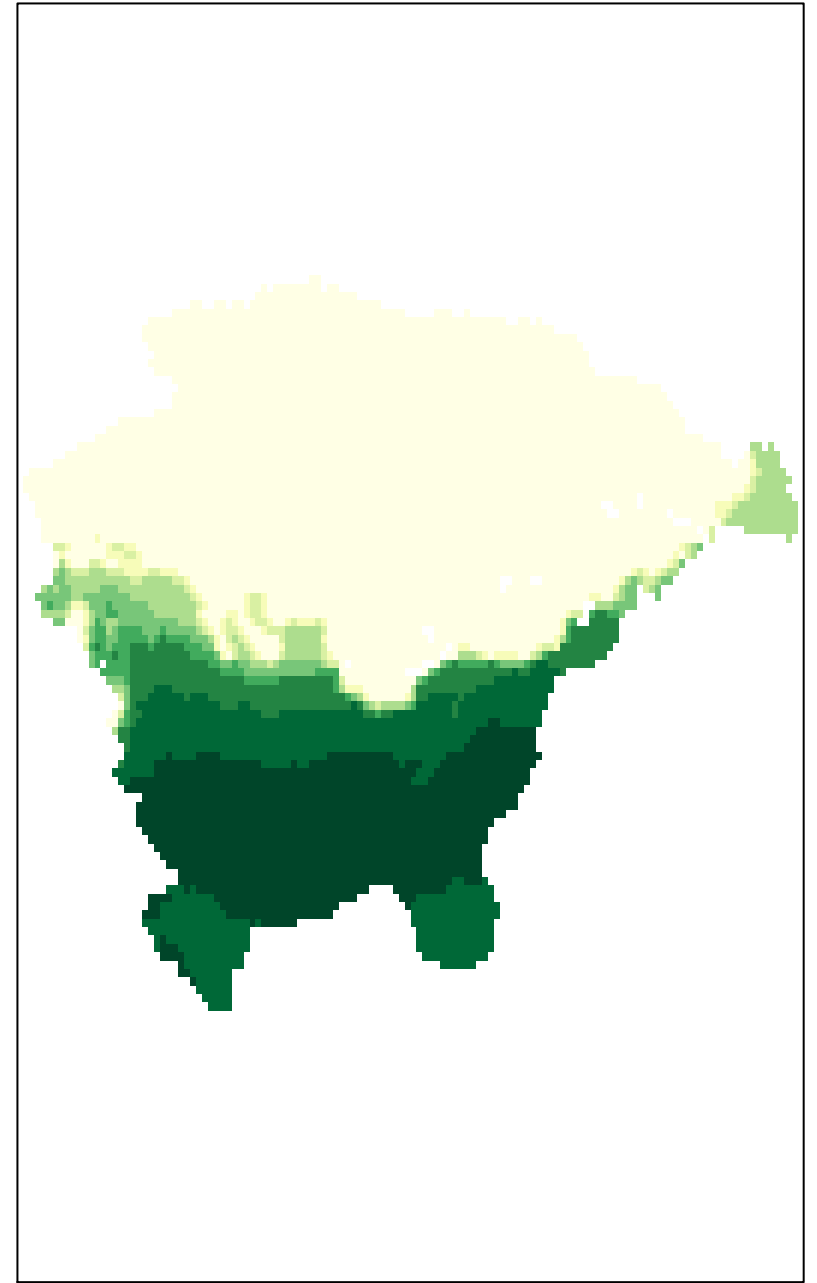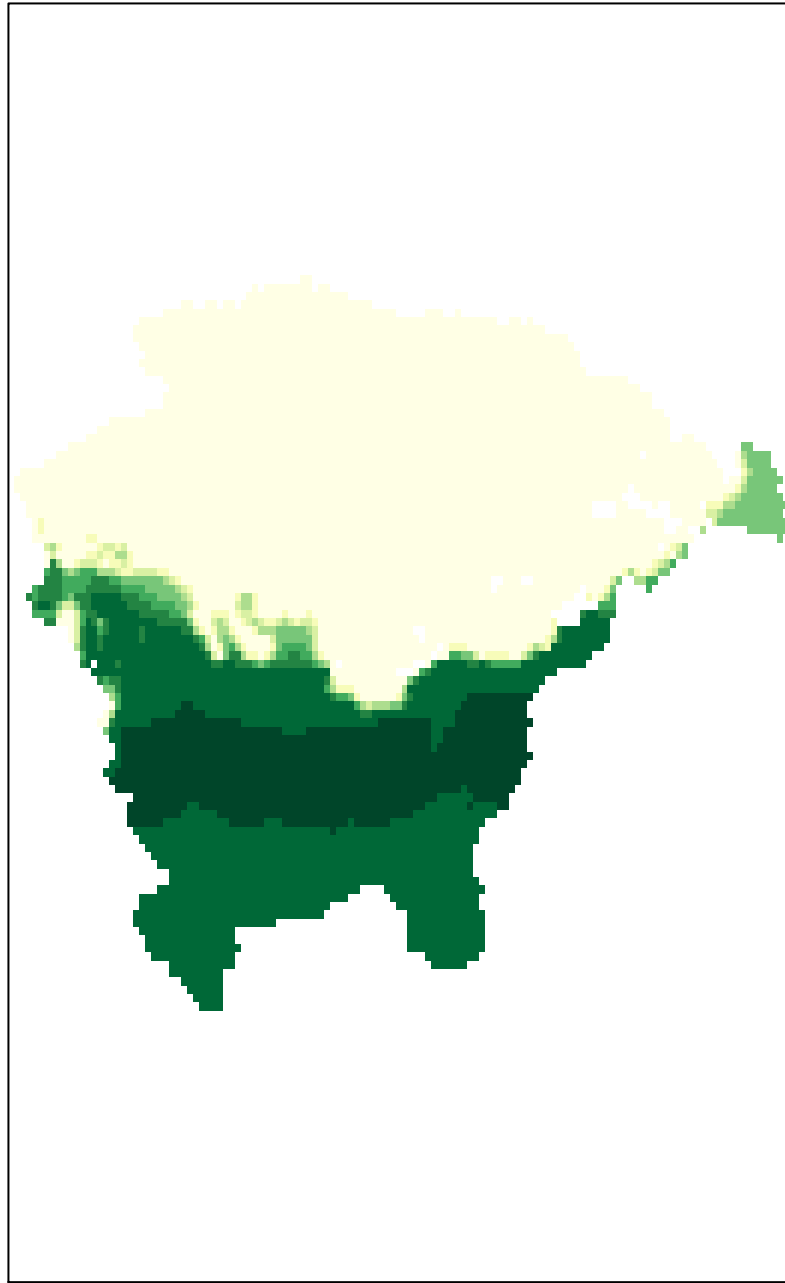
SUM, X14000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt
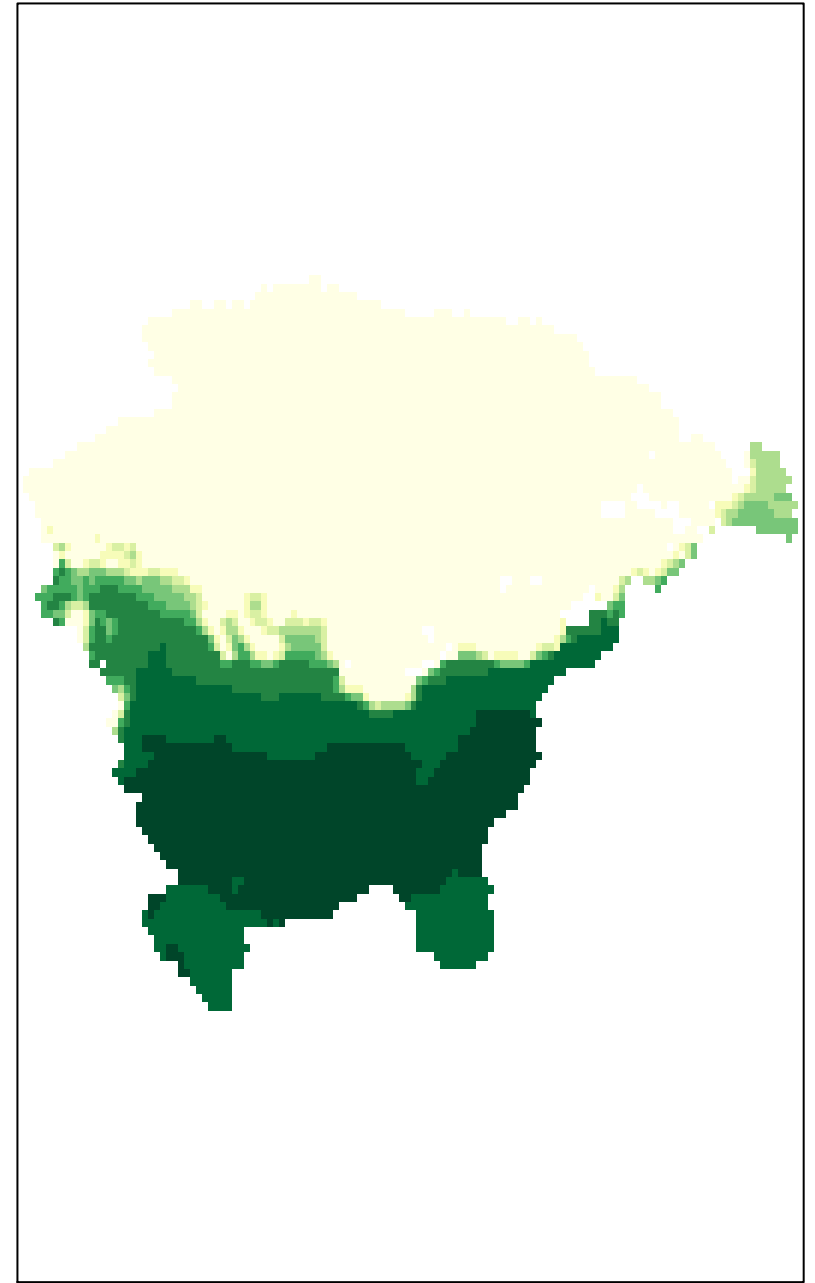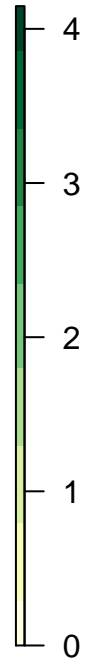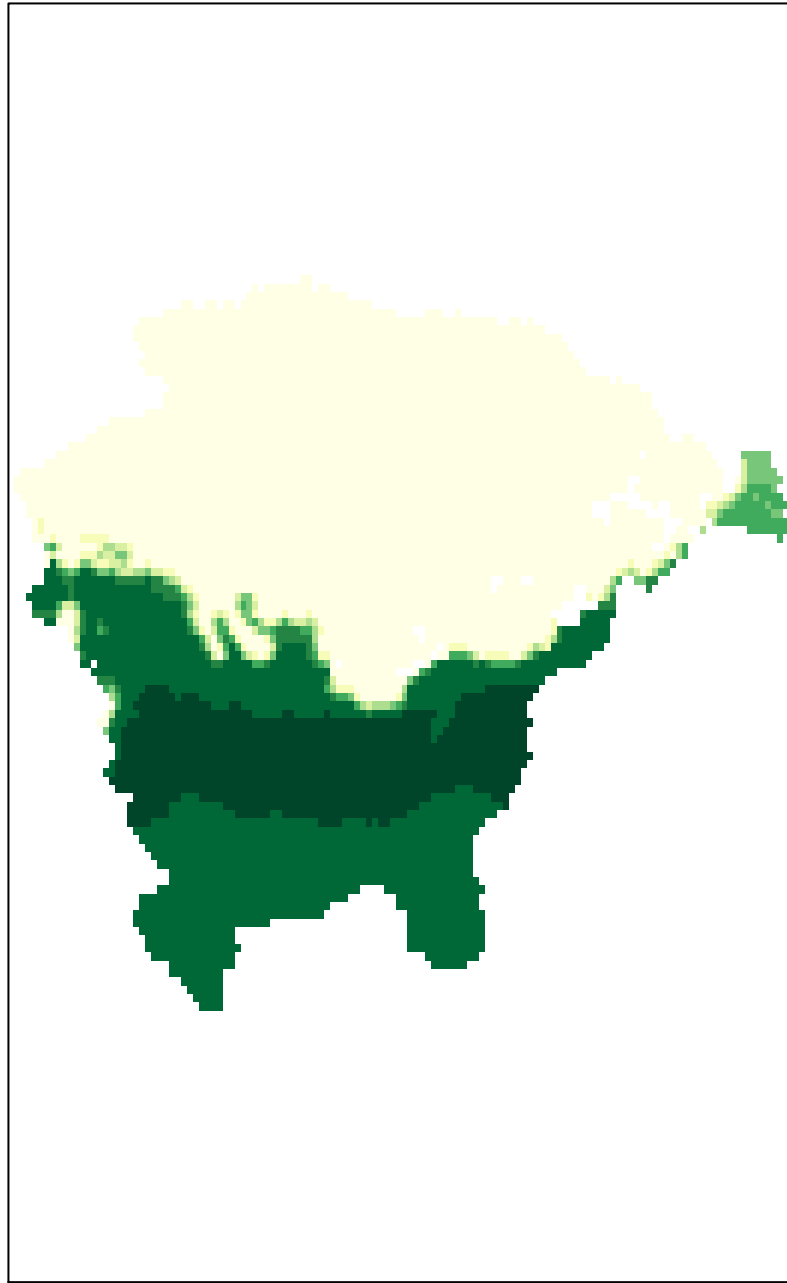
SUM, X13000.ybp   **SUM, X13000.ybp**

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X12000.ybp

SUM, X12000.ybp
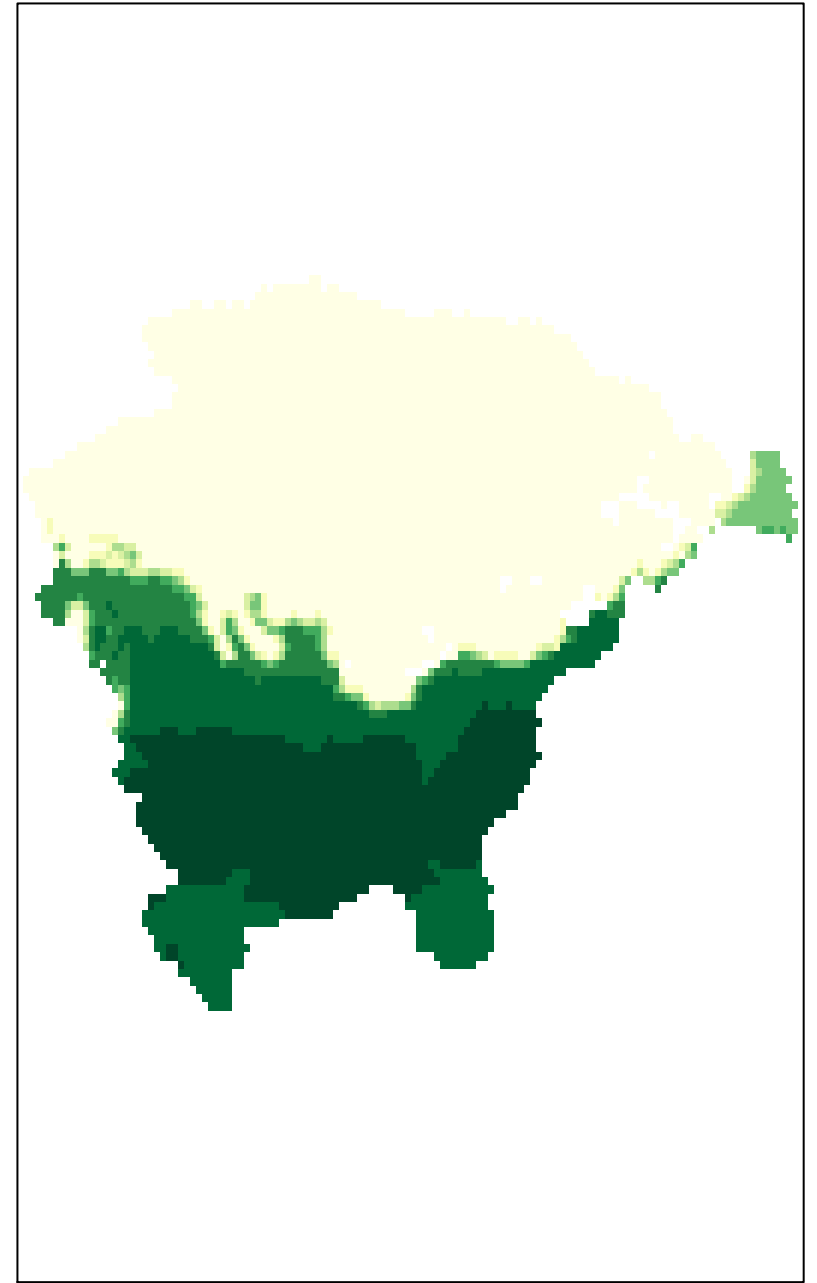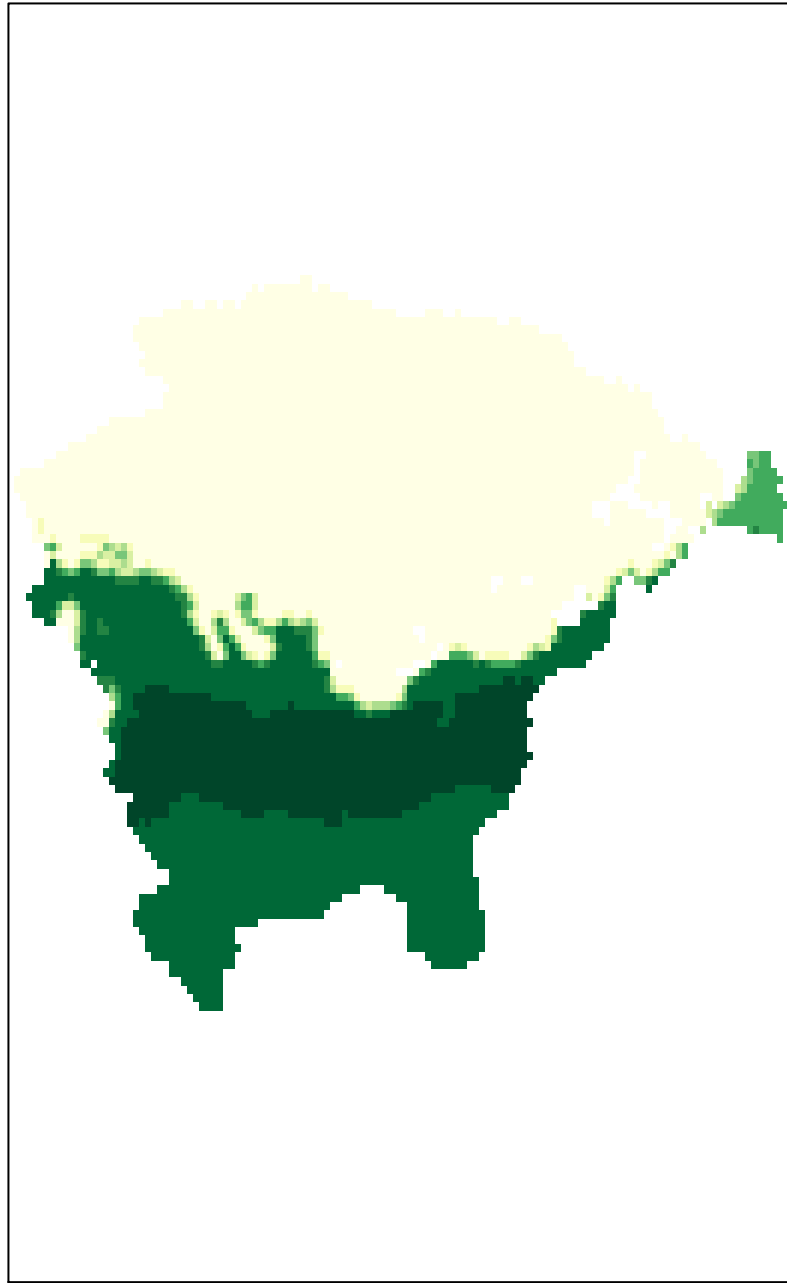

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X11000.ybp

SUM, X11000.ybp

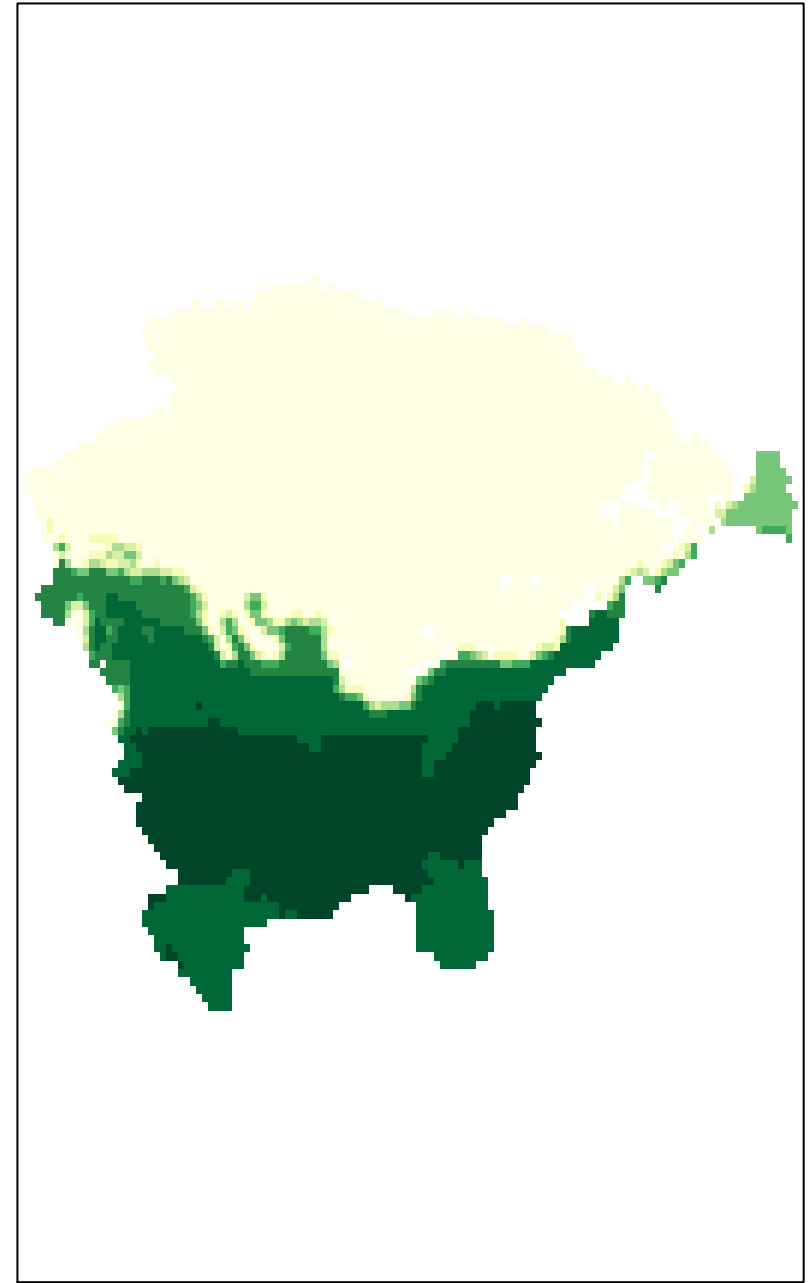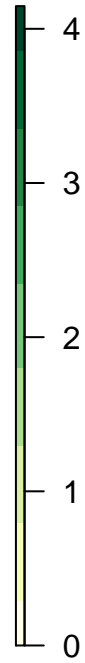
Species skipped = Fraxinus caroliniana, GCM = ecbilt

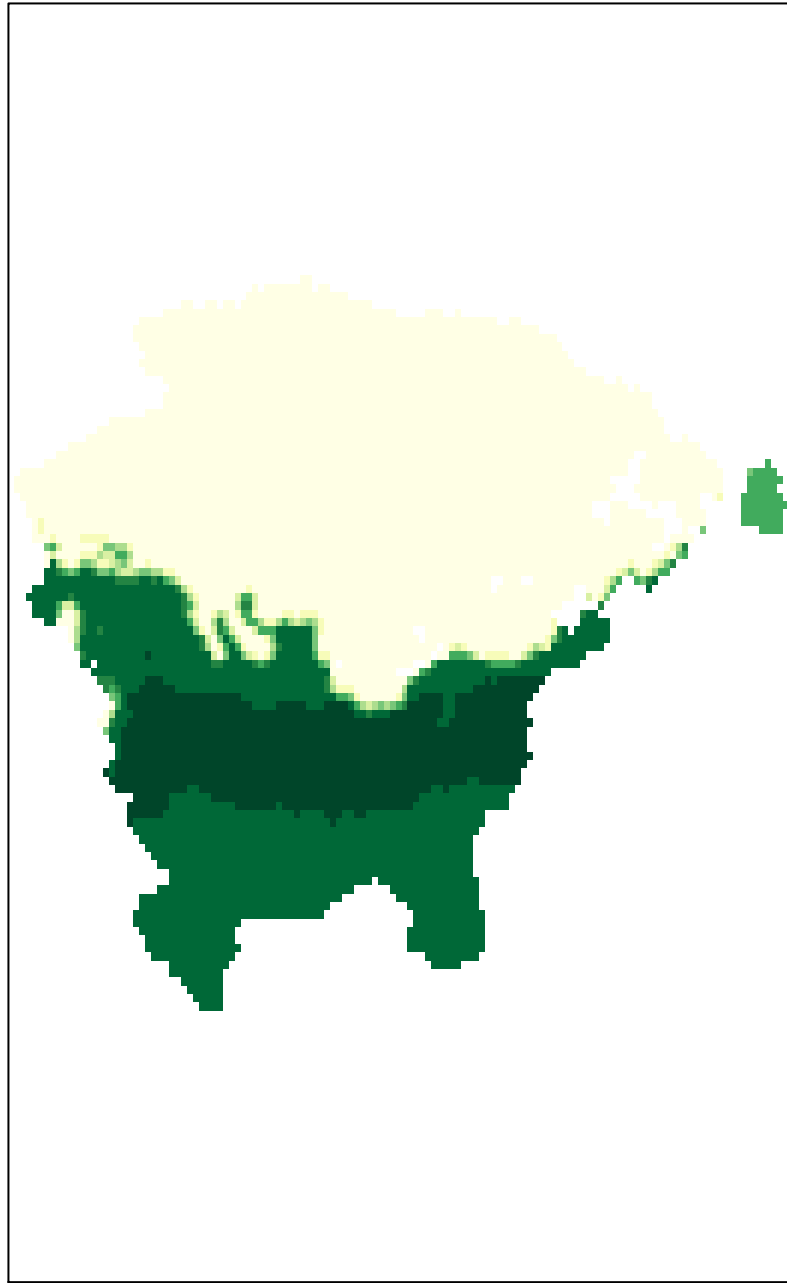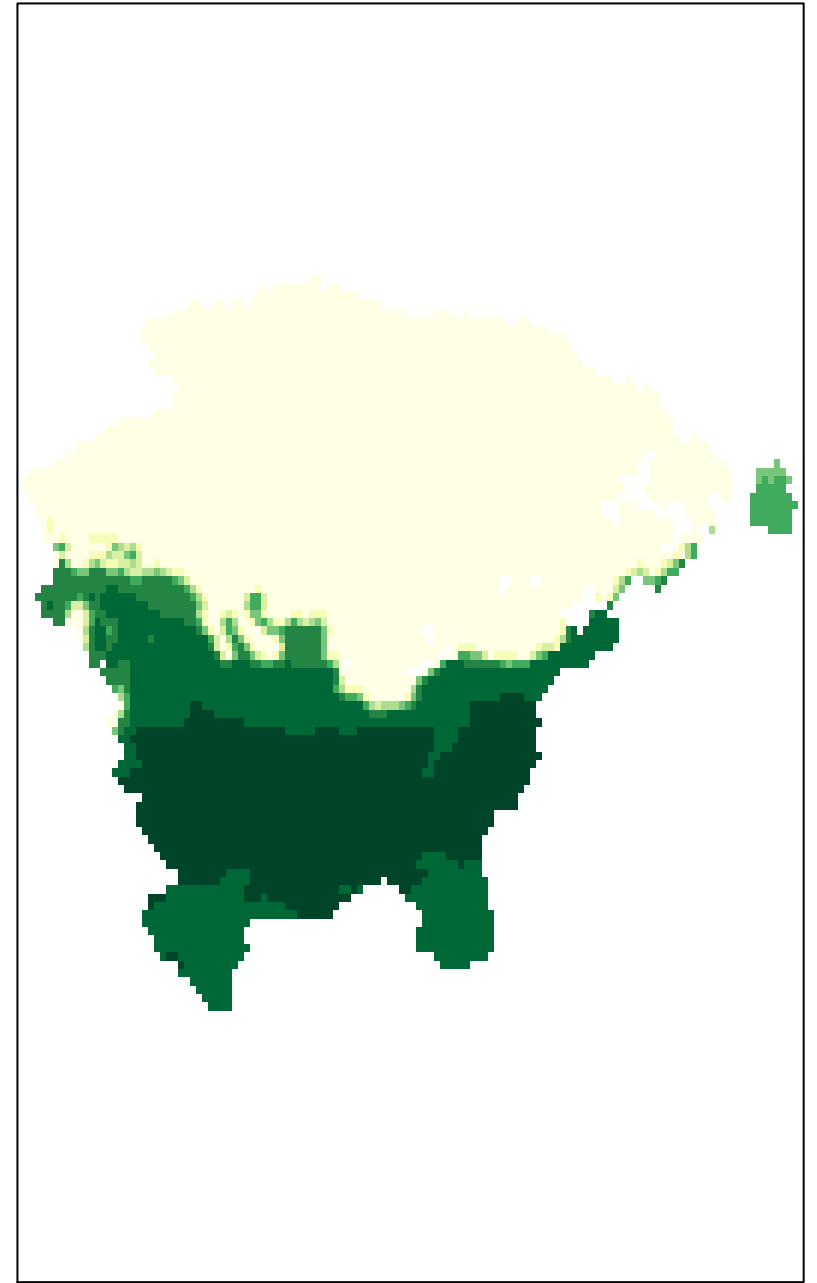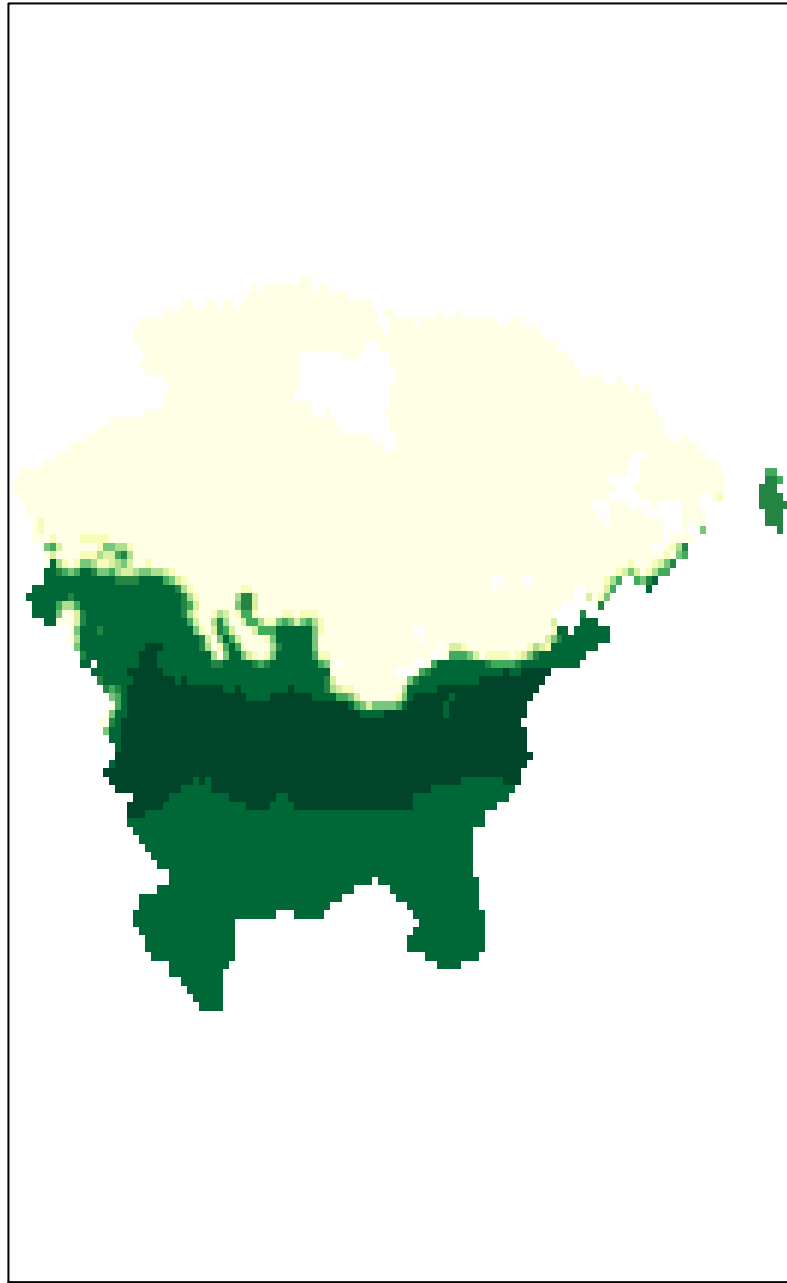**SUM, X10000.ybp** **SUM, X10000.ybp**

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X9000.ybp

SUM, X9000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

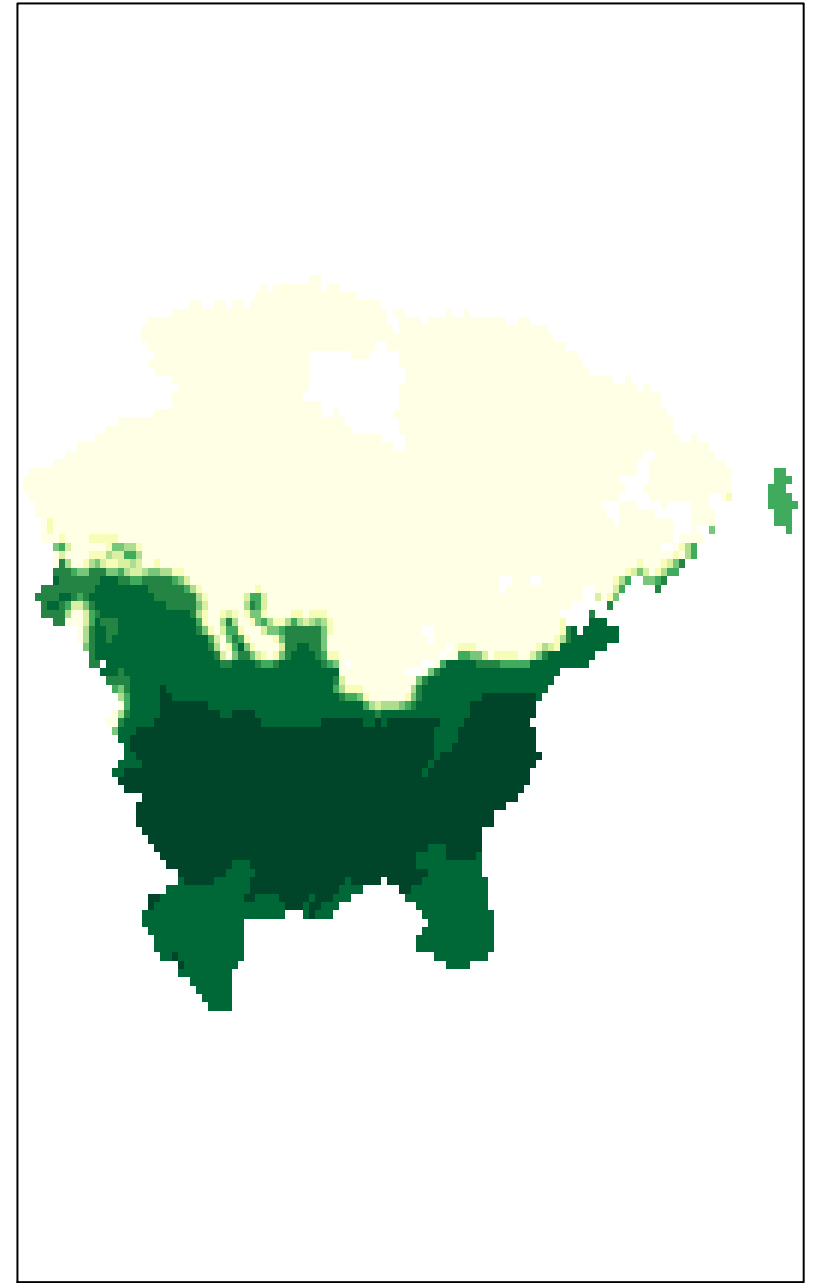**SUM, X8000.ybp**

SUM, X8000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X7000.ybp

SUM, X7000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt
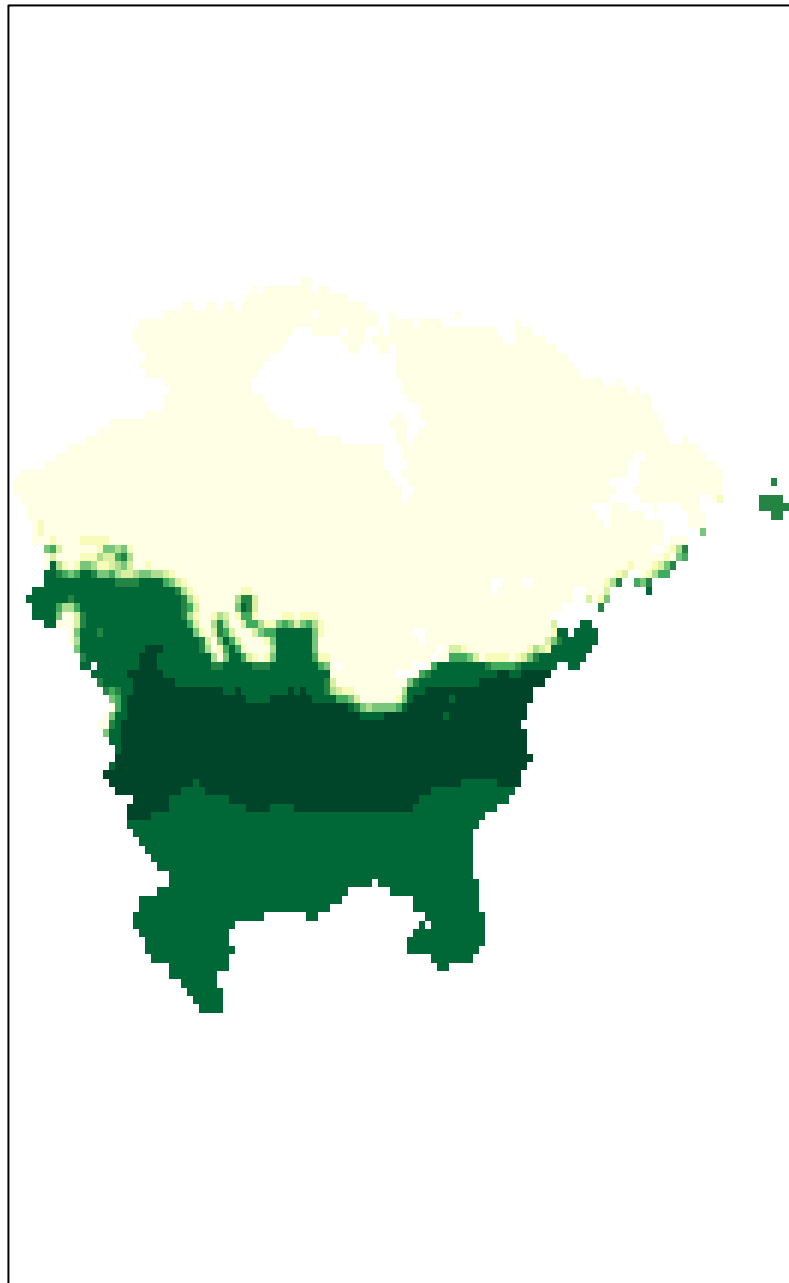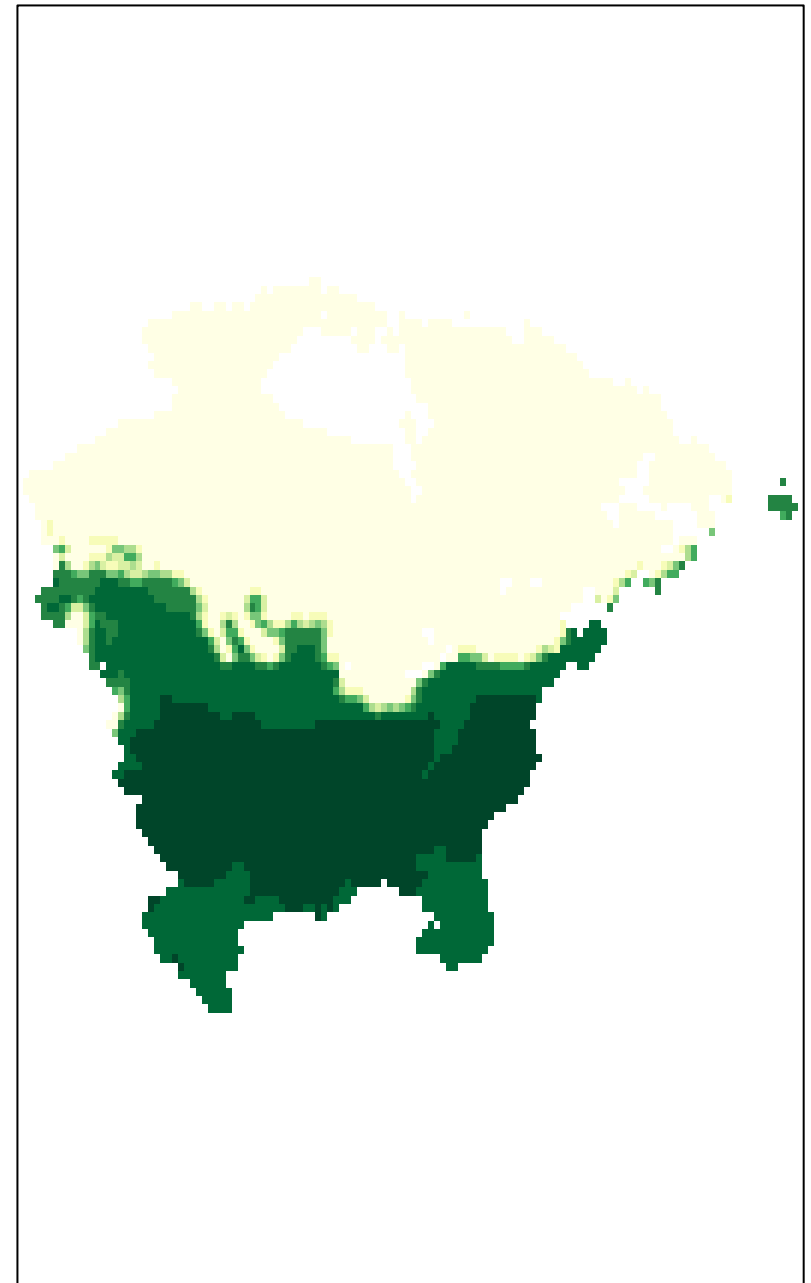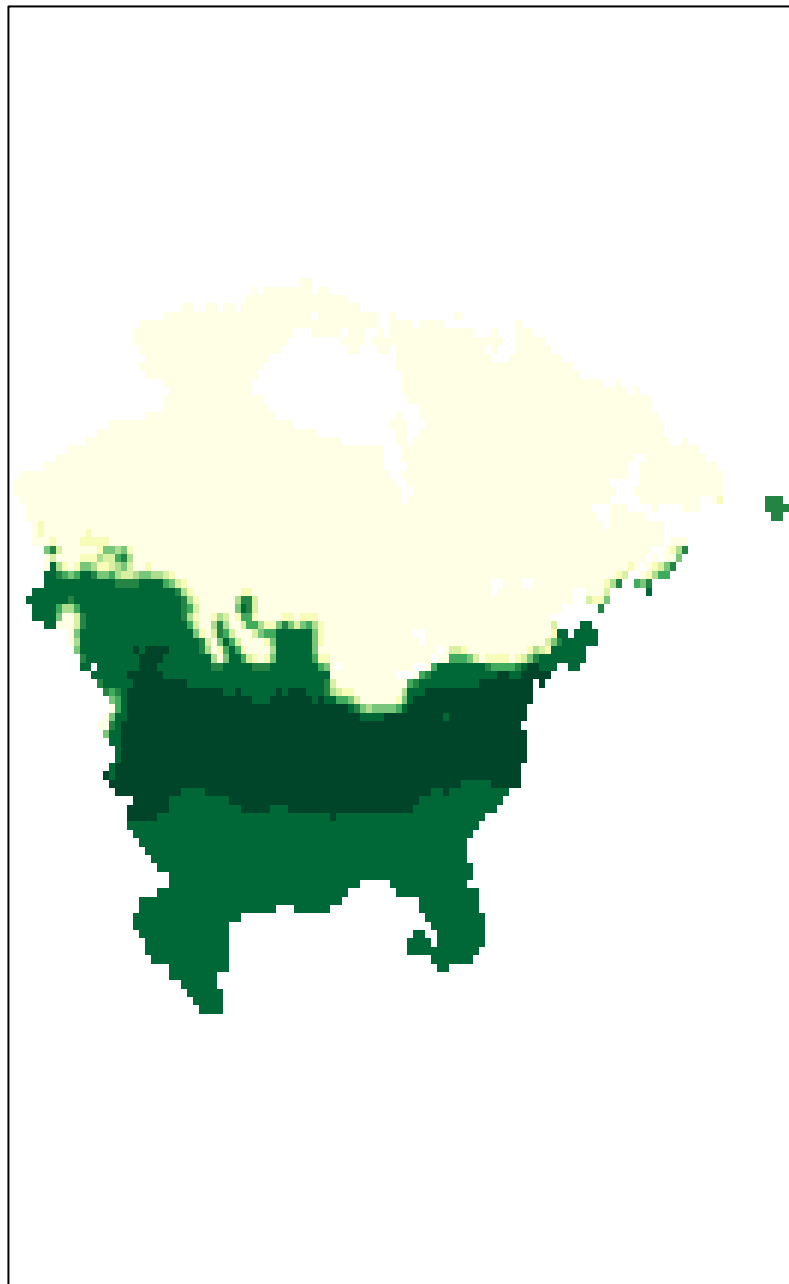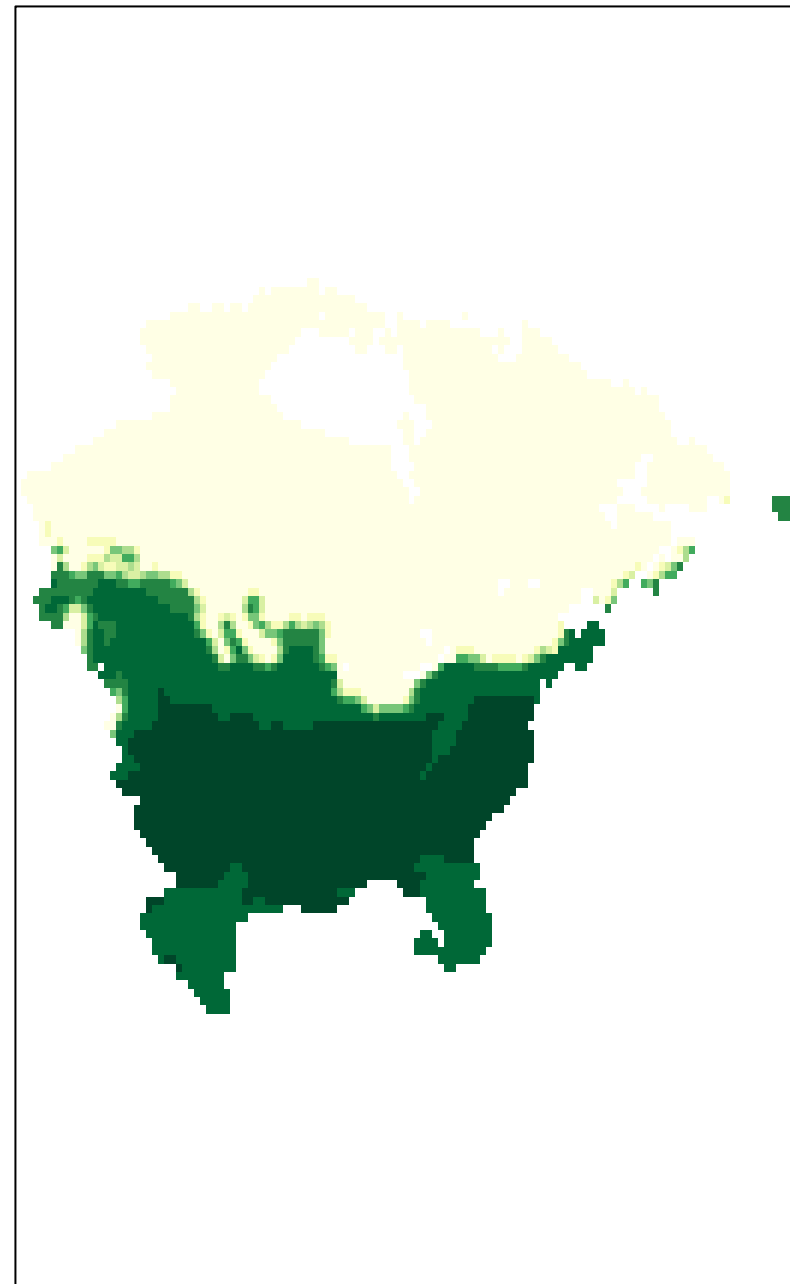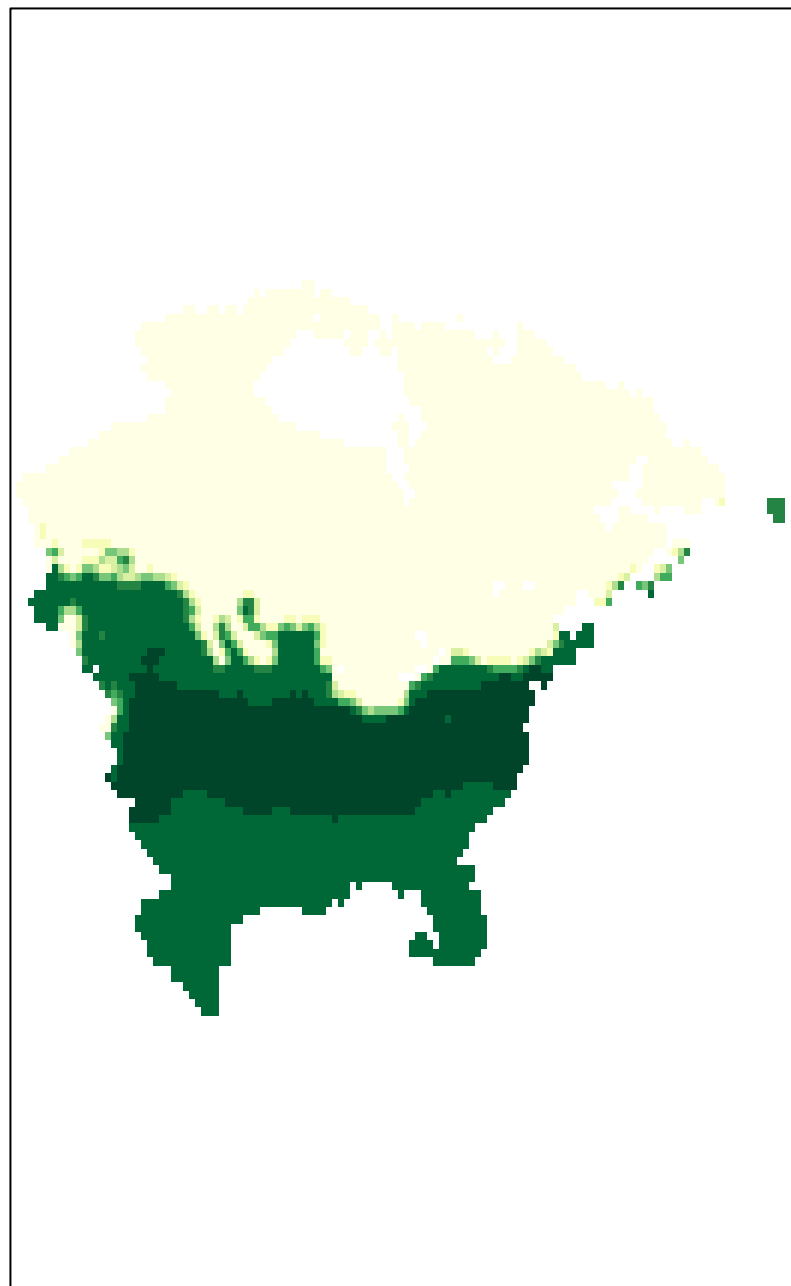
SUM, X6000.ybp
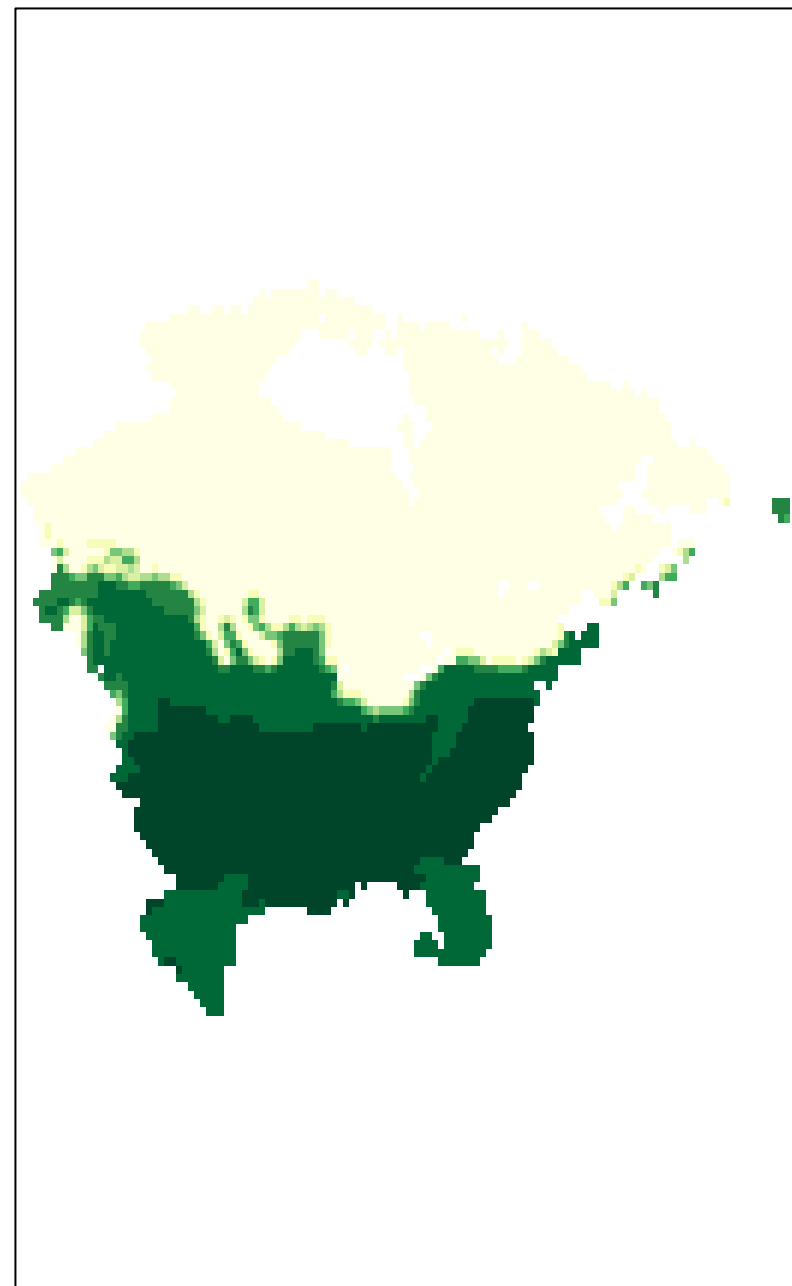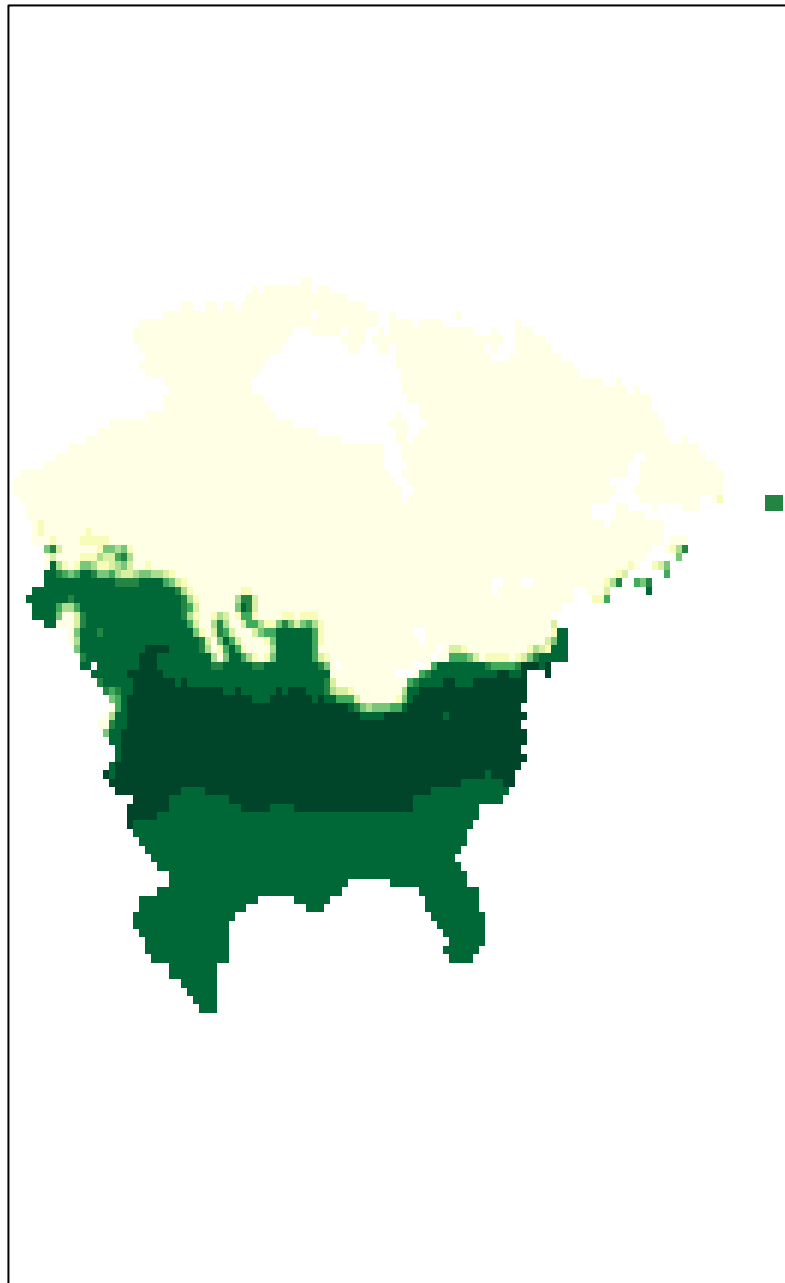
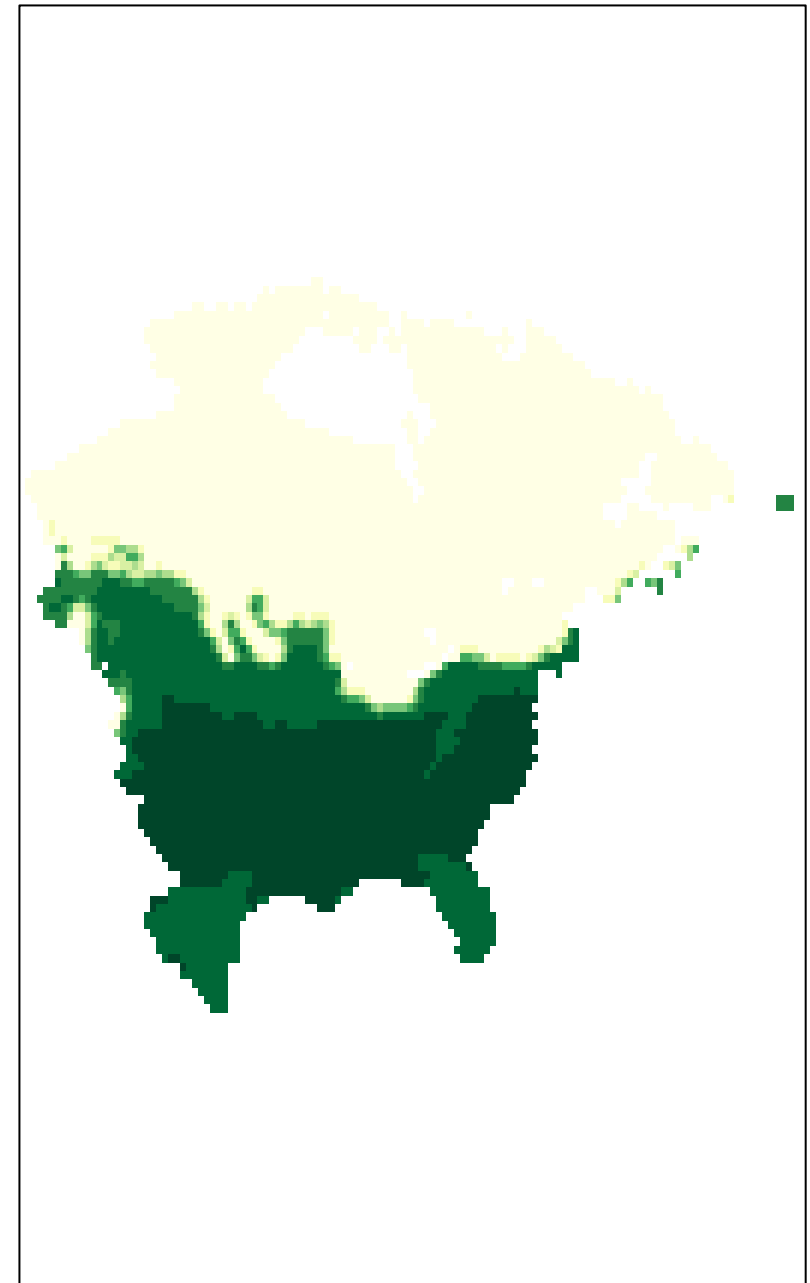**SUM, X6000.ybp**

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X5000.ybp

SUM, X5000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt
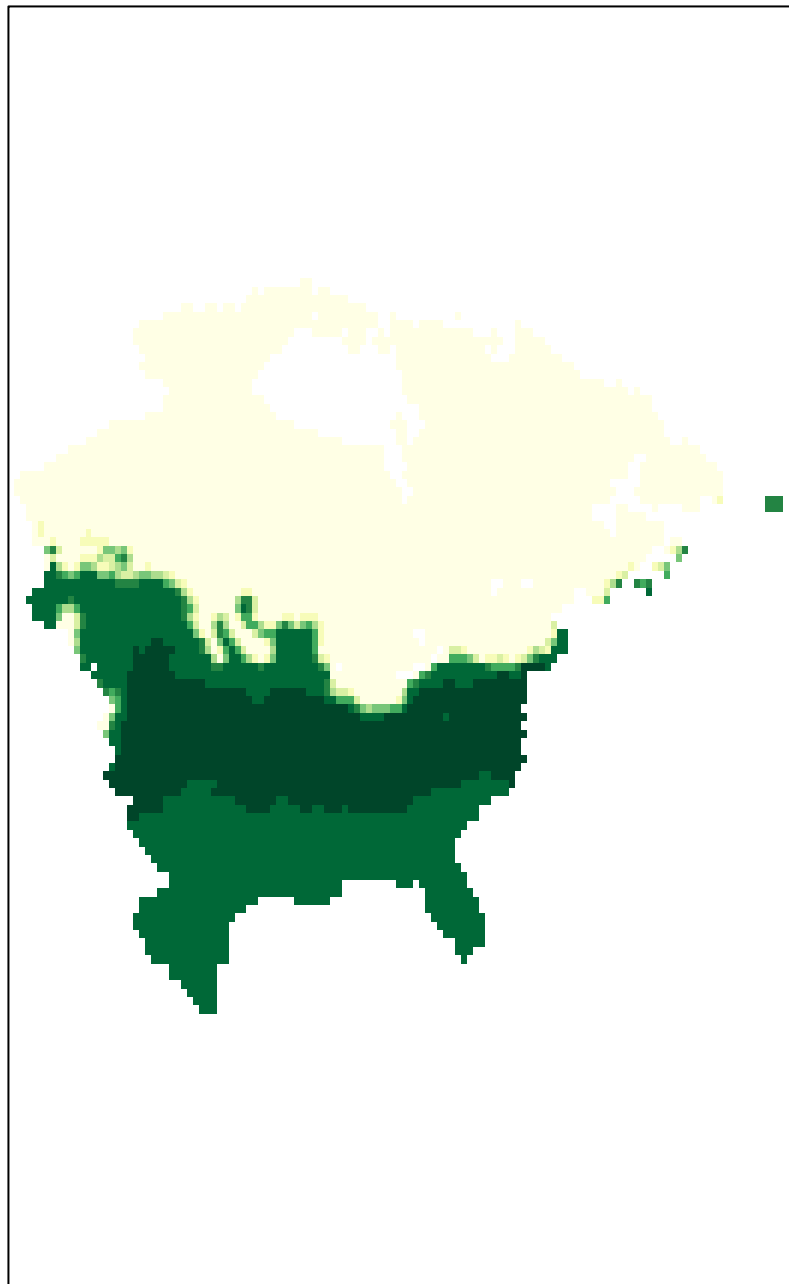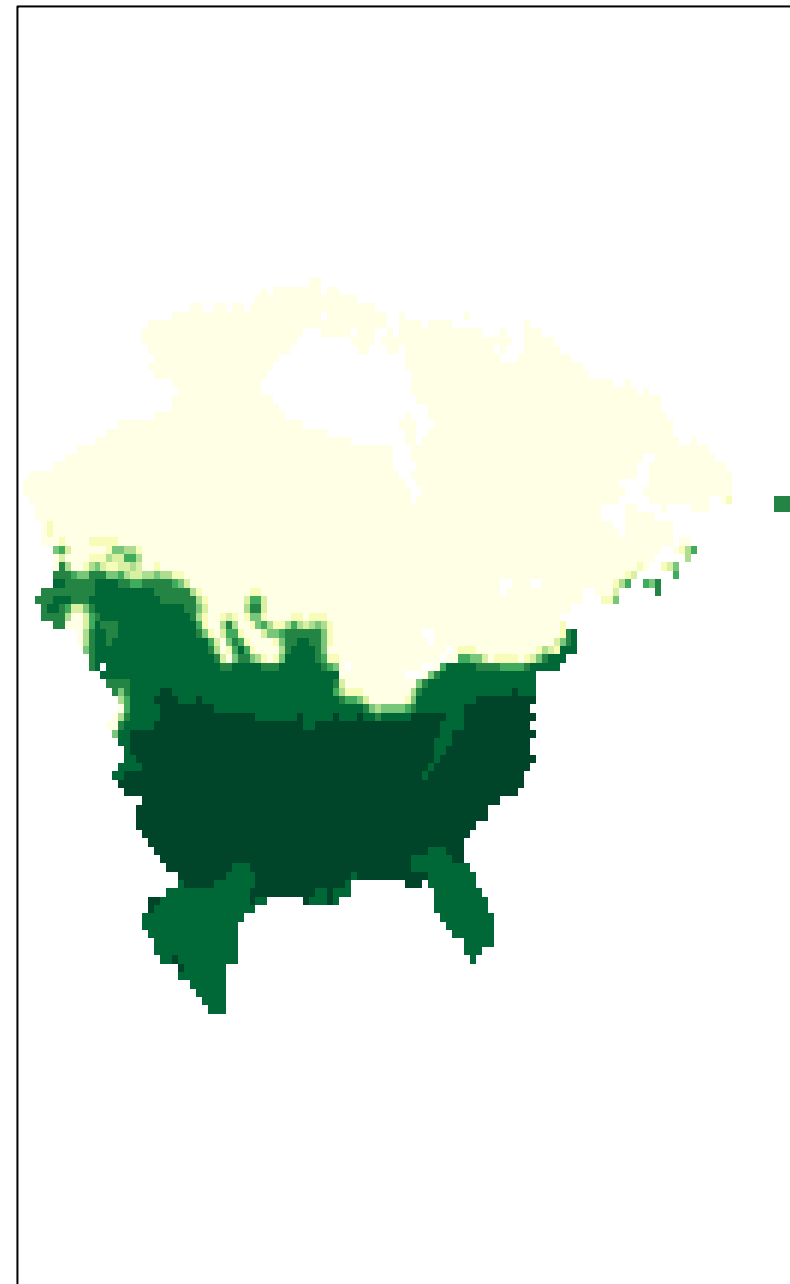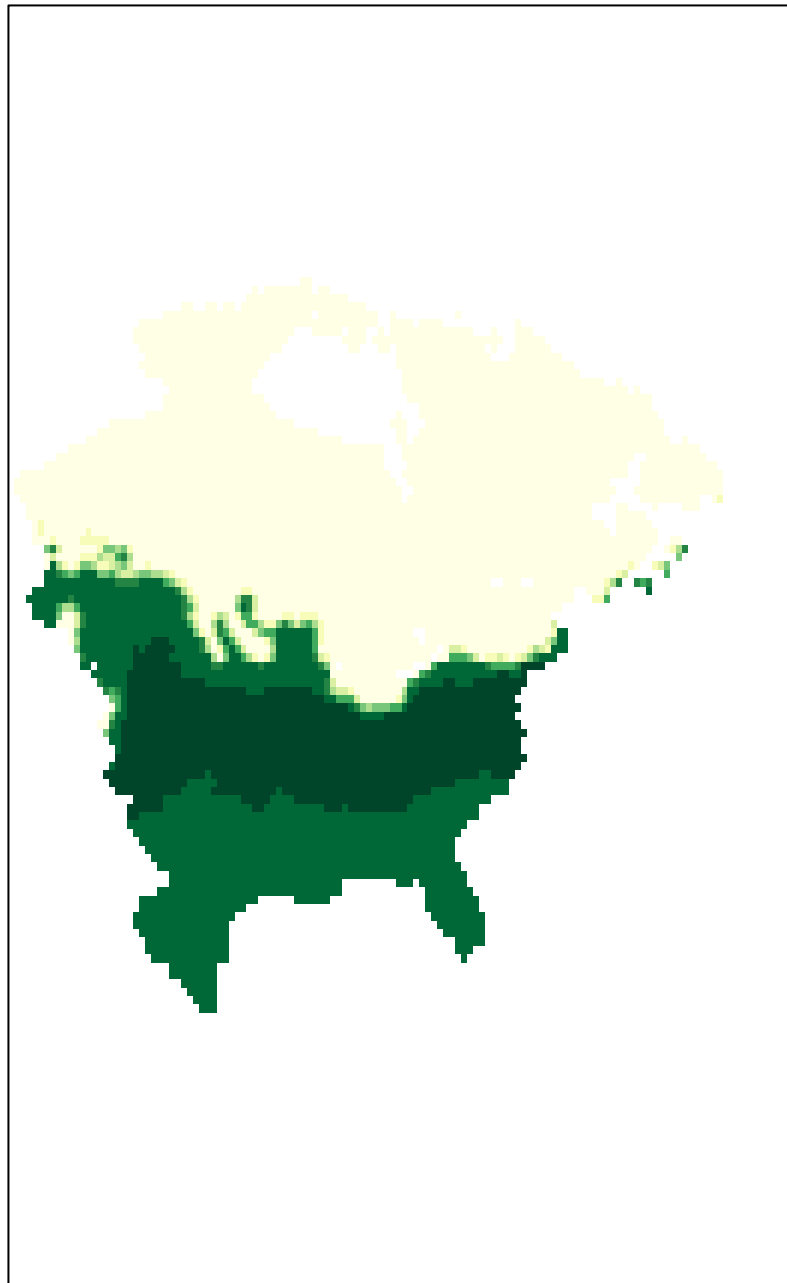
SUM, X4000.ybp

SUM, X4000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X3000.ybp
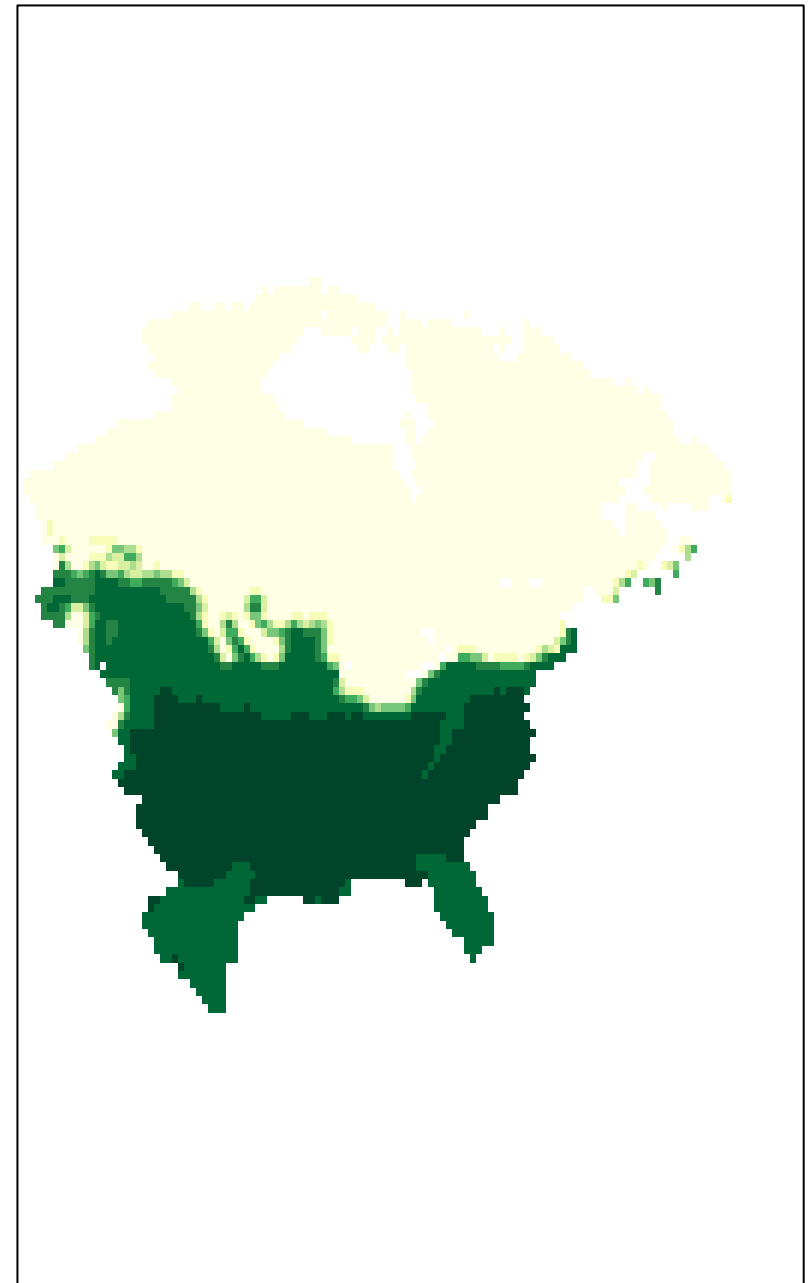
SUM, X3000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X2000.ybp

SUM, X2000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt

SUM, X1000.ybp

SUM, X1000.ybp

Species skipped = Fraxinus caroliniana, GCM = ecbilt
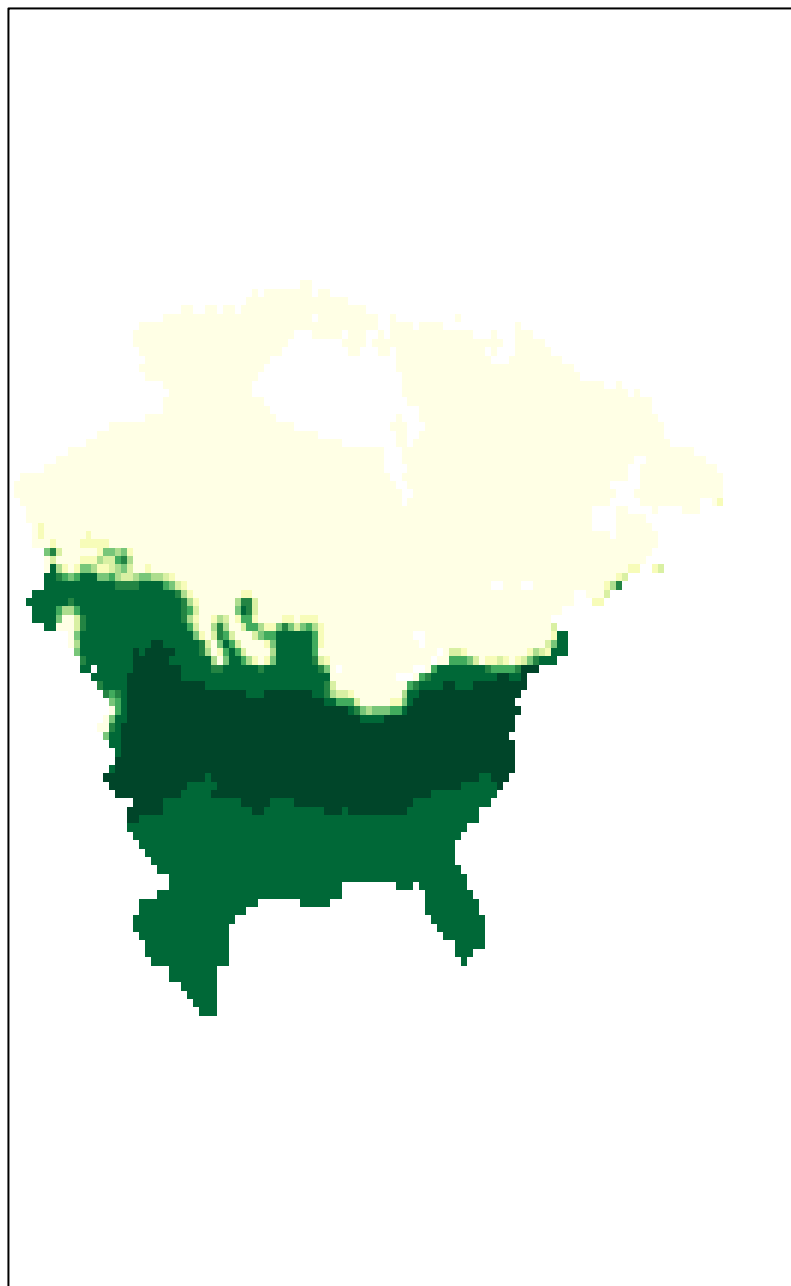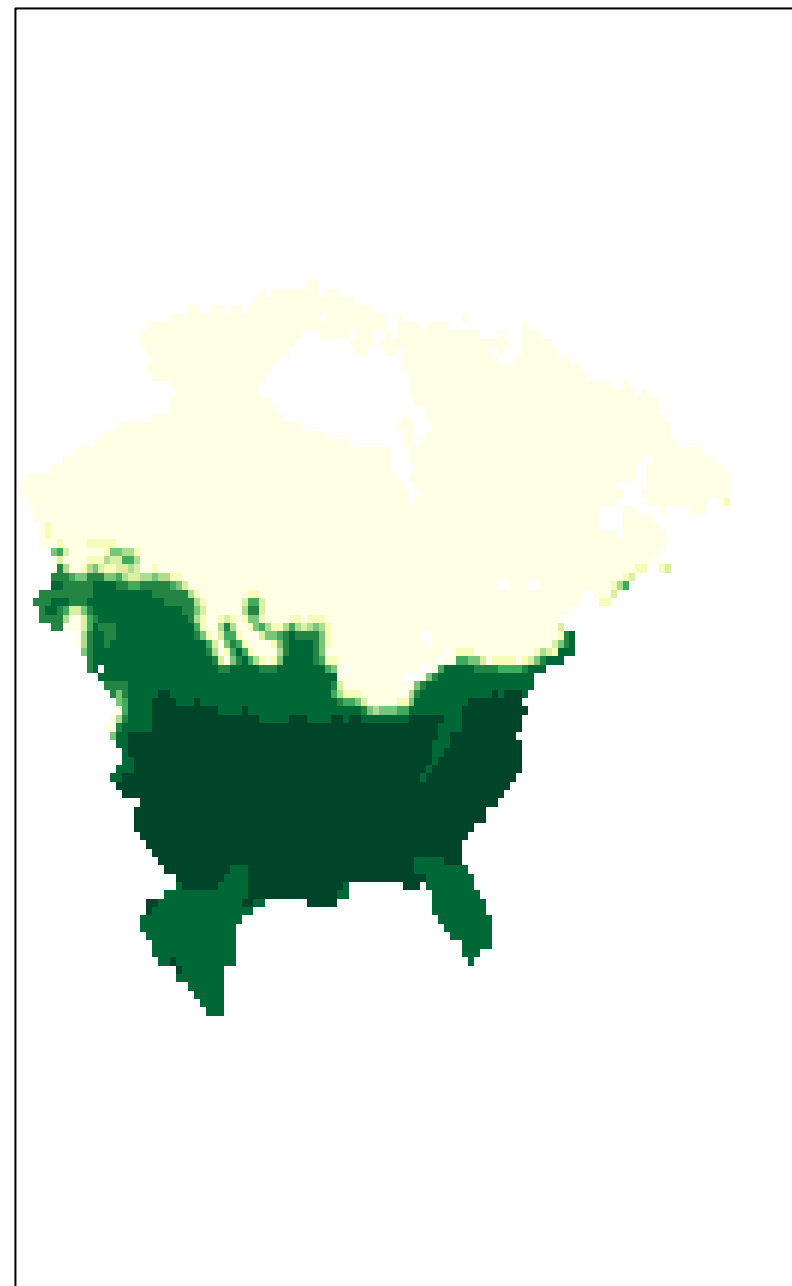
SUM, X0.ybp   **SUM, X0.ybp**

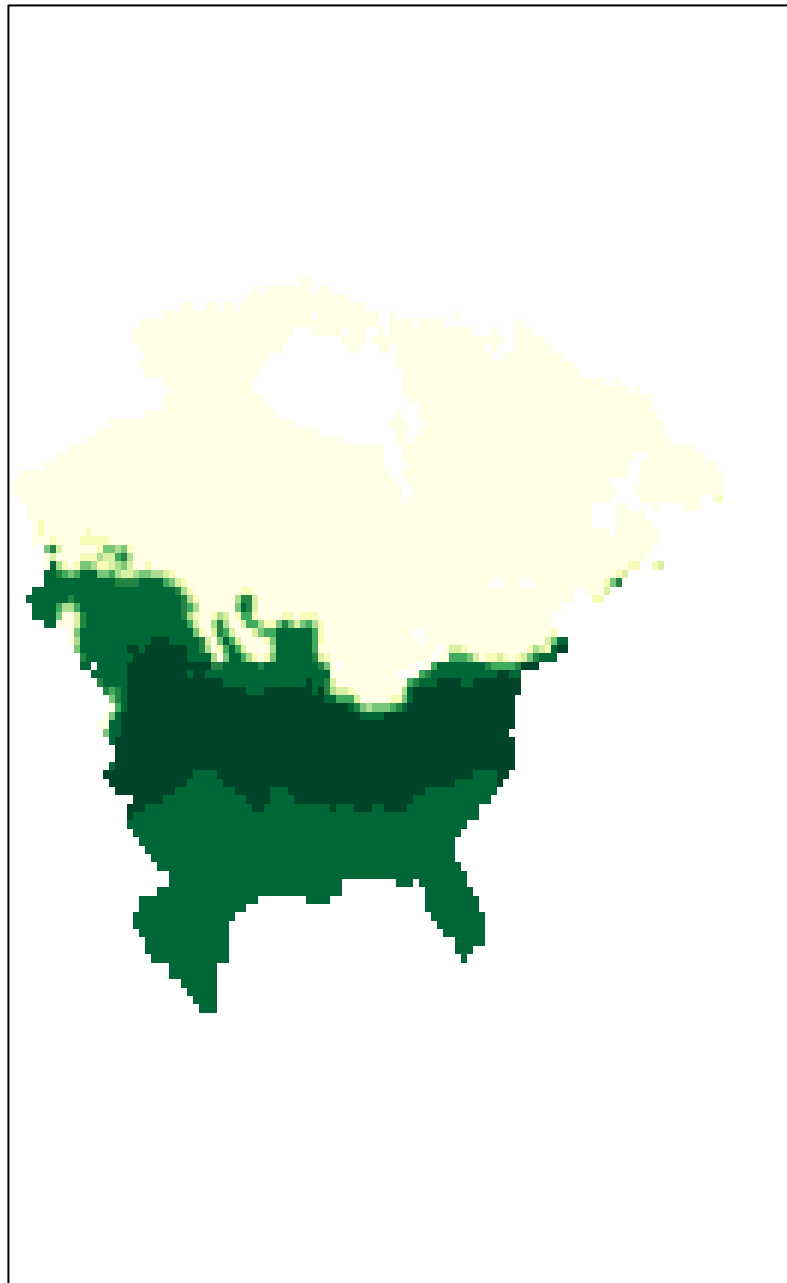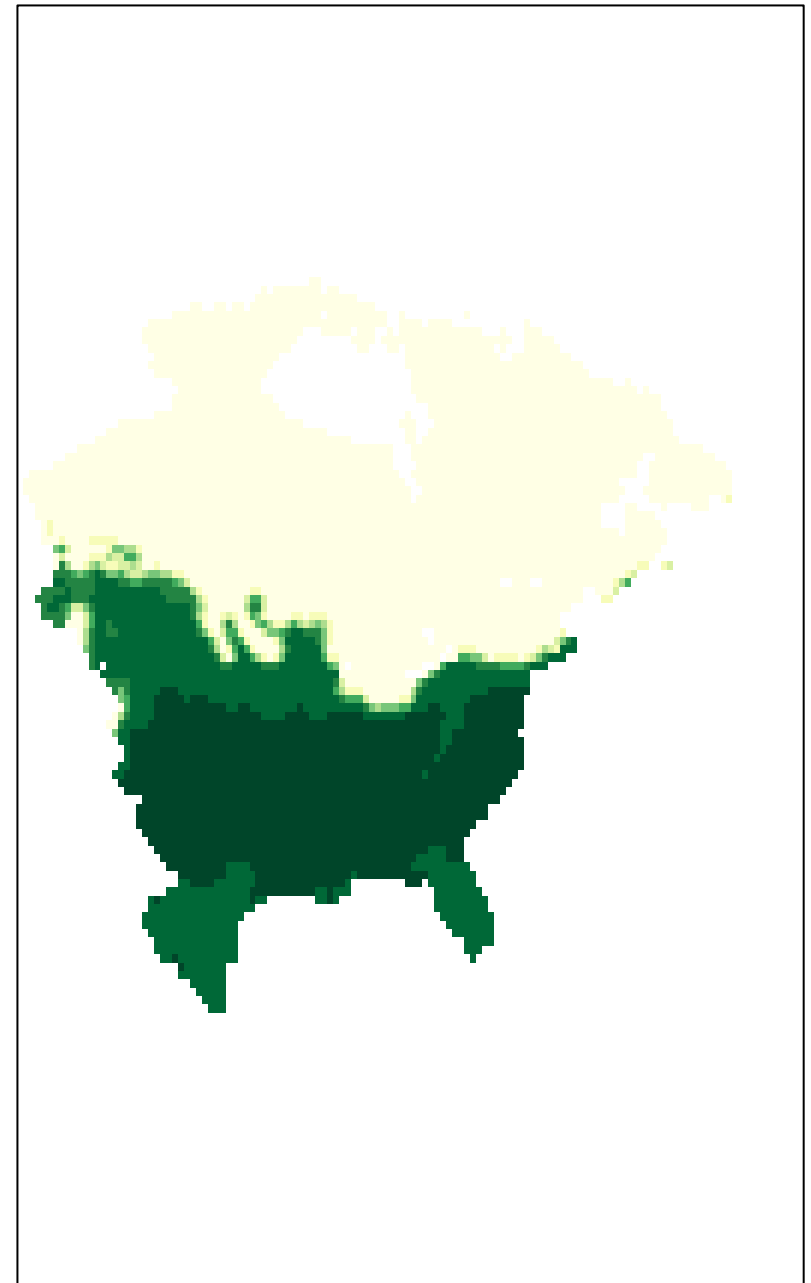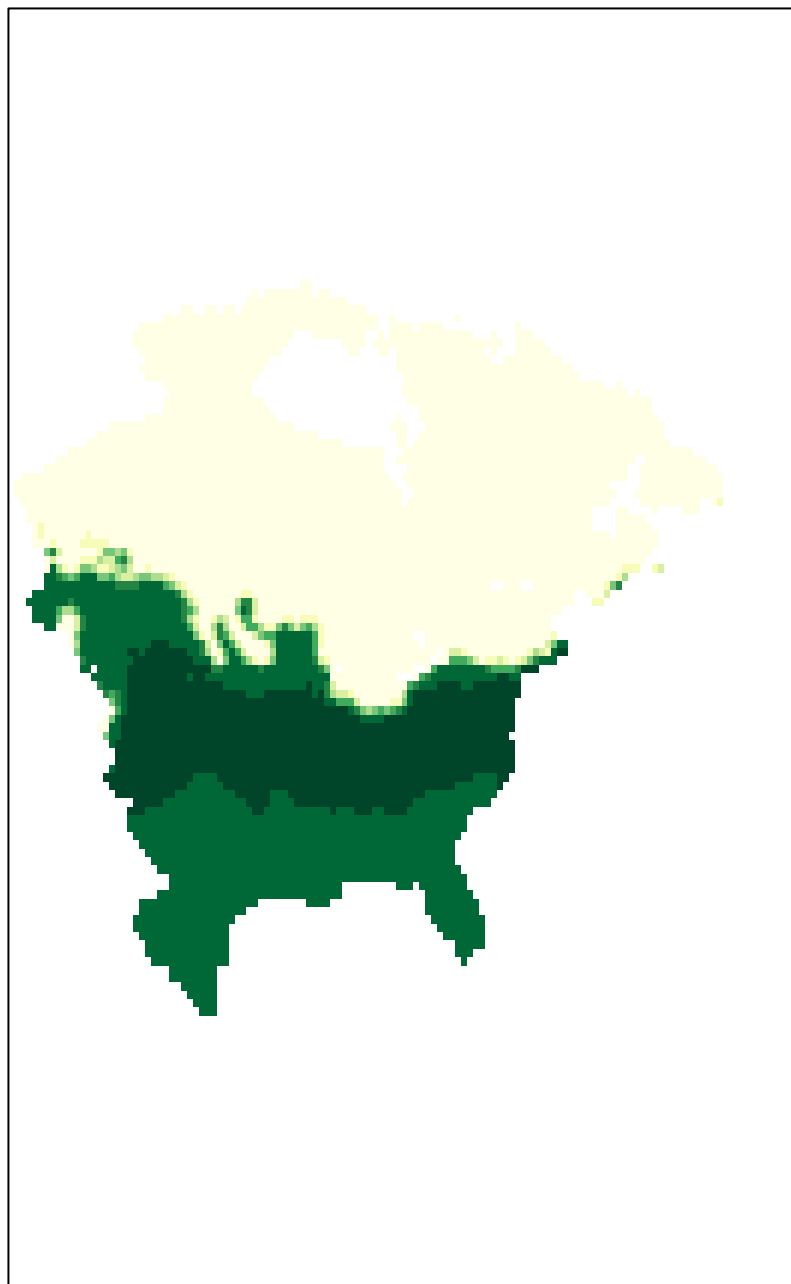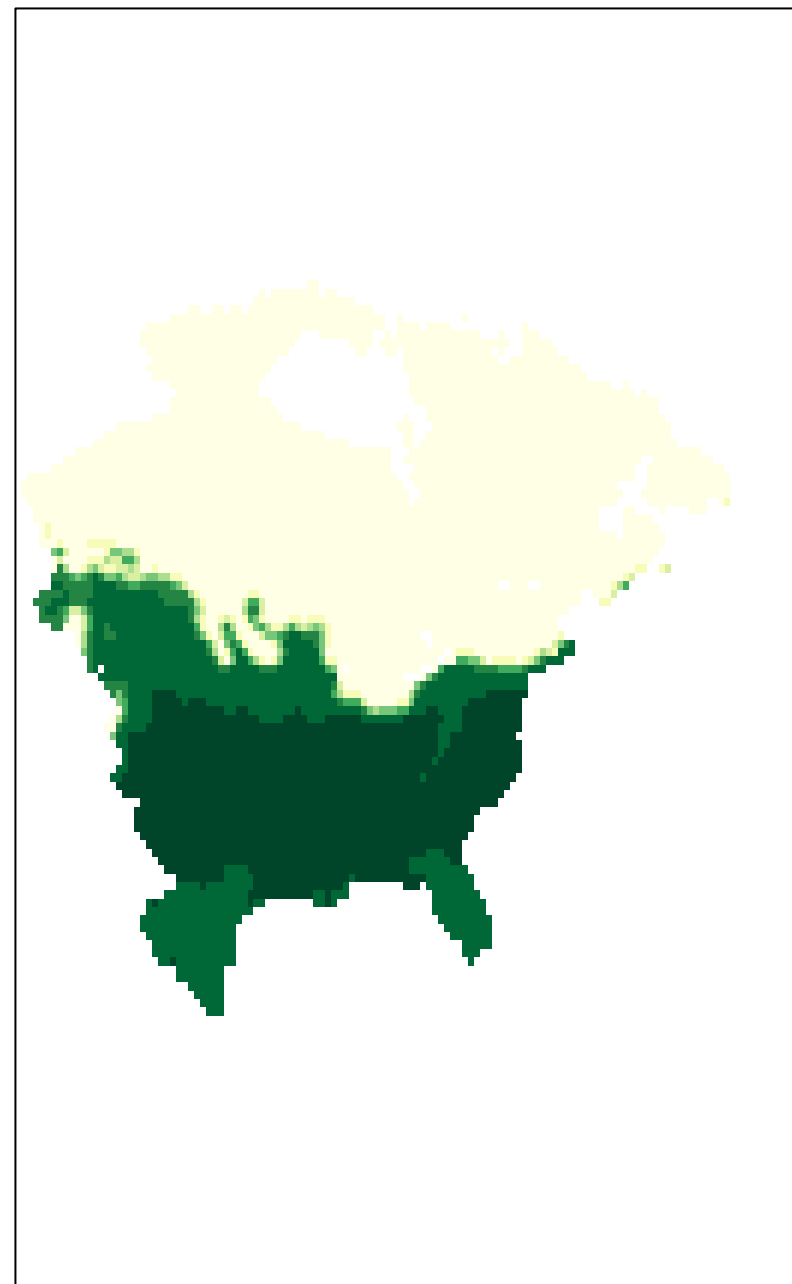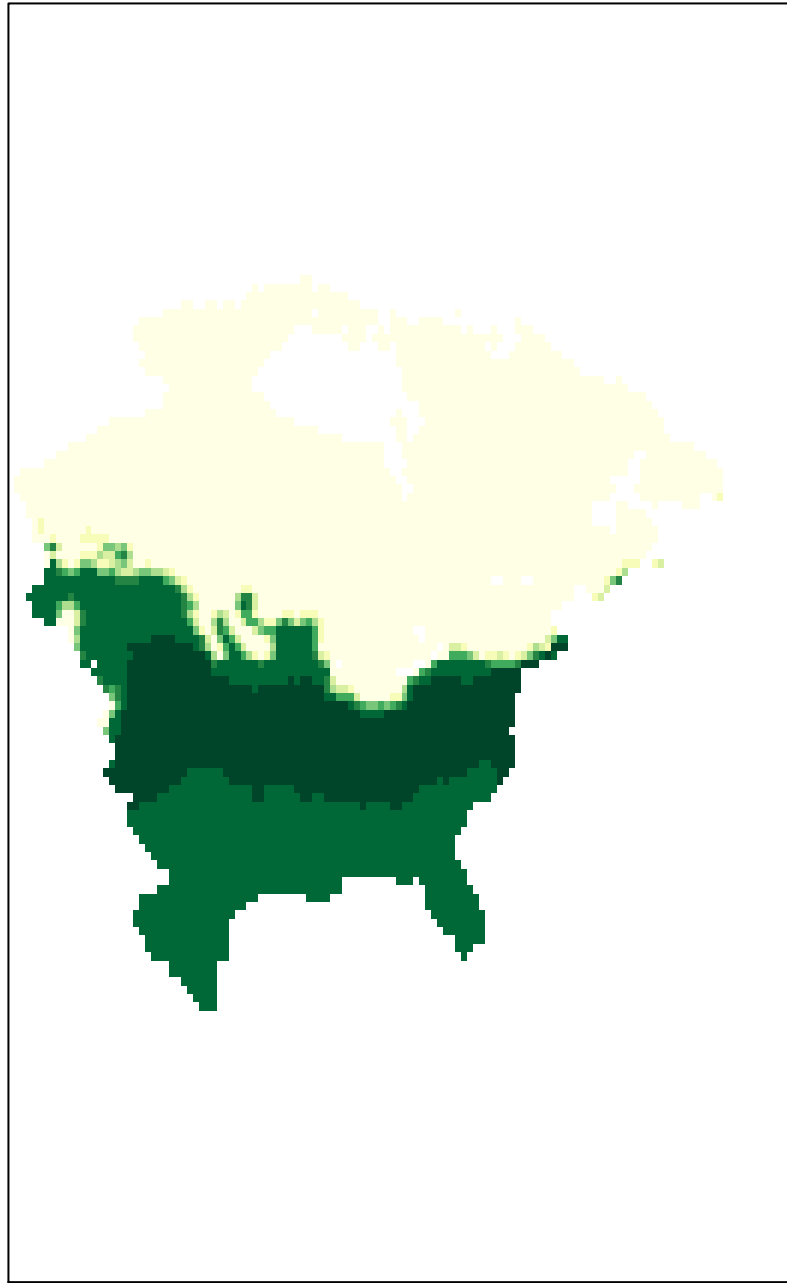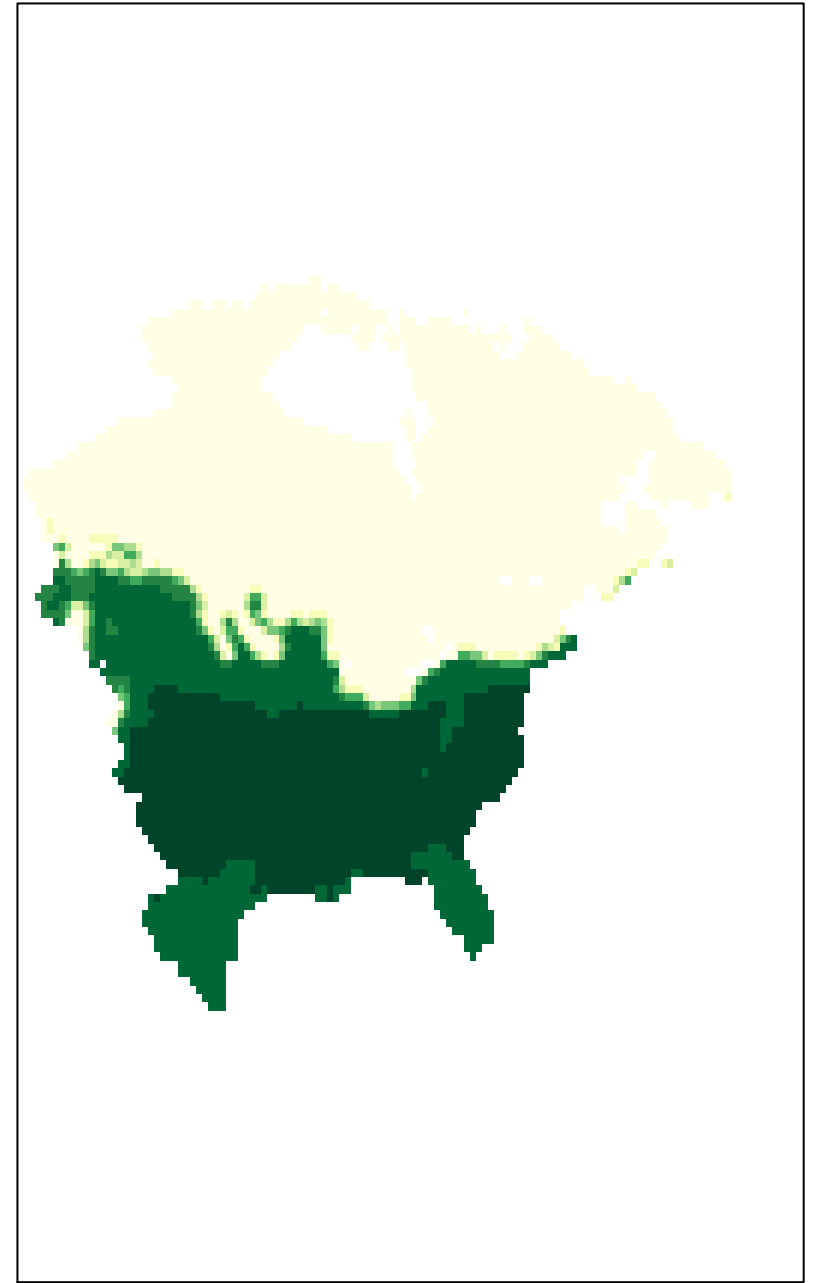Species skipped = Fraxinus caroliniana, GCM = ecbilt