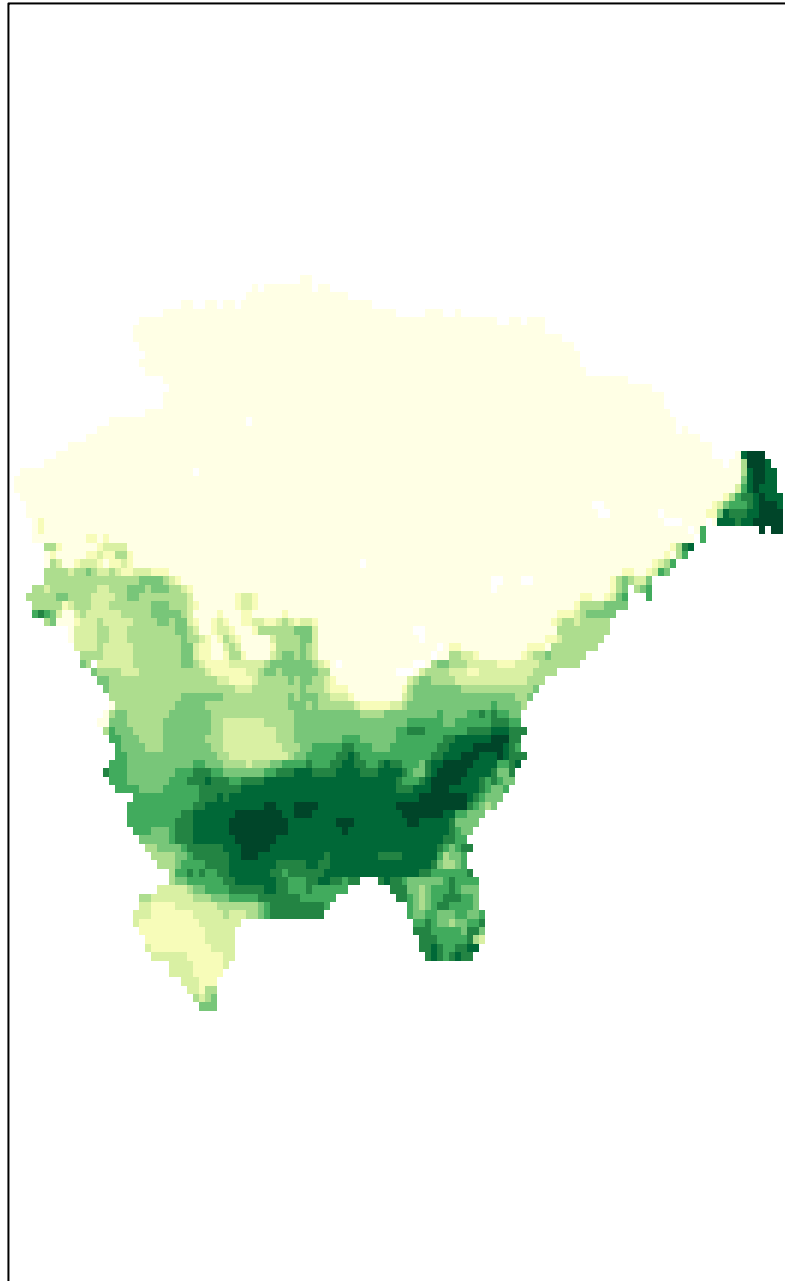
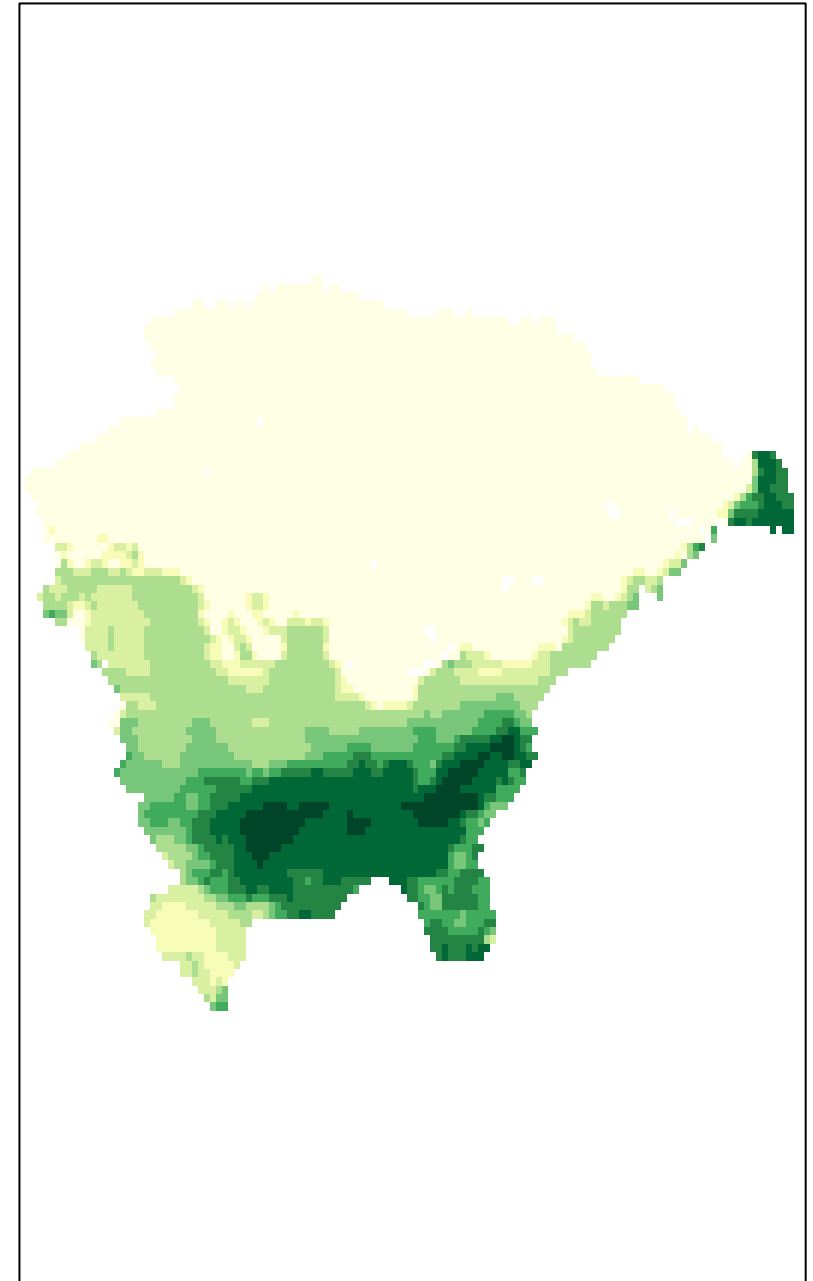


SUM, X21000.ybp

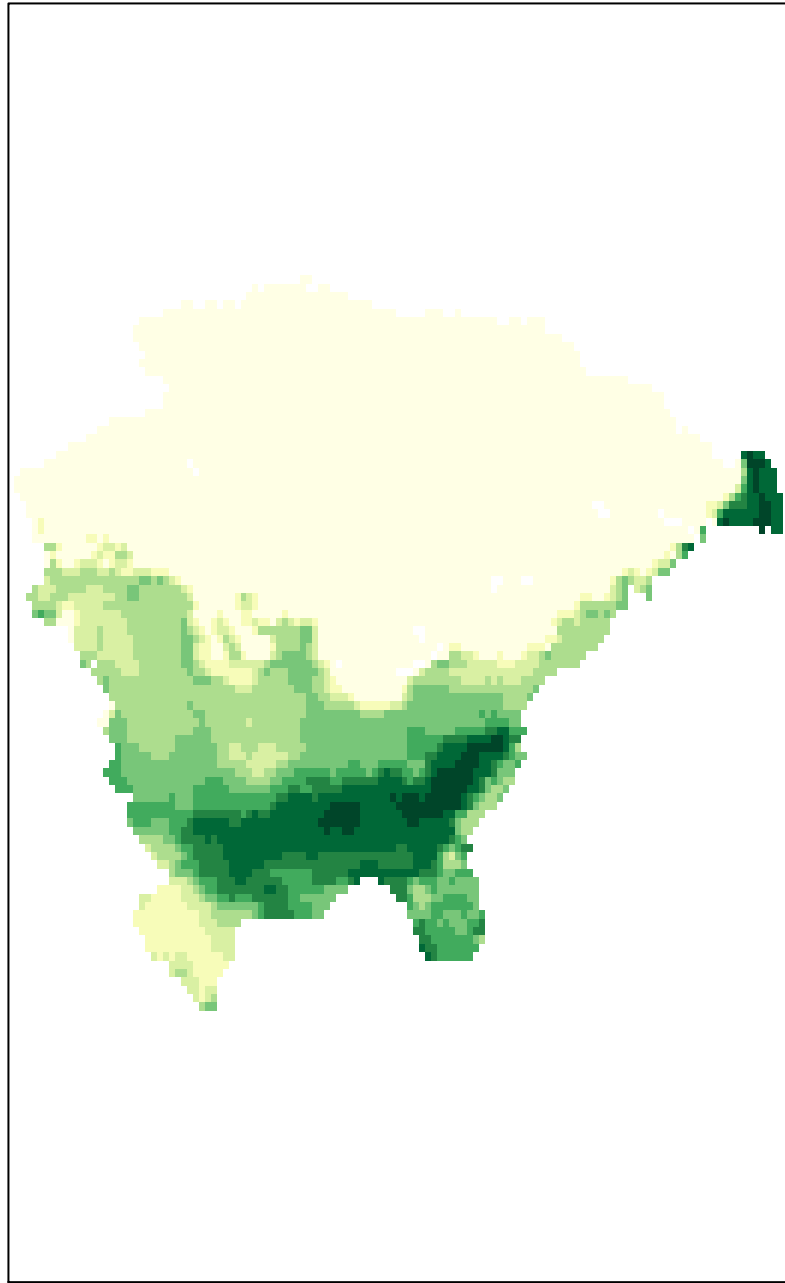


SUM, X21000.ybp

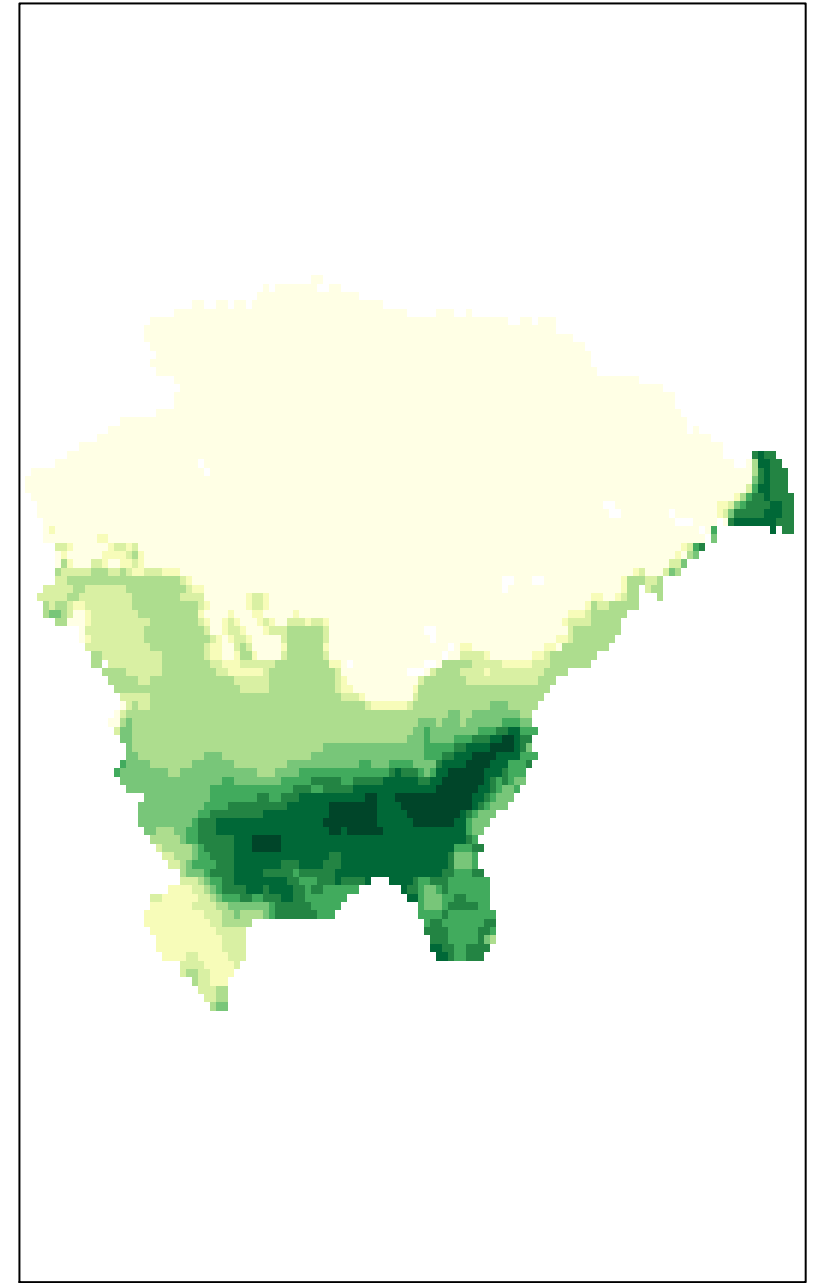


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X20000.ybp

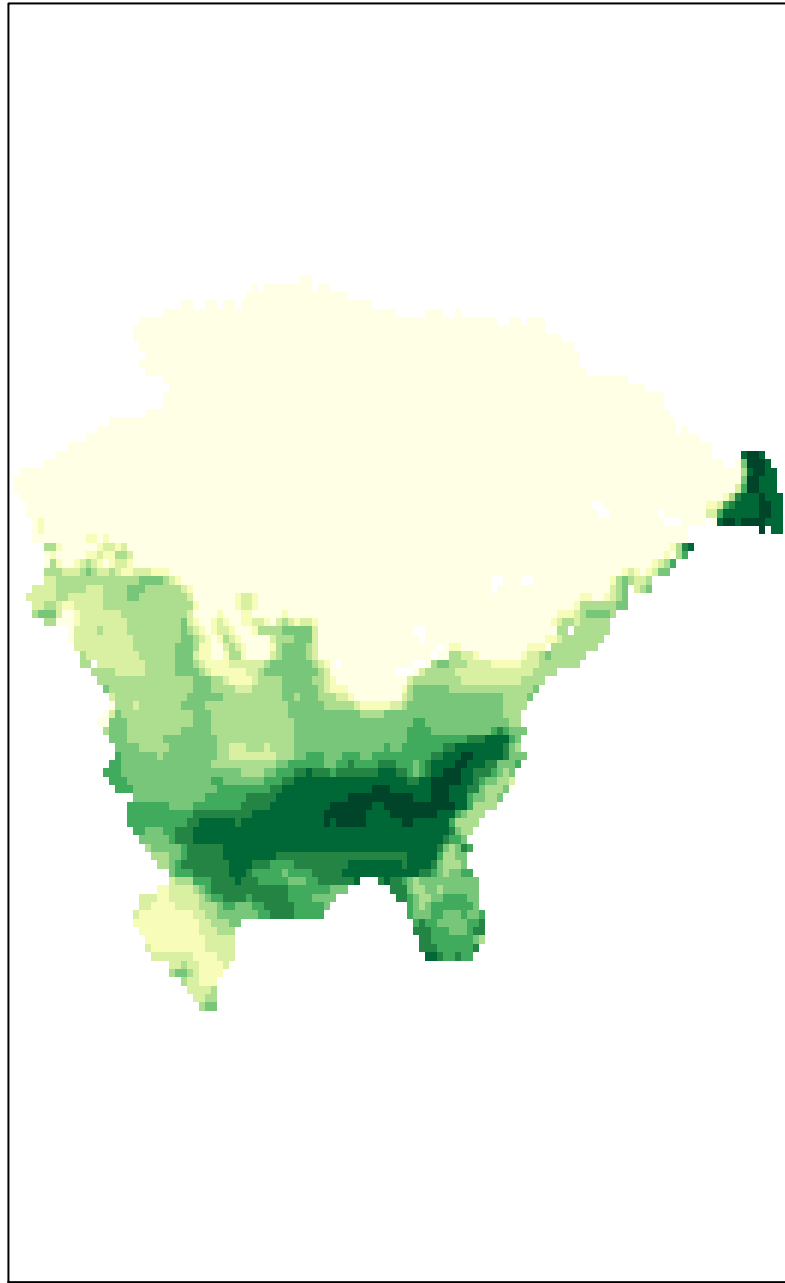


SUM, X20000.ybp

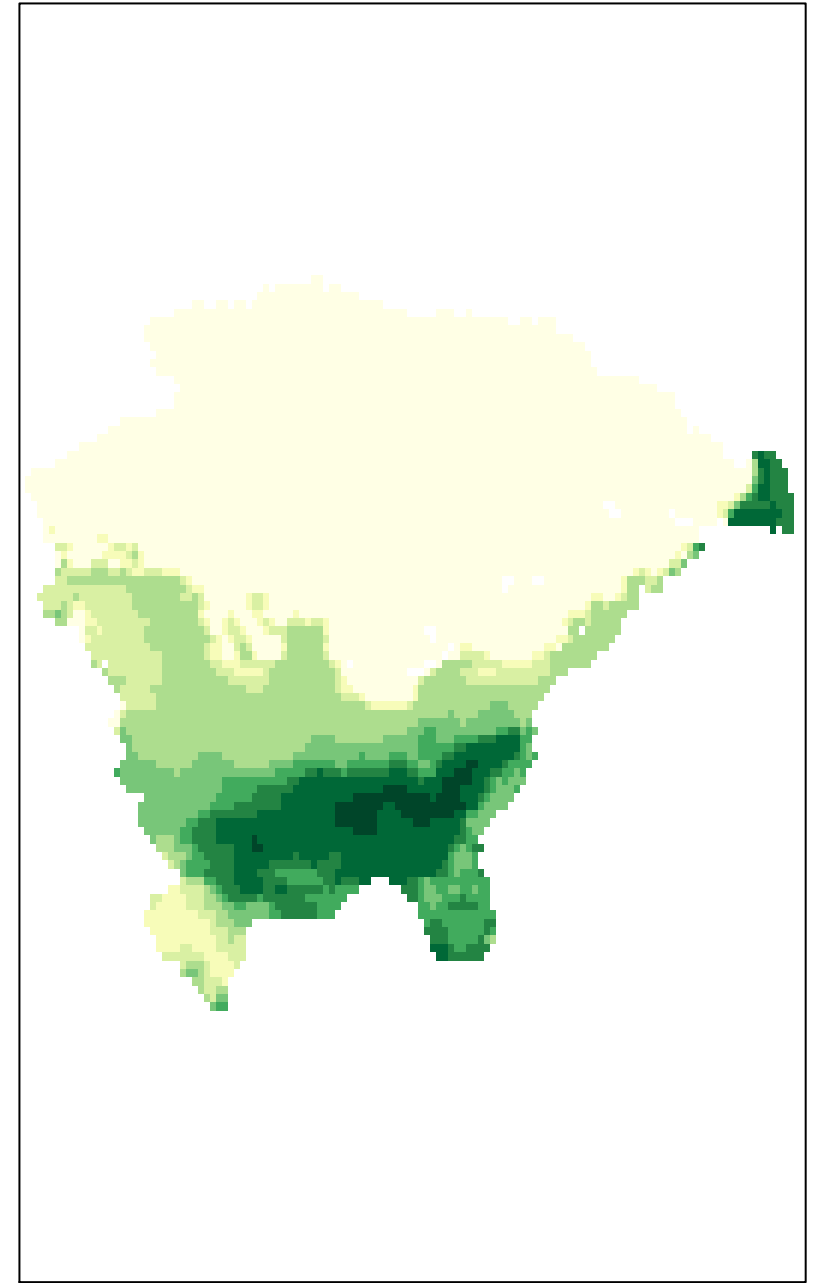


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X19000.ybp

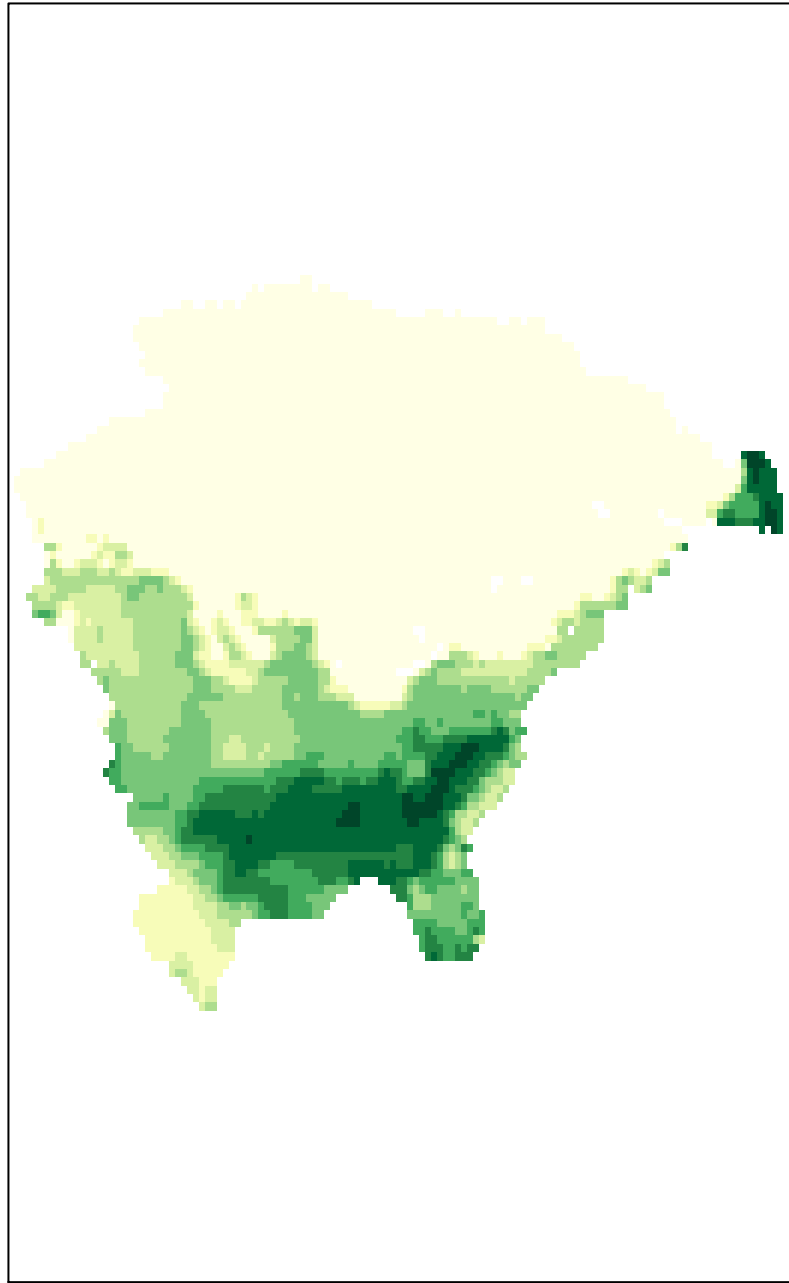


SUM, X19000.ybp

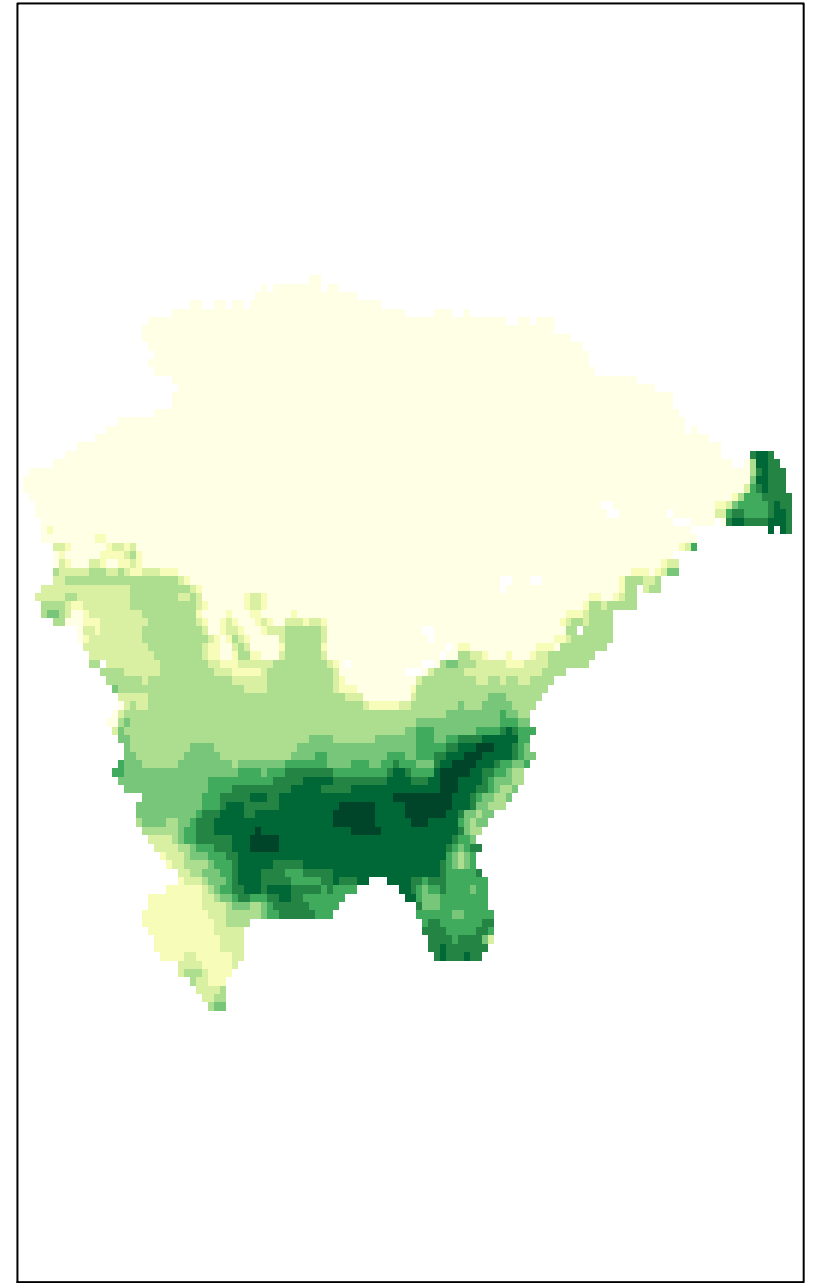


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X18000.ybp

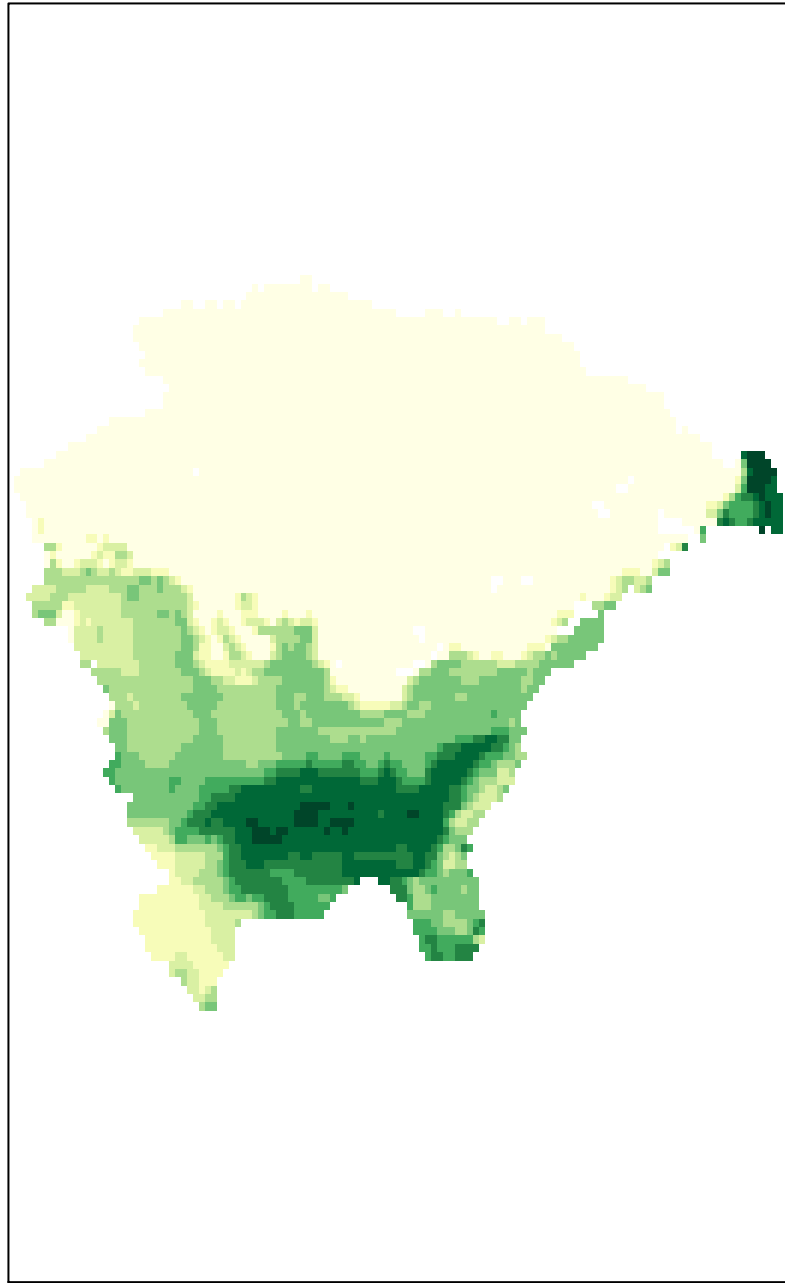


SUM, X18000.ybp

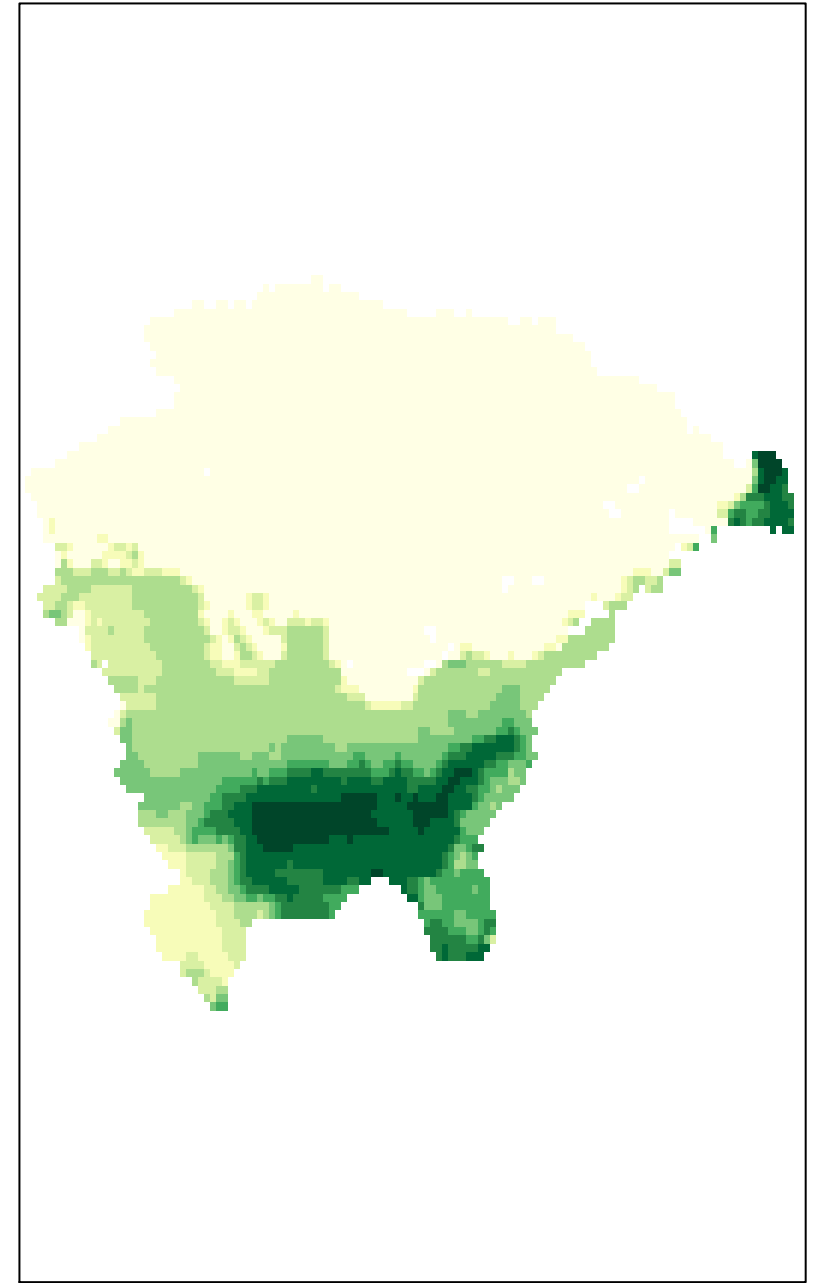


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X17000.ybp

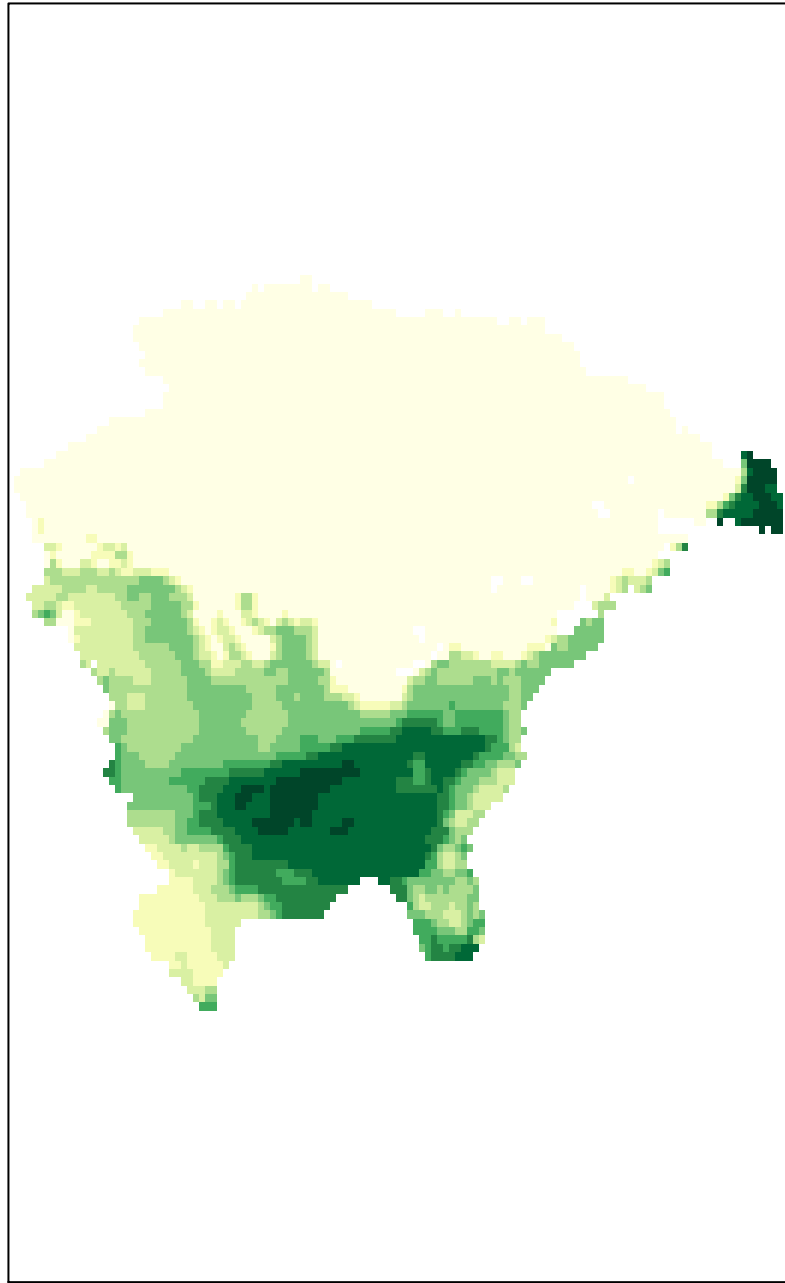


SUM, X17000.ybp

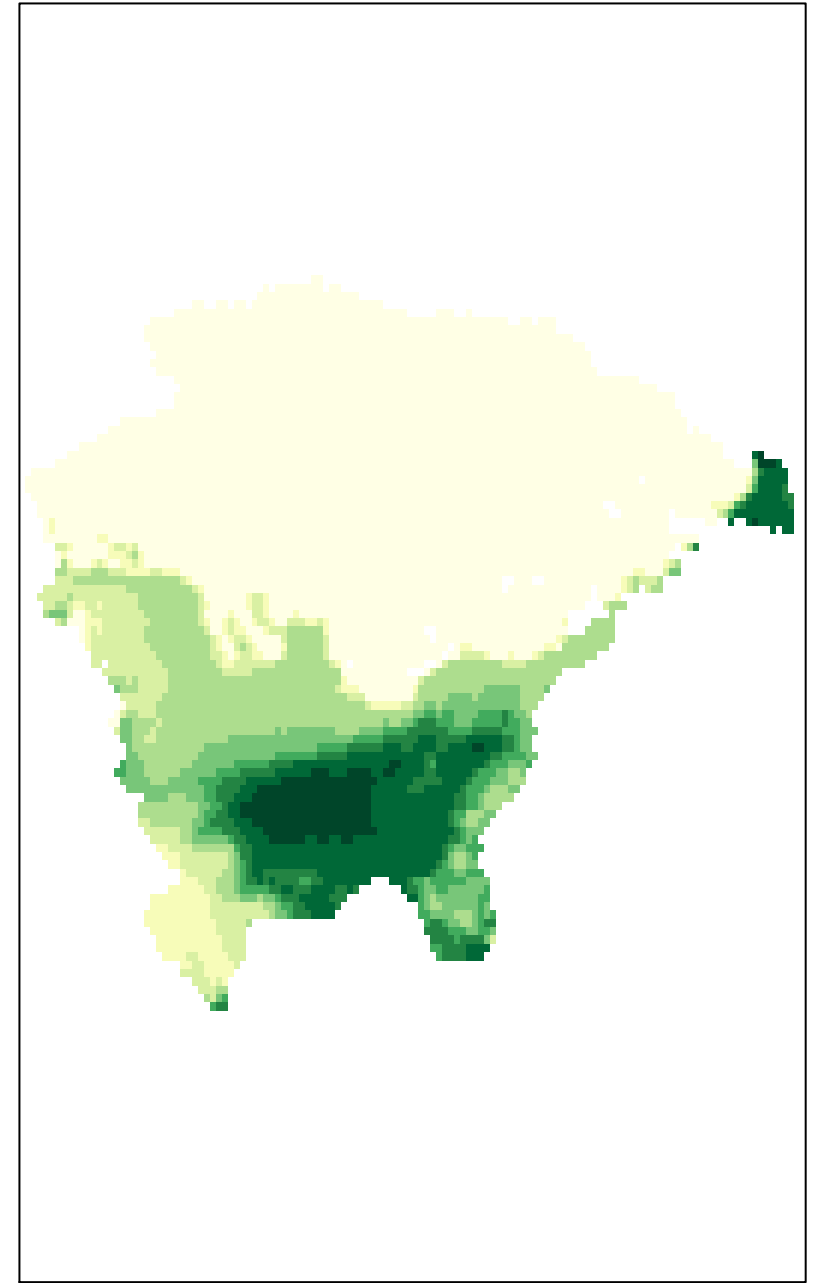


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X16000.ybp

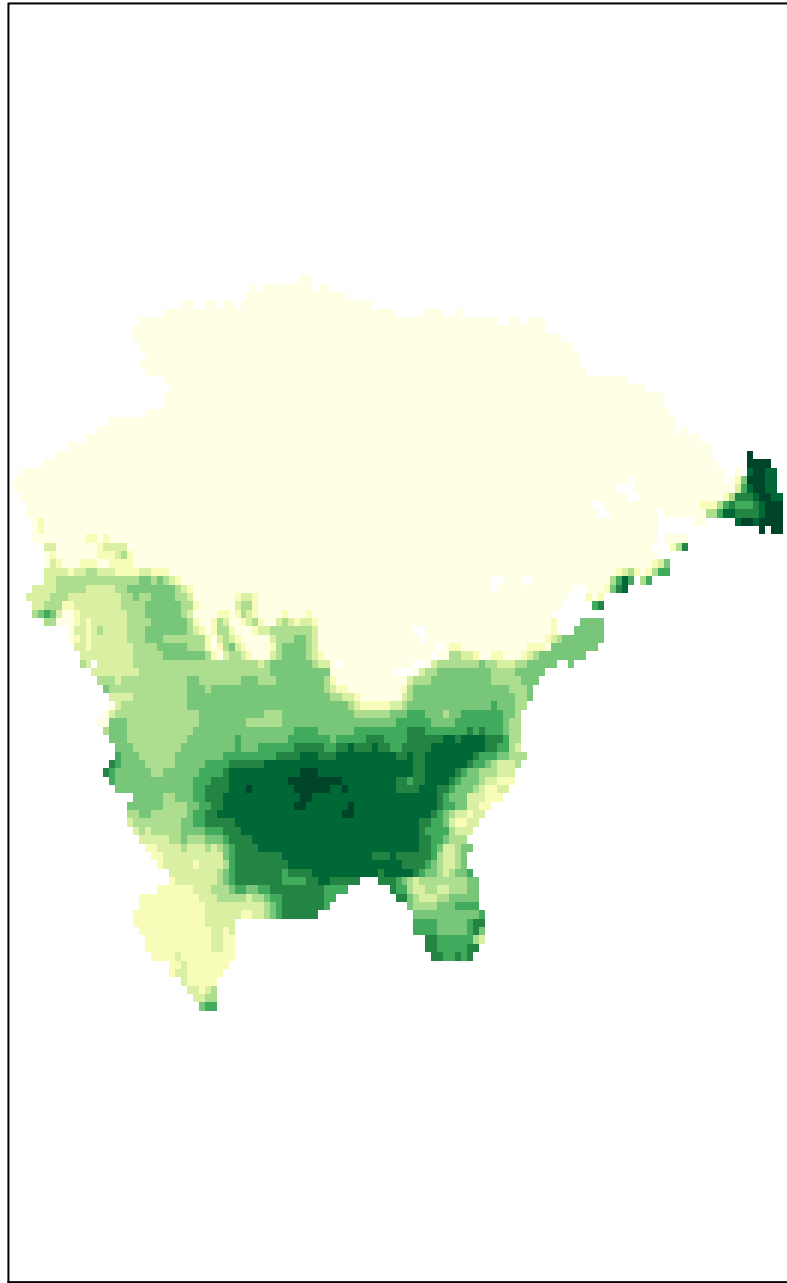


SUM, X16000.ybp

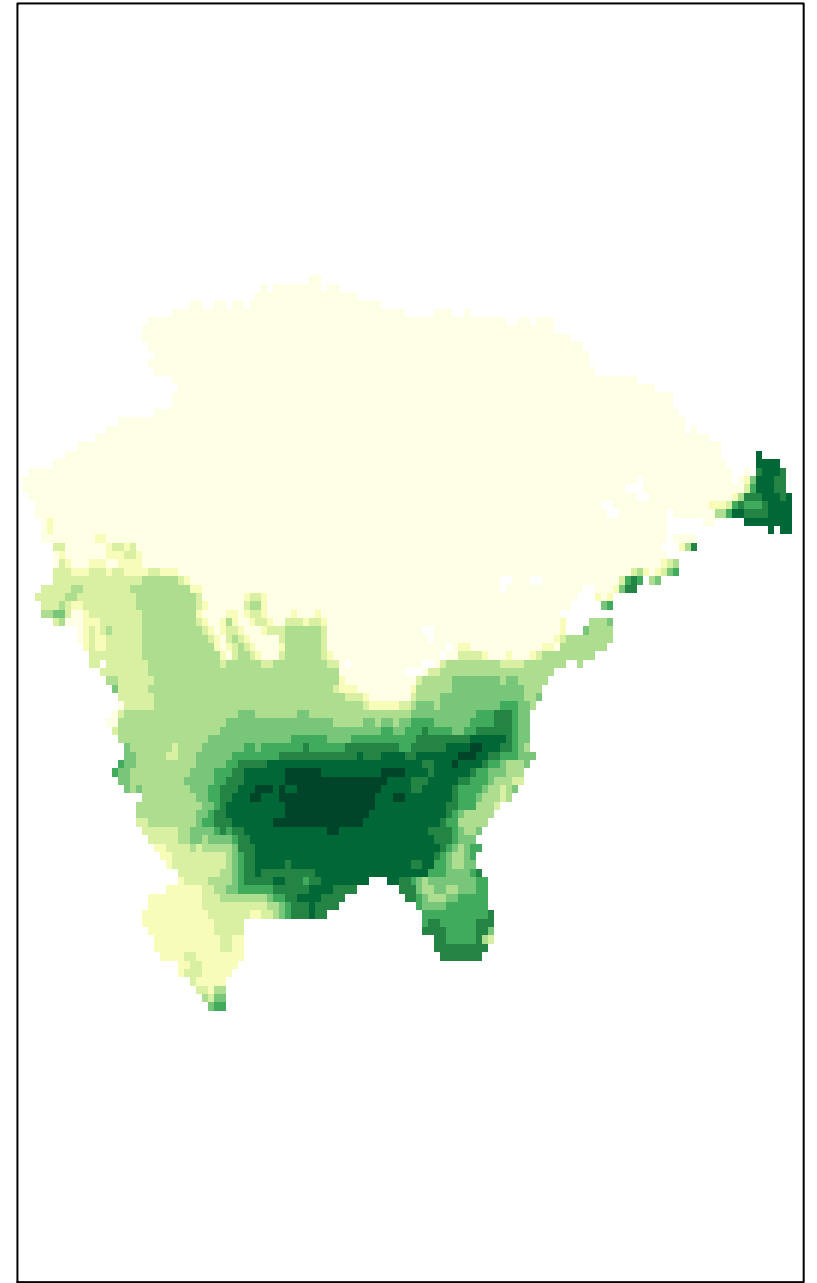


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X15000.ybp

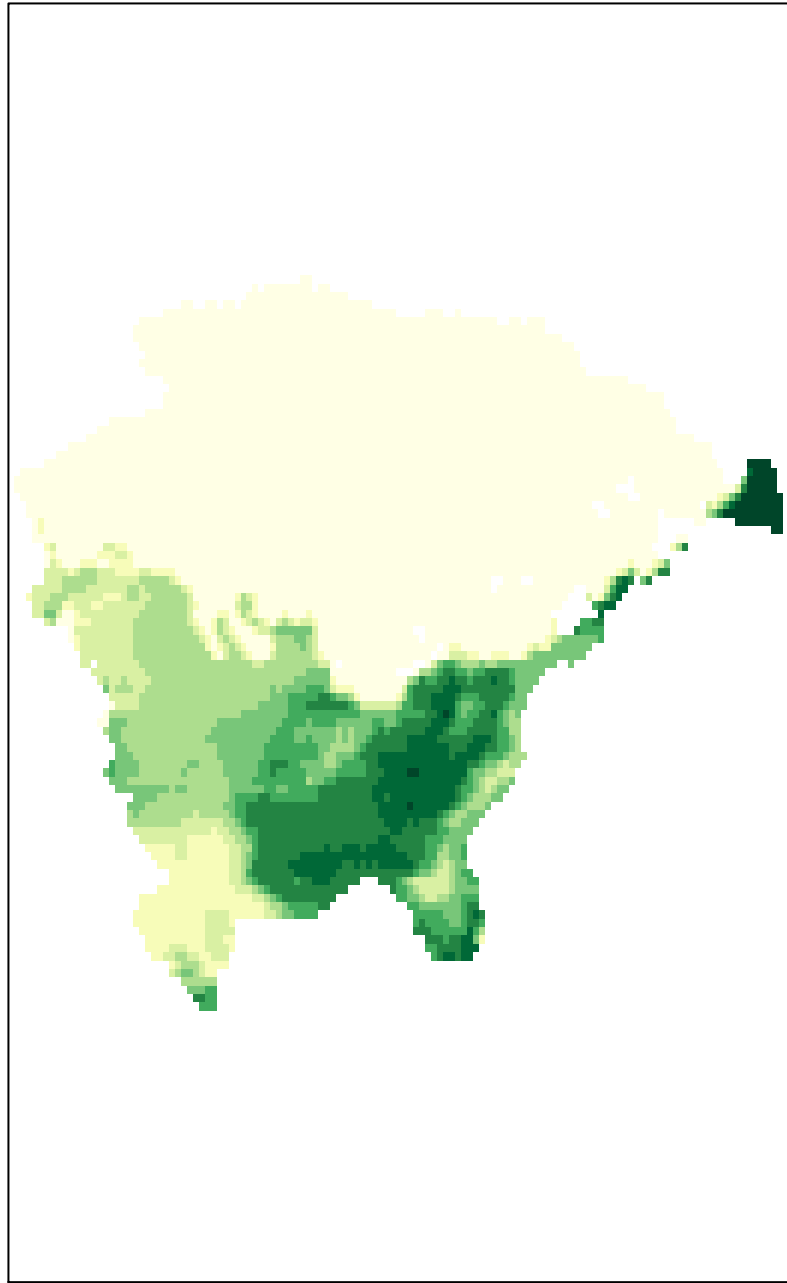


SUM, X15000.ybp

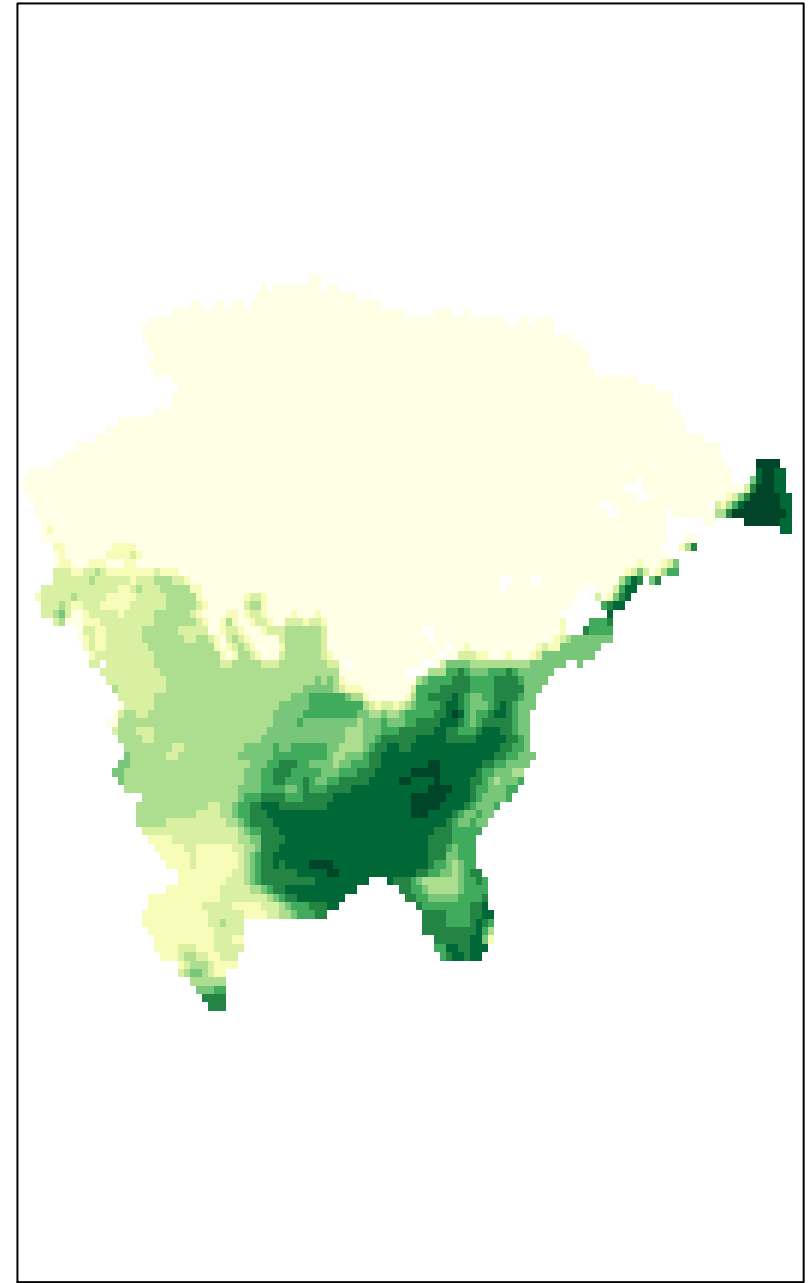


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X14000.ybp



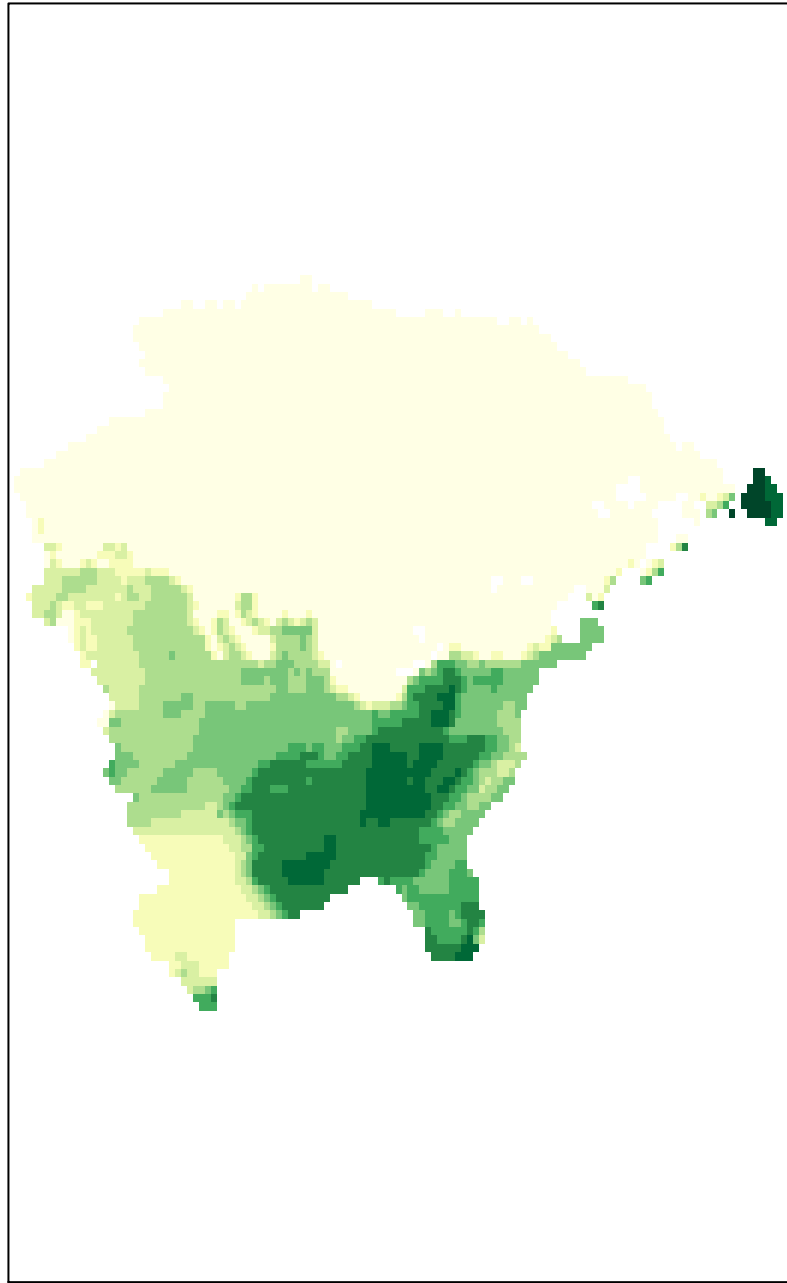
SUM, X14000.ybp



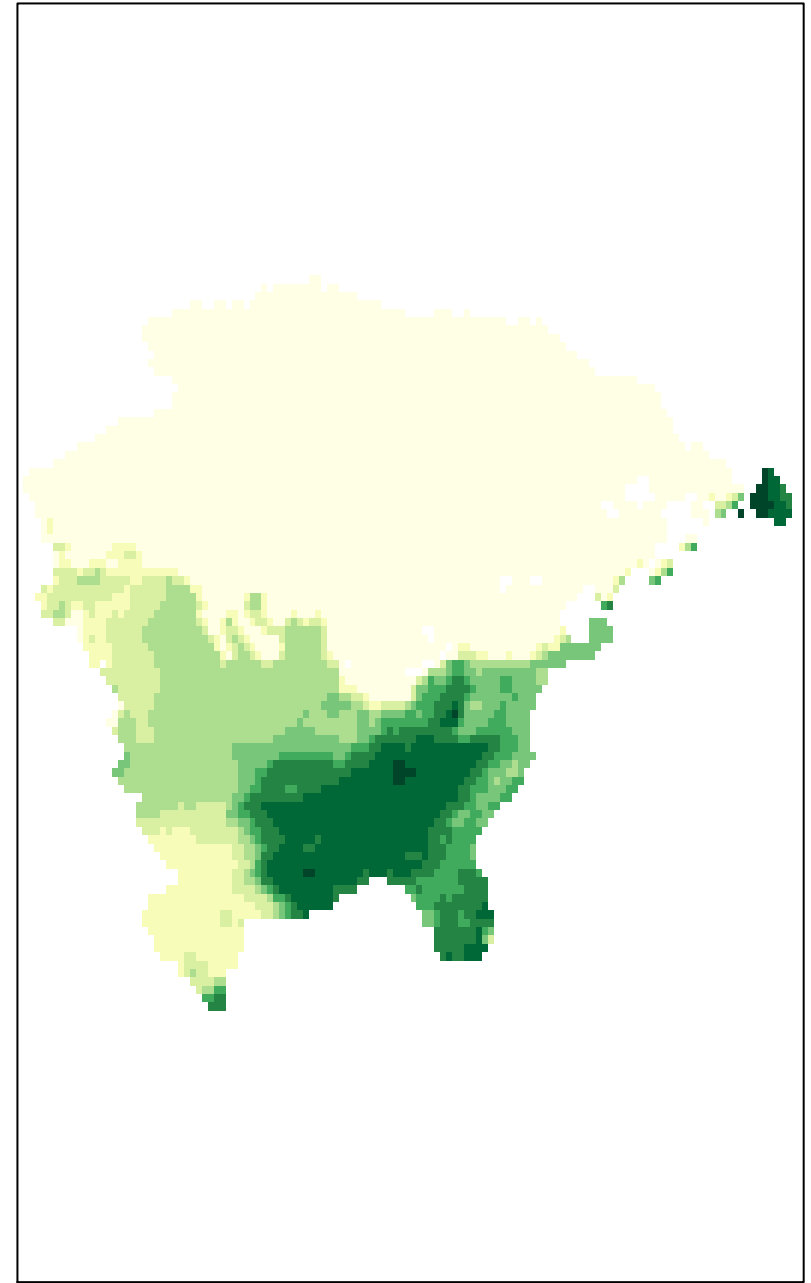
Species skipped = *Fraxinus profunda*, GCM = Beyer



SUM, X13000.ybp

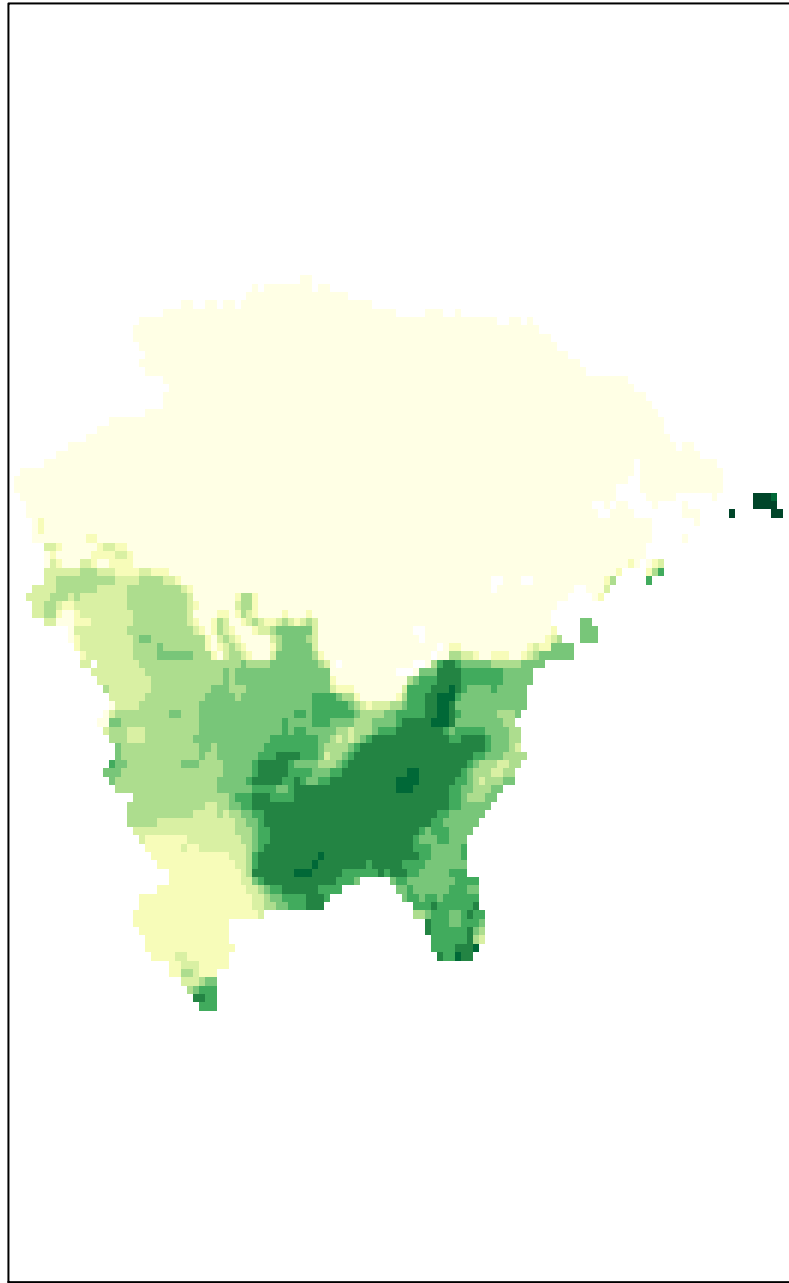


SUM, X13000.ybp

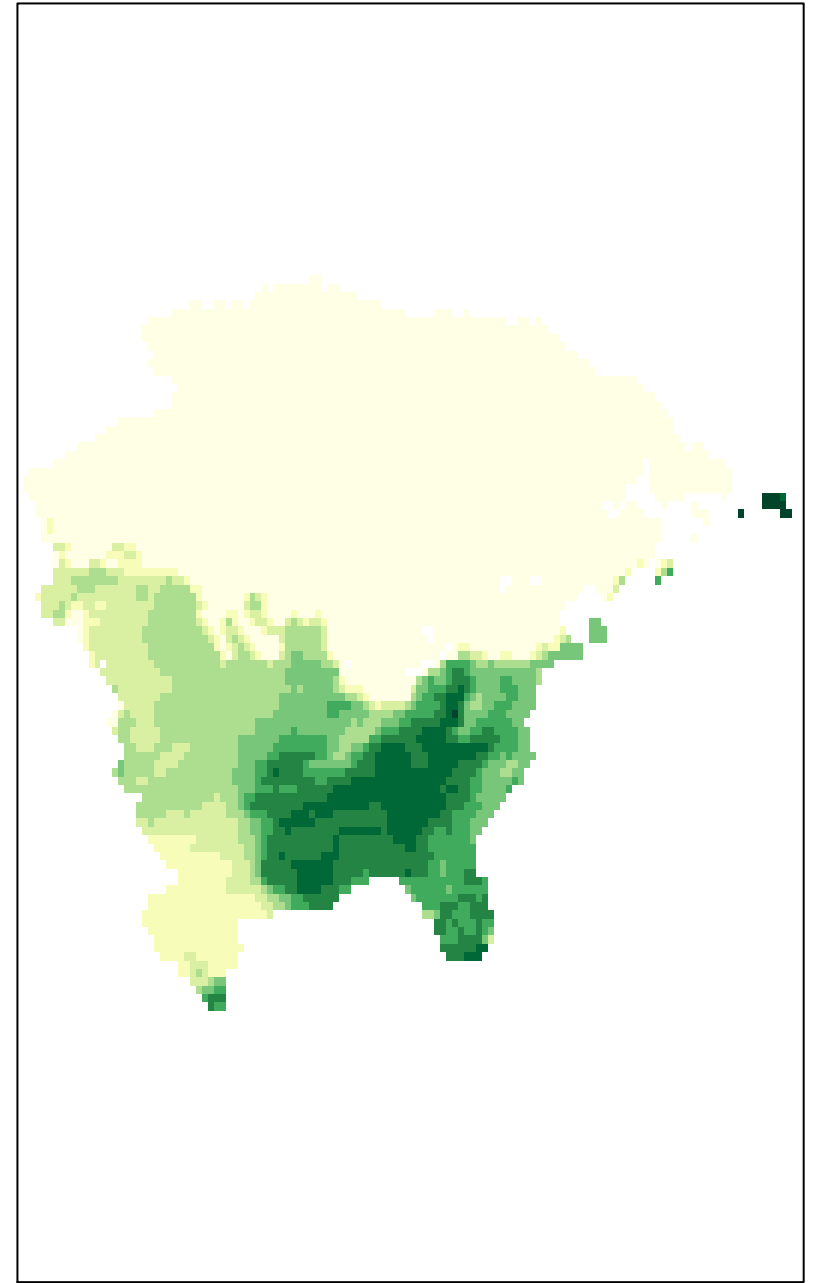


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X12000.ybp

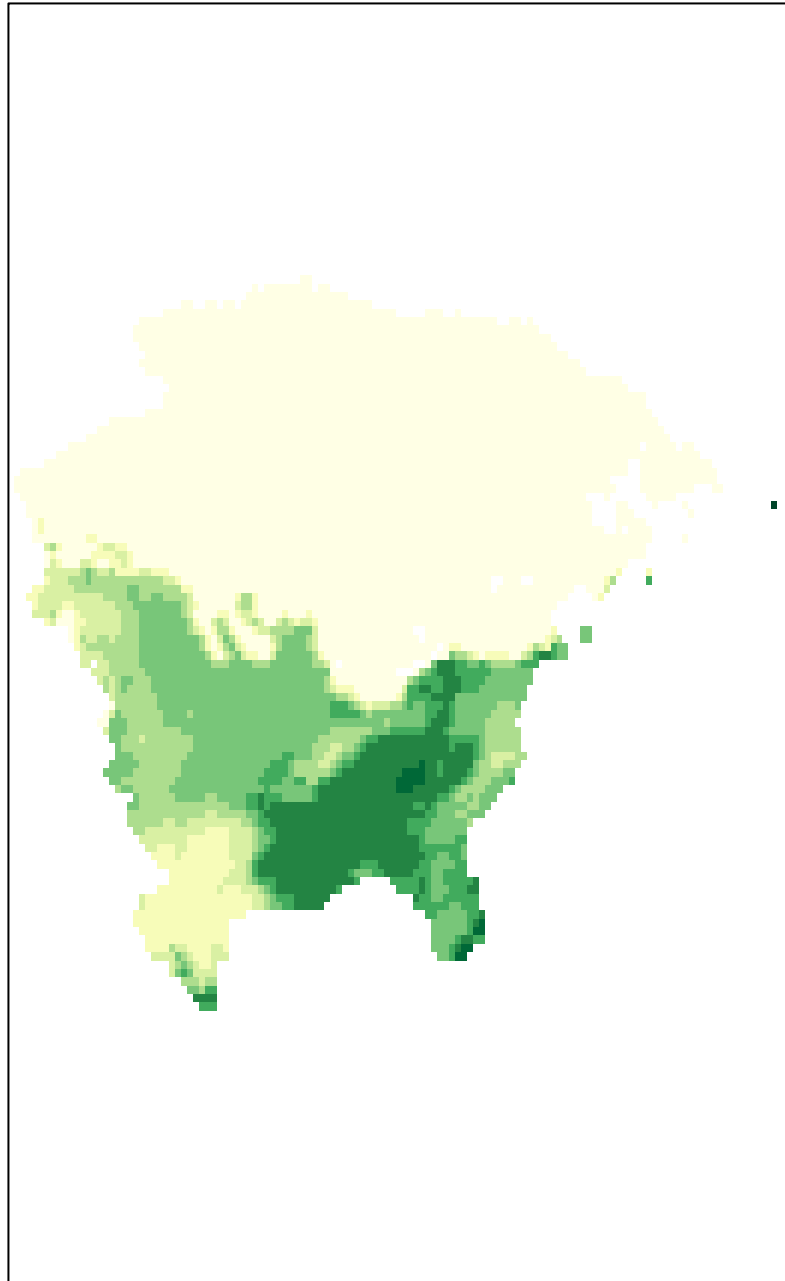


SUM, X12000.ybp

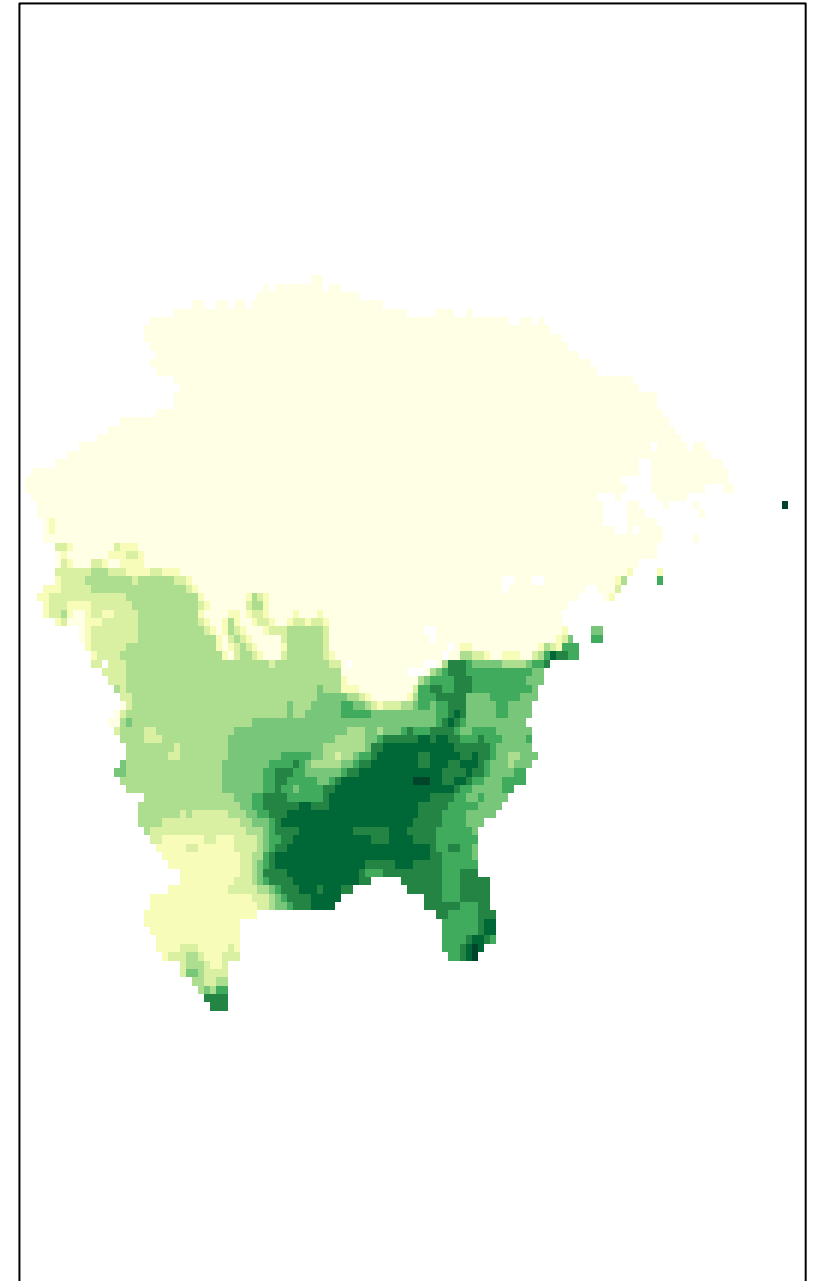


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X11000.ybp

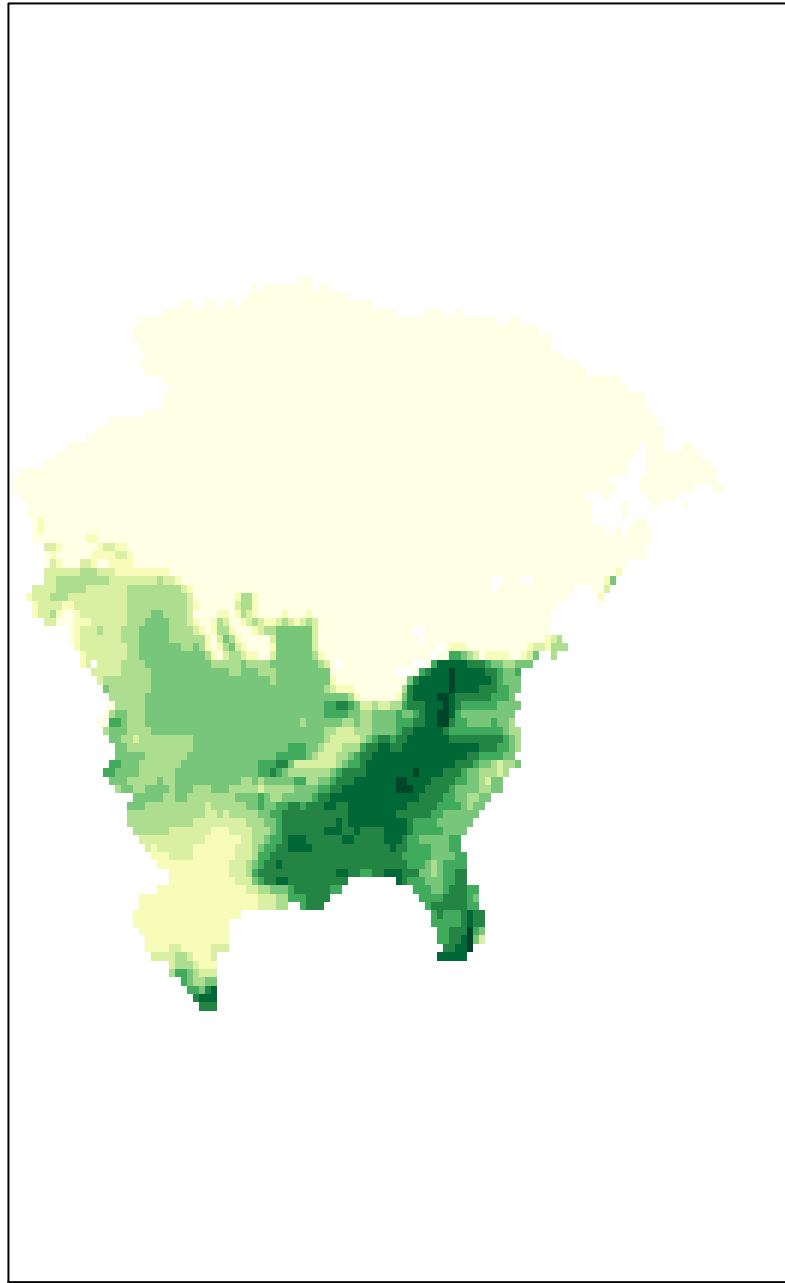


SUM, X11000.ybp

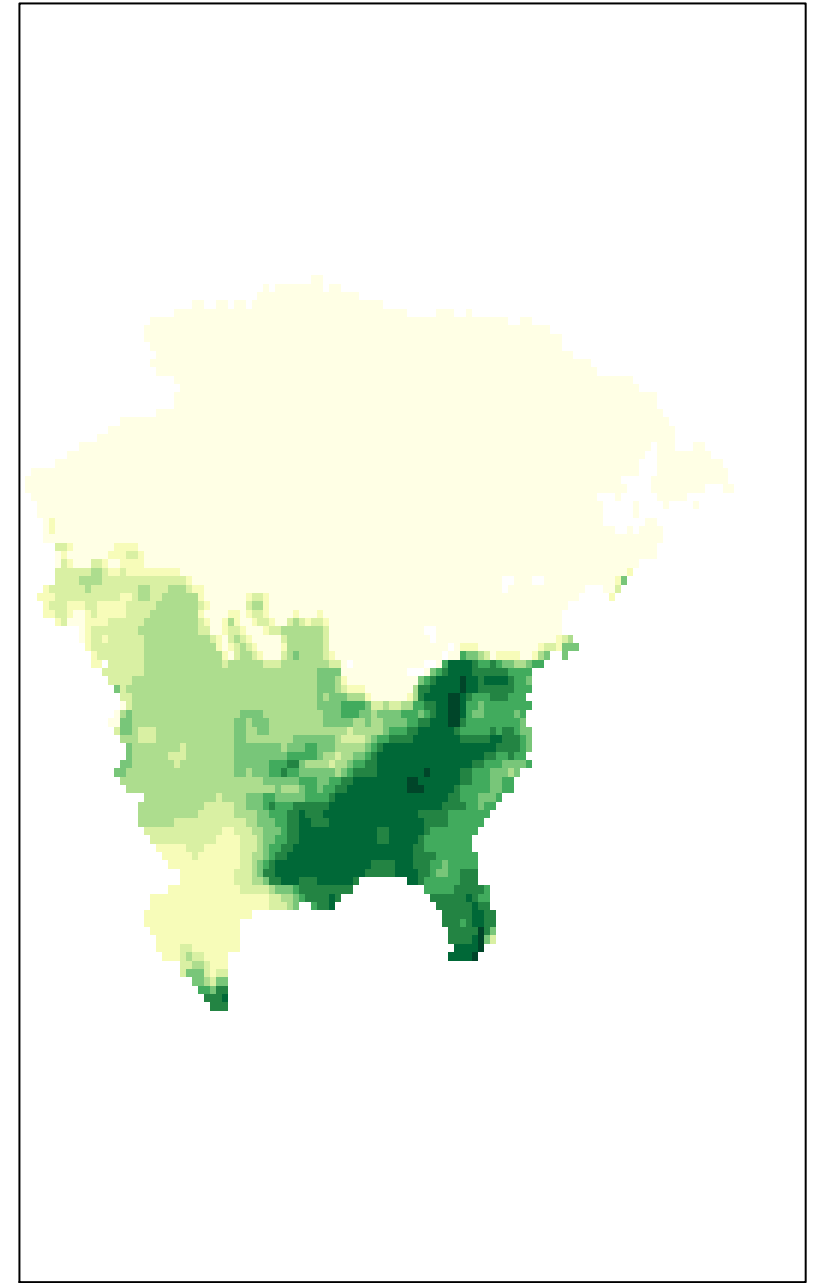


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X10000.ybp

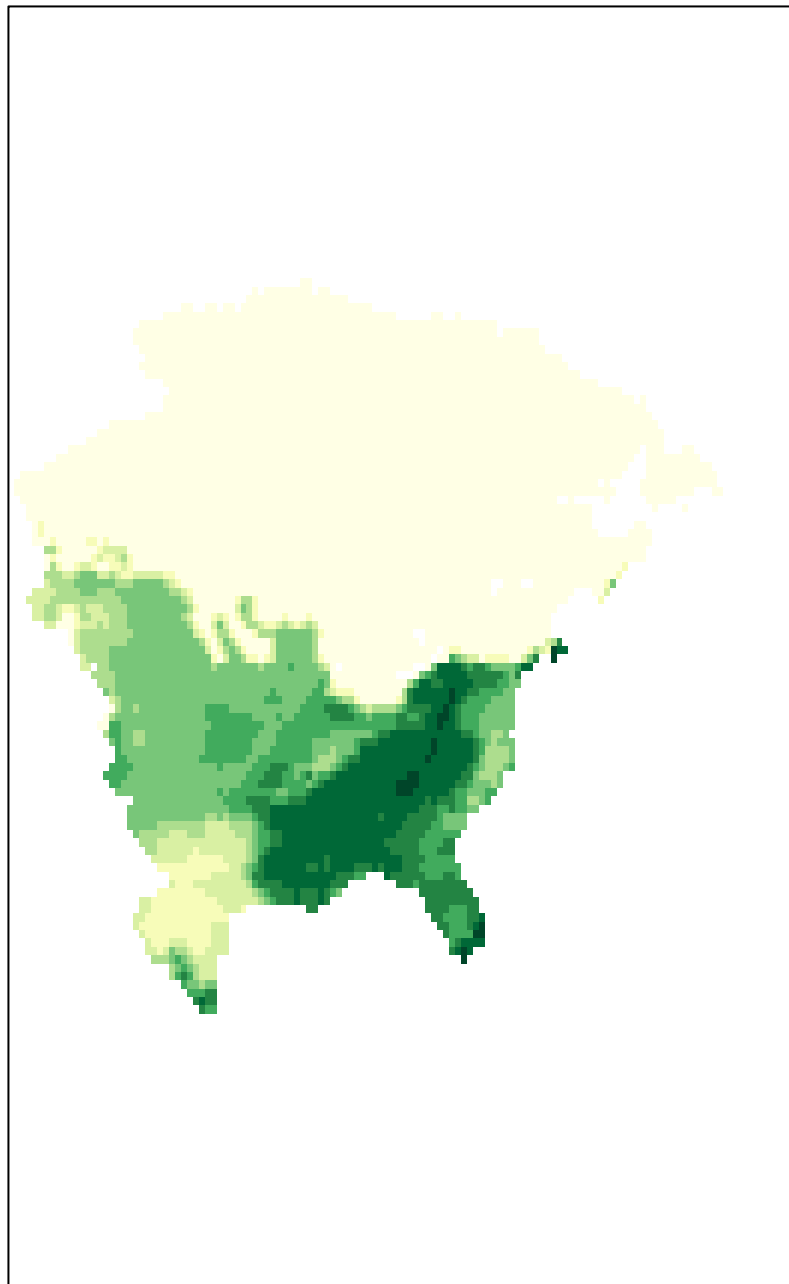


SUM, X10000.ybp

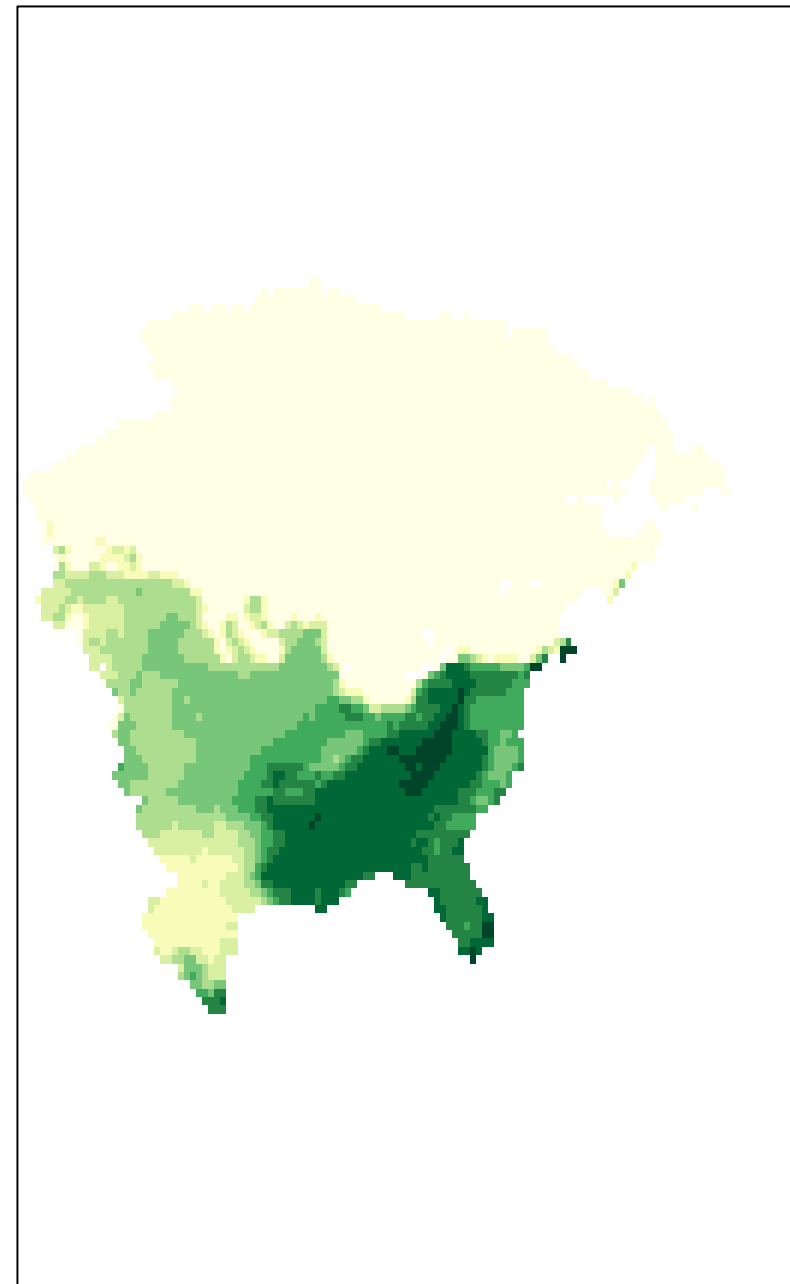


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X9000.ybp

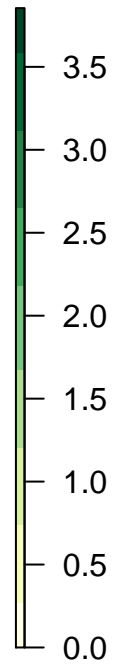
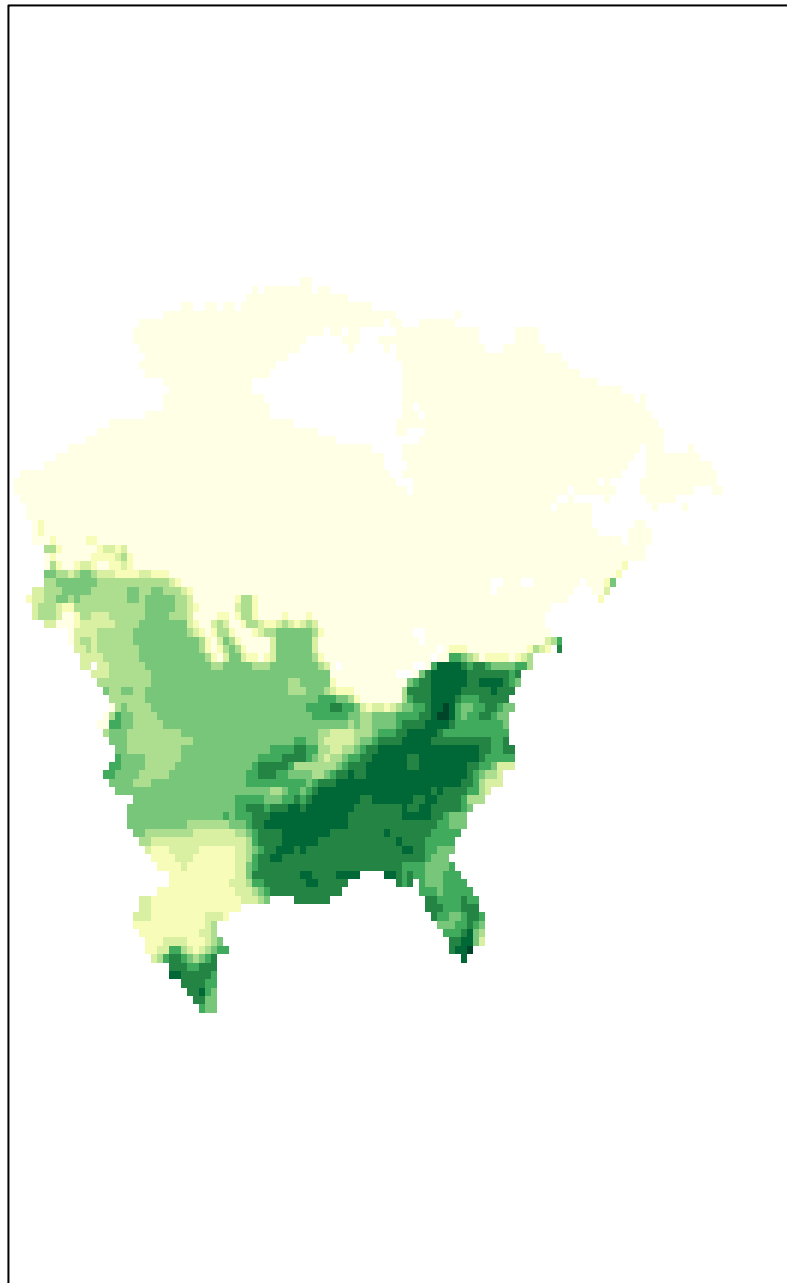


SUM, X9000.ybp

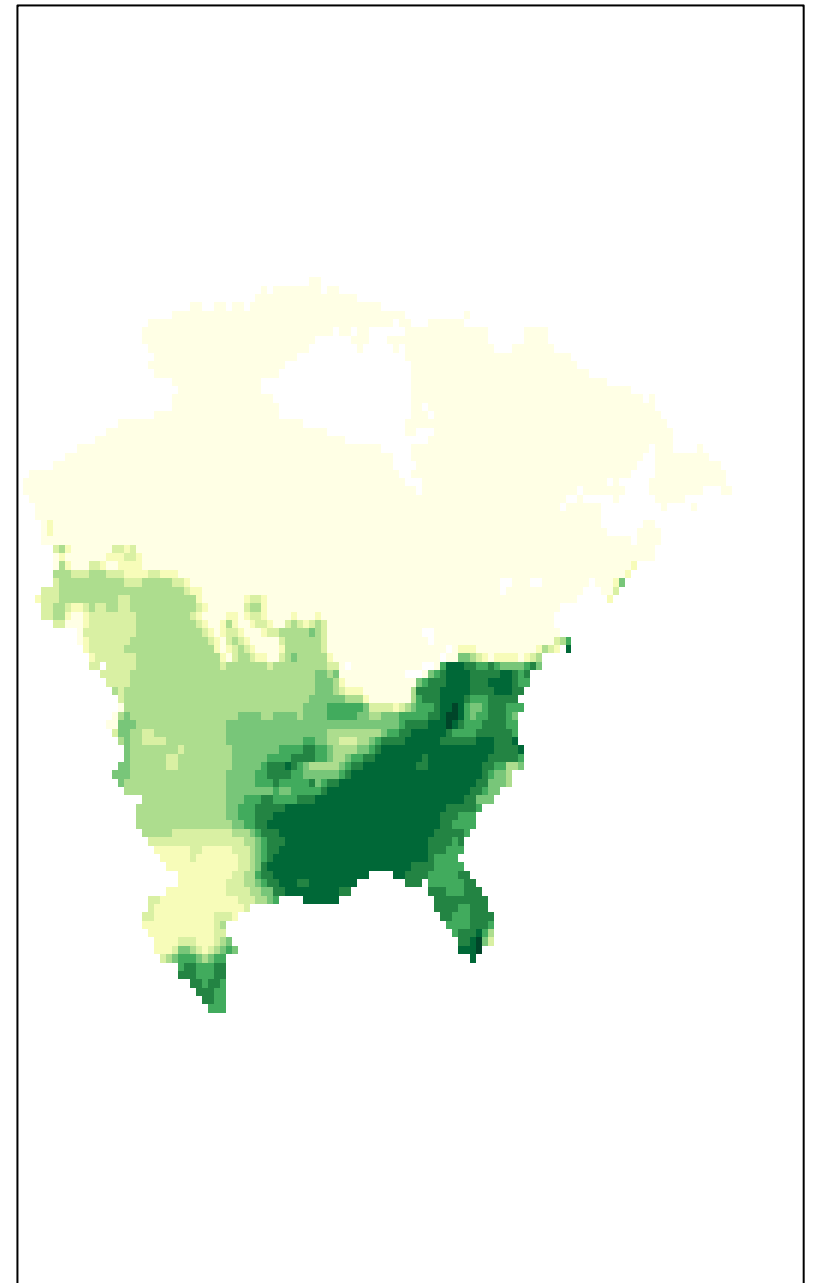


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X8000.ybp

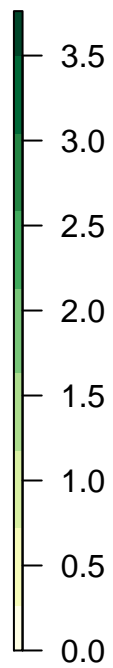
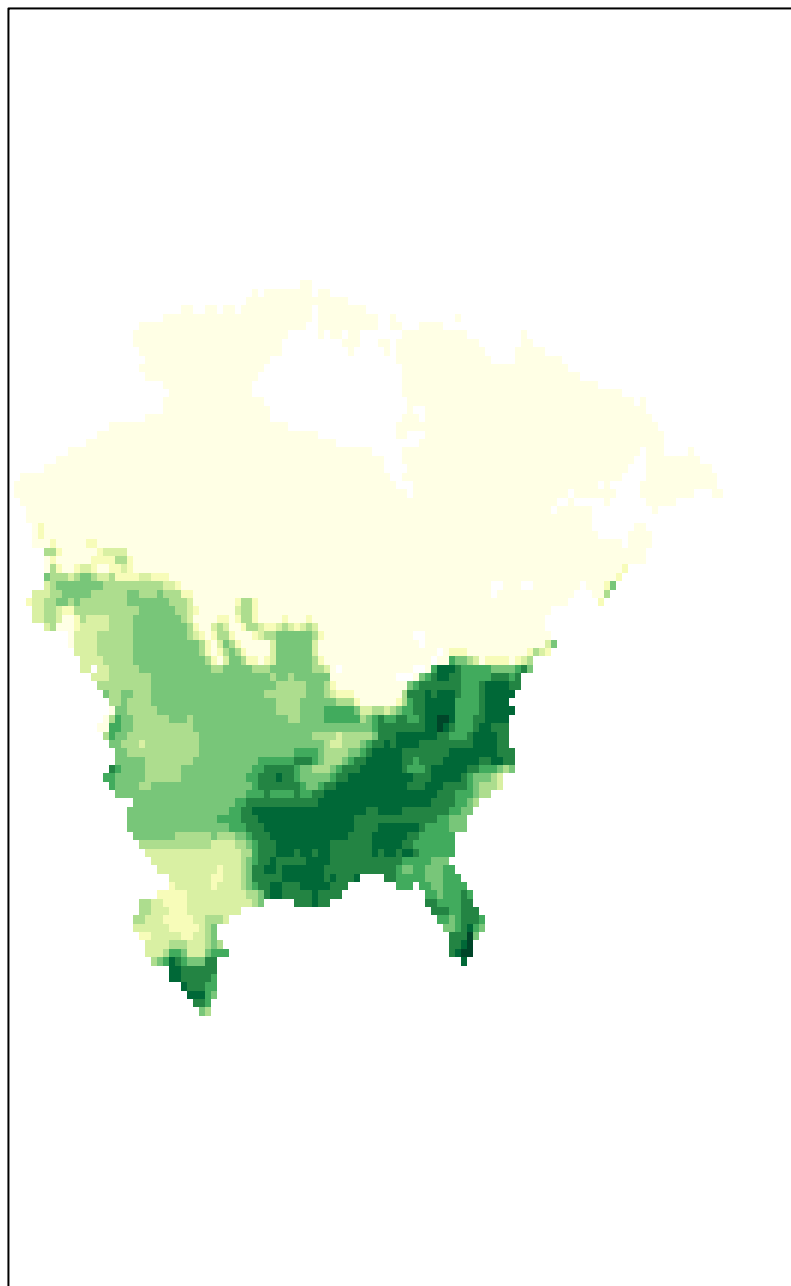


SUM, X8000.ybp

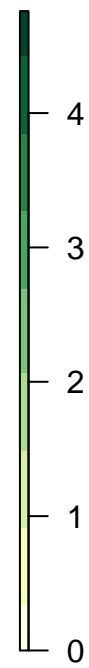
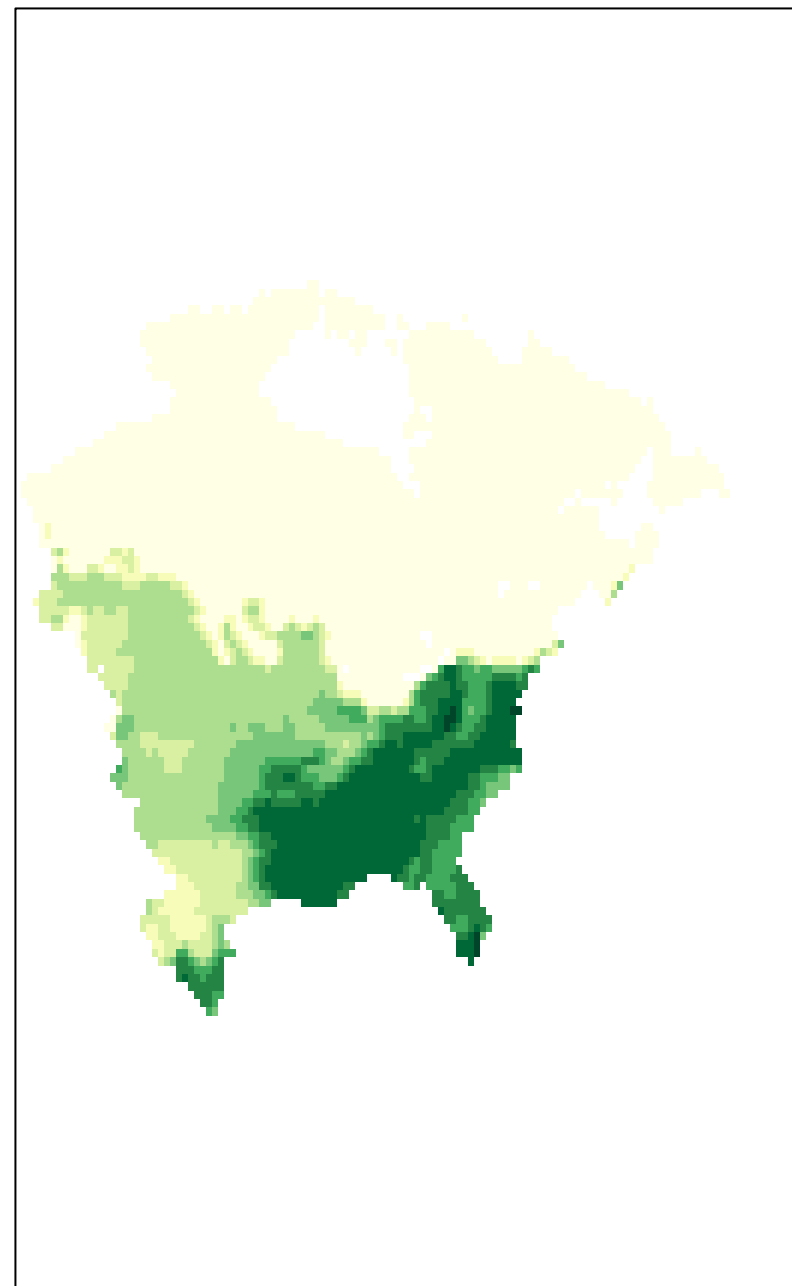


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X7000.ybp

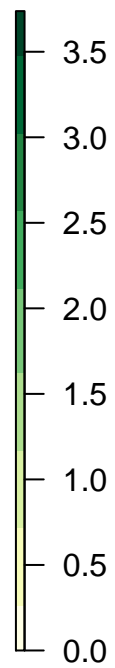
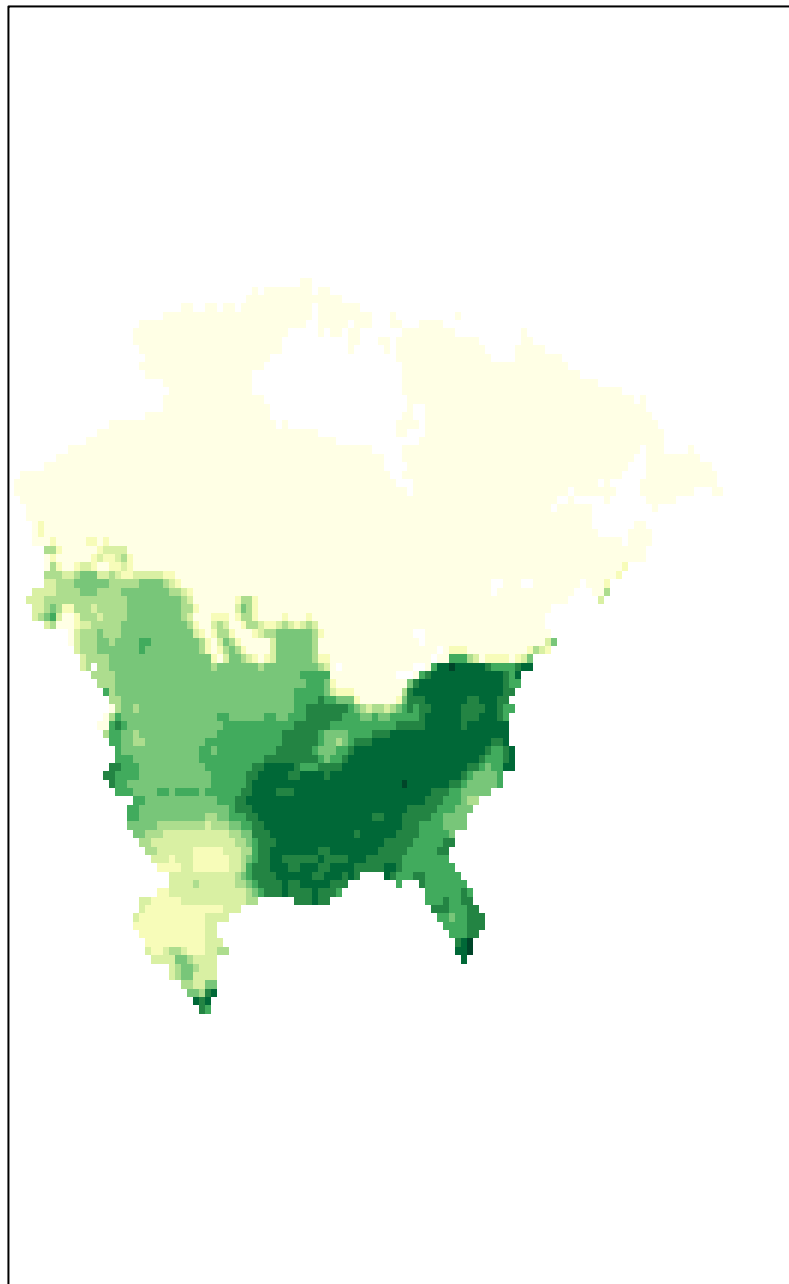


SUM, X7000.ybp

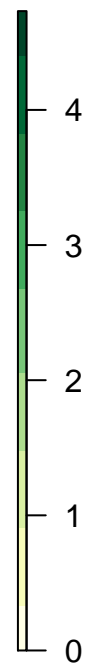
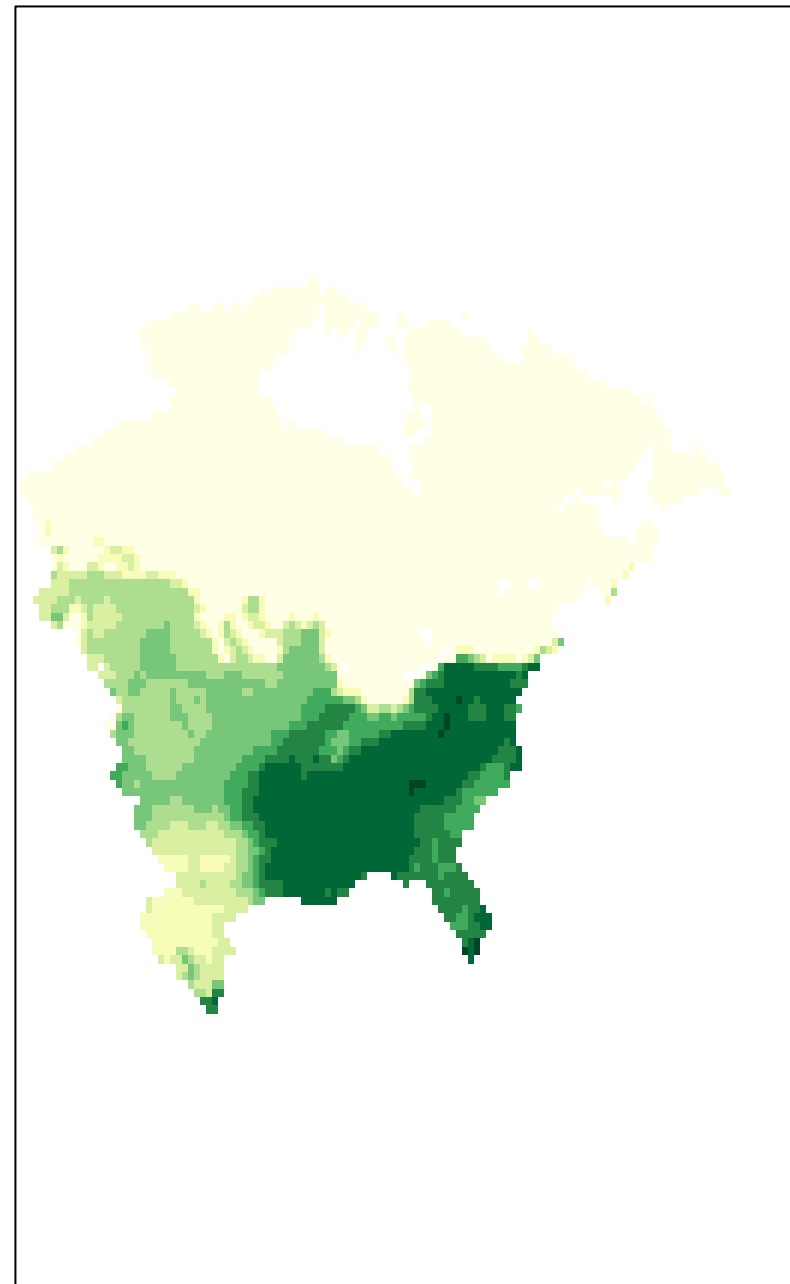


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X6000.ybp

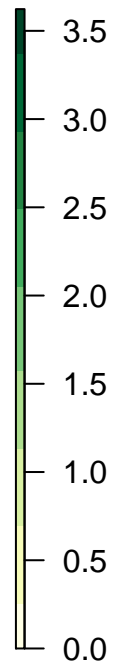
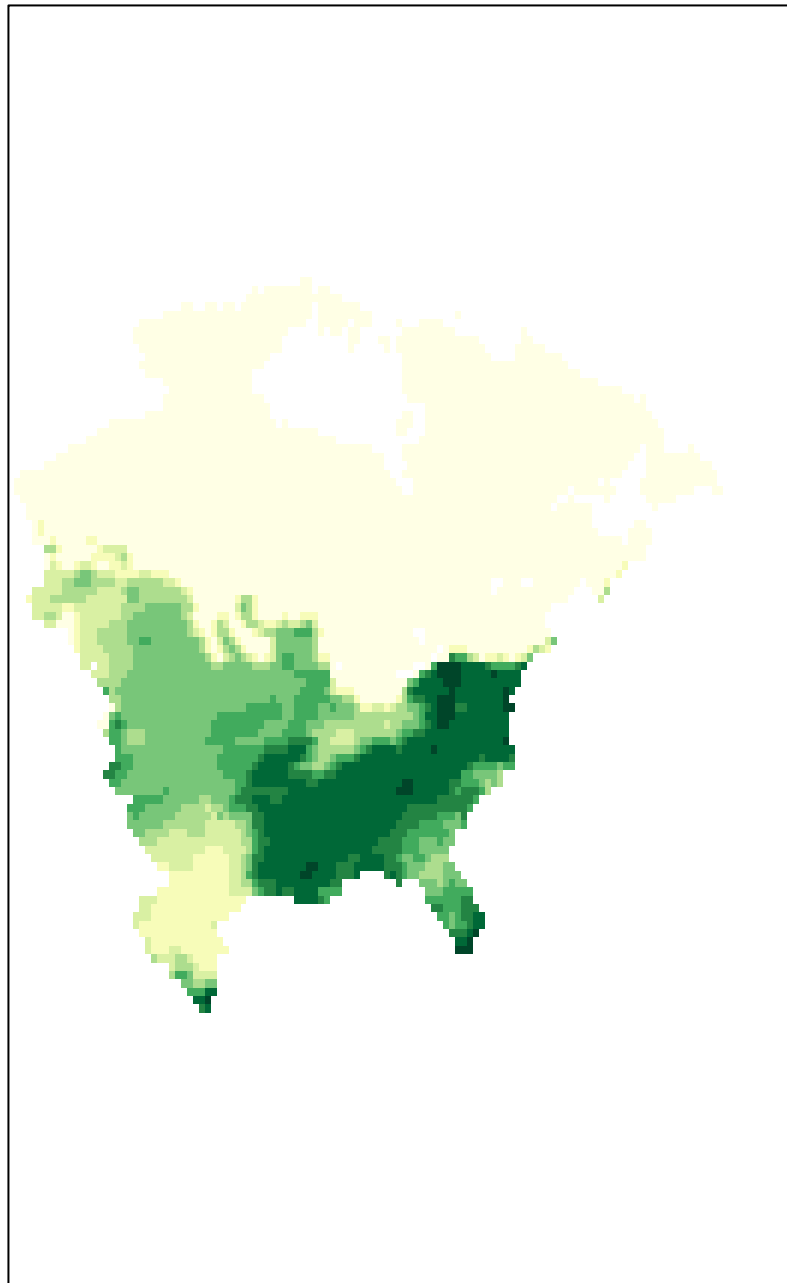


SUM, X6000.ybp

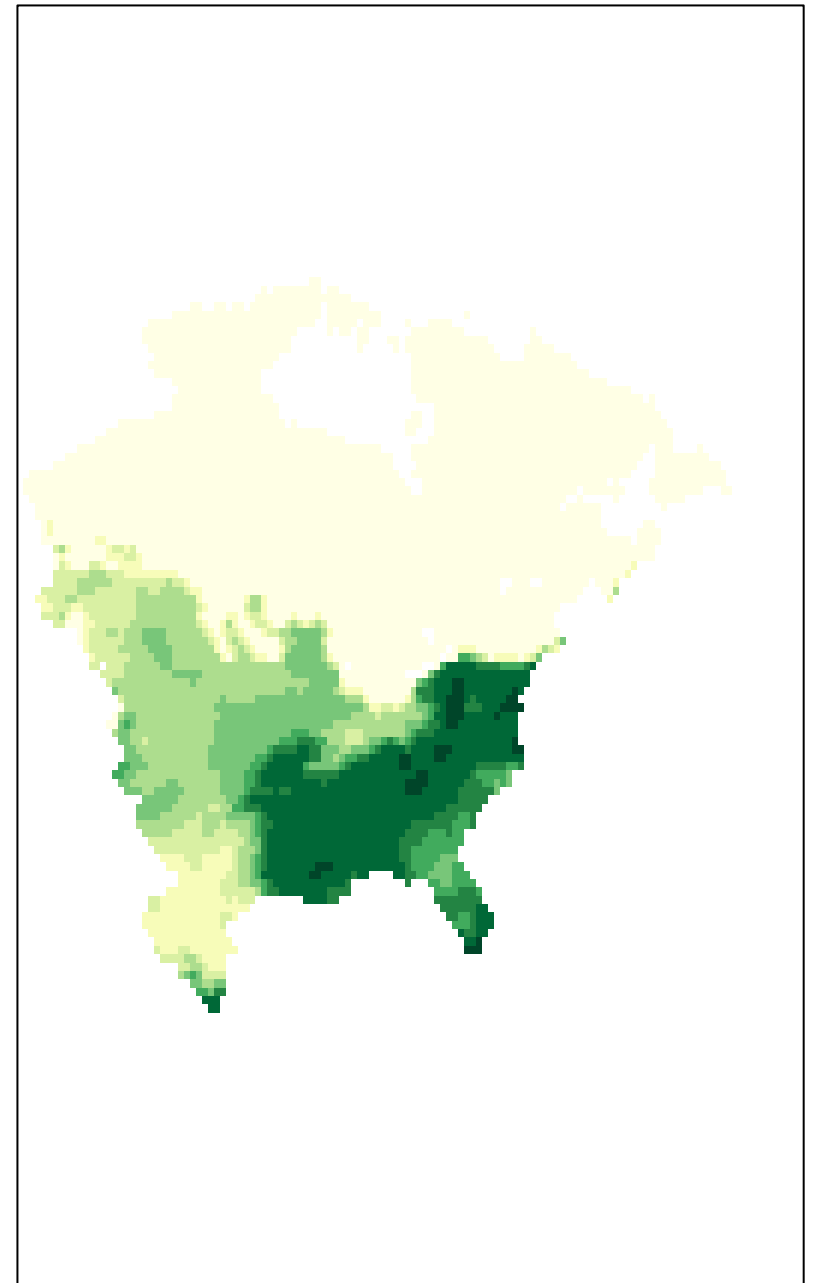




SUM, X5000.ybp

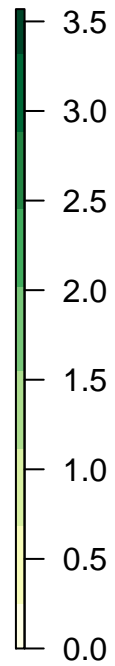
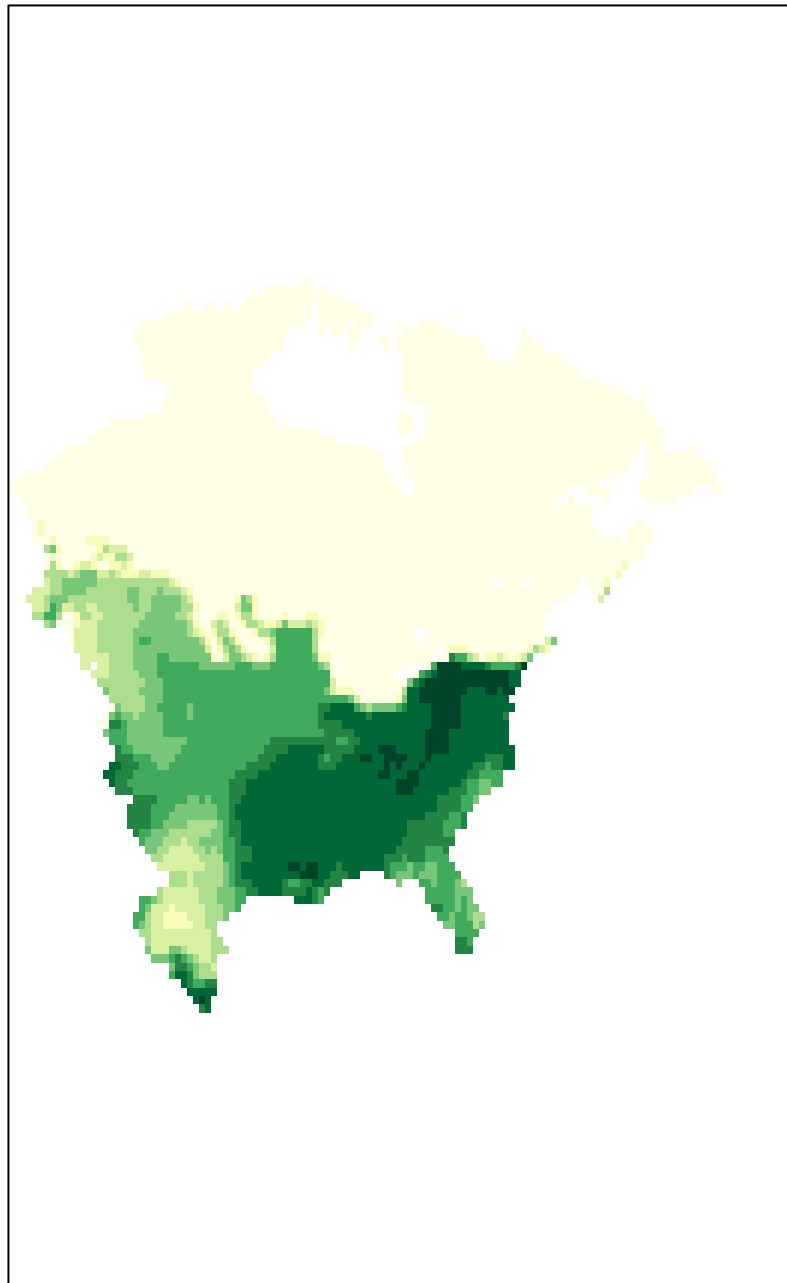


SUM, X5000.ybp

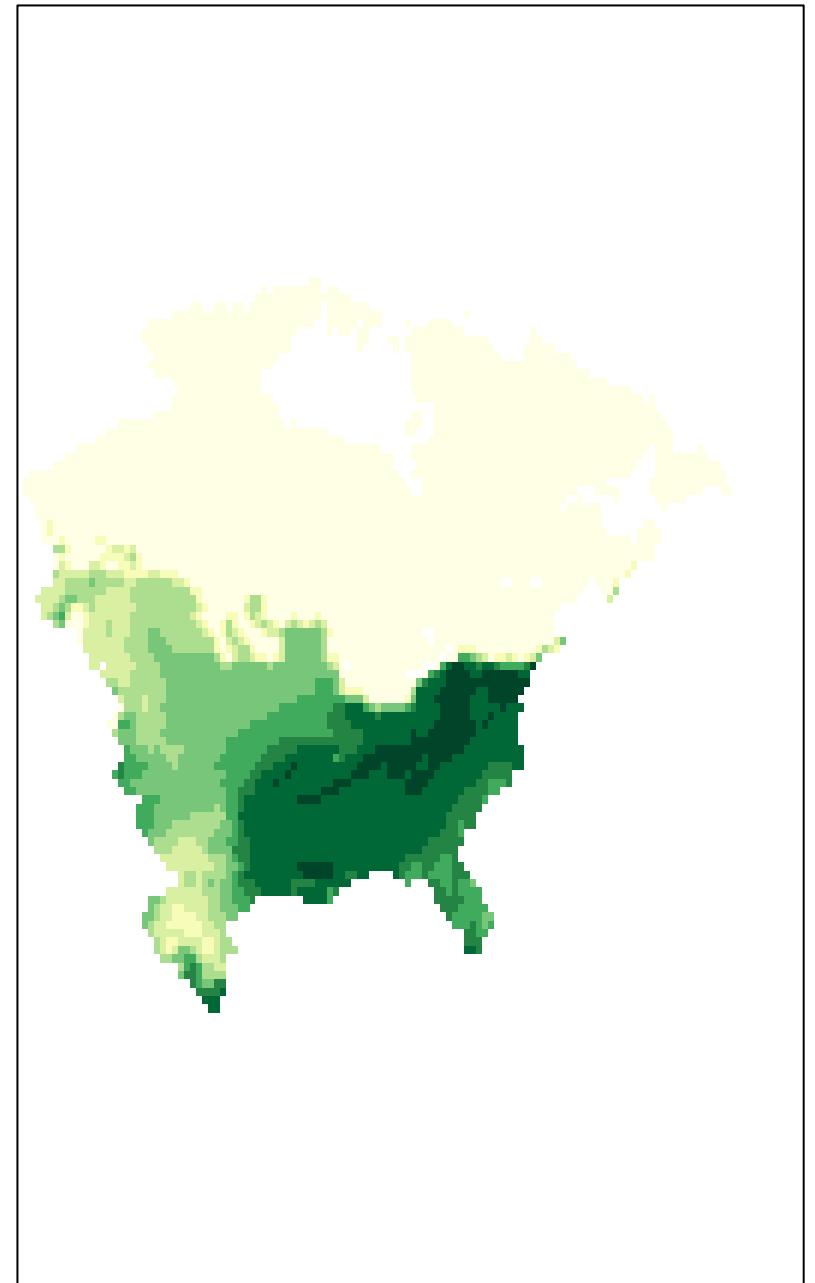


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X4000.ybp

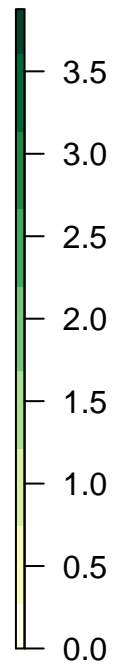
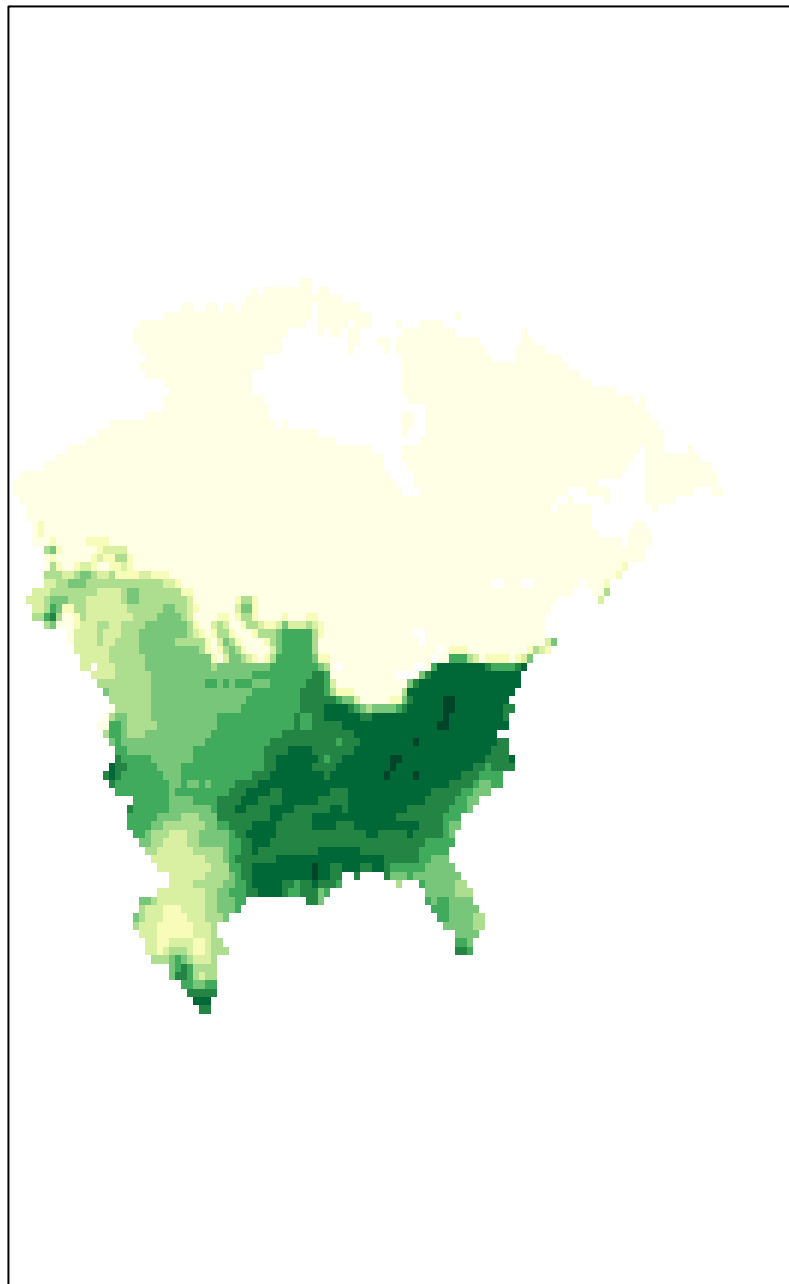


SUM, X4000.ybp

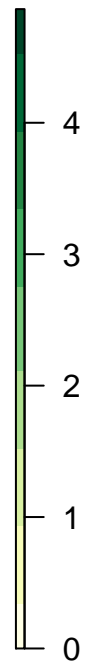
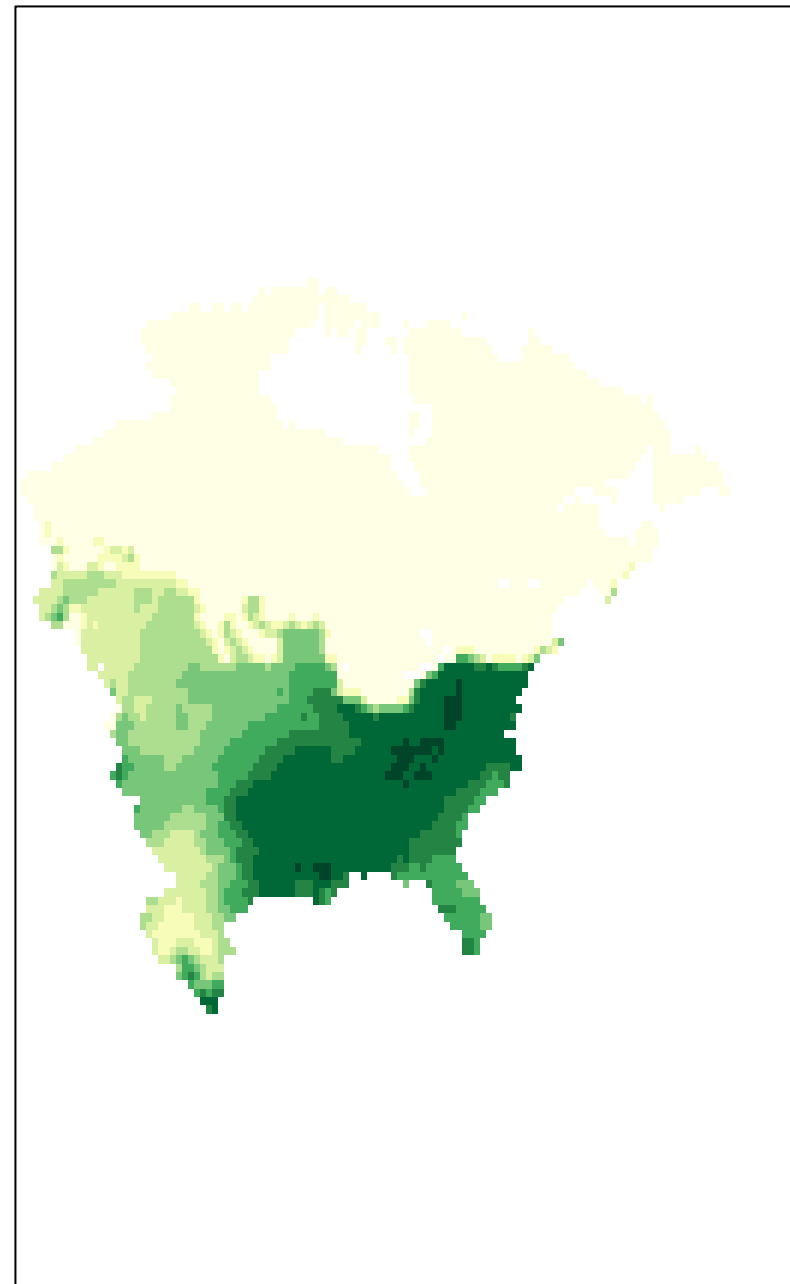


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X3000.ybp

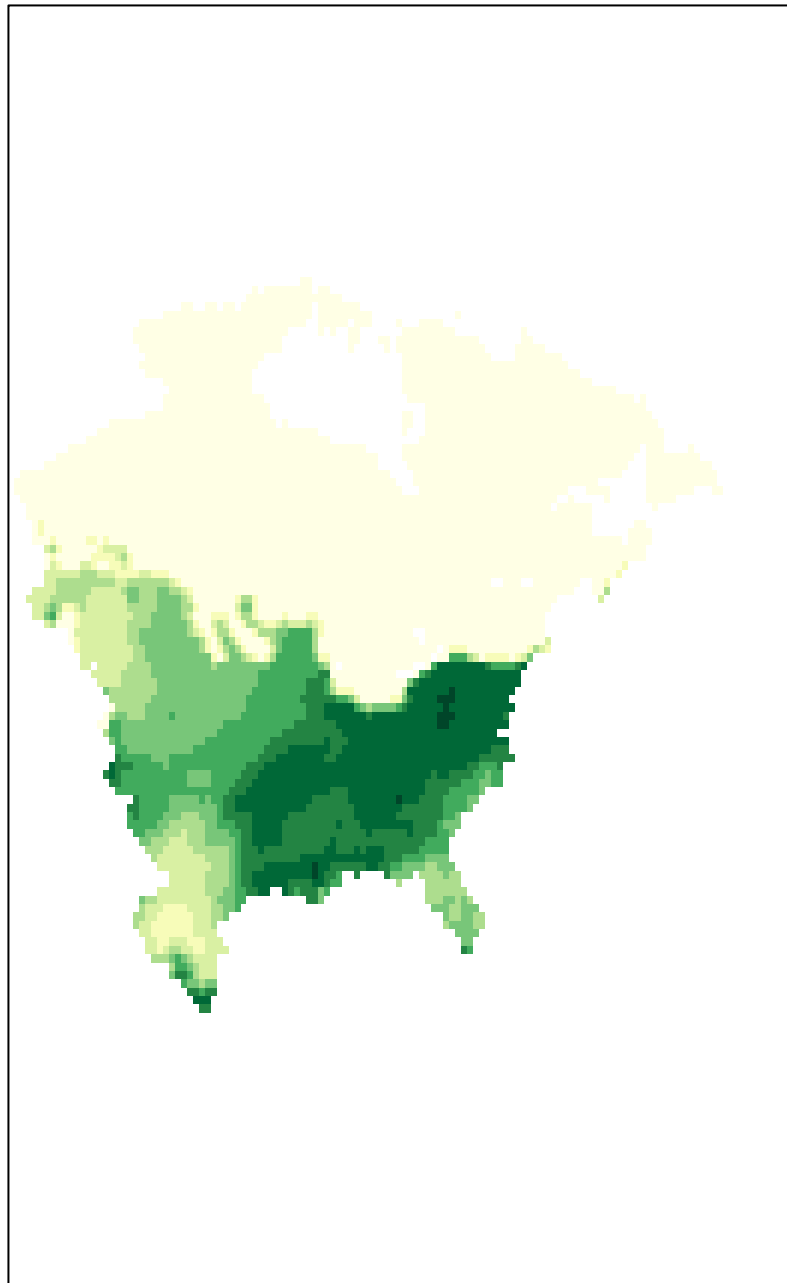


SUM, X3000.ybp

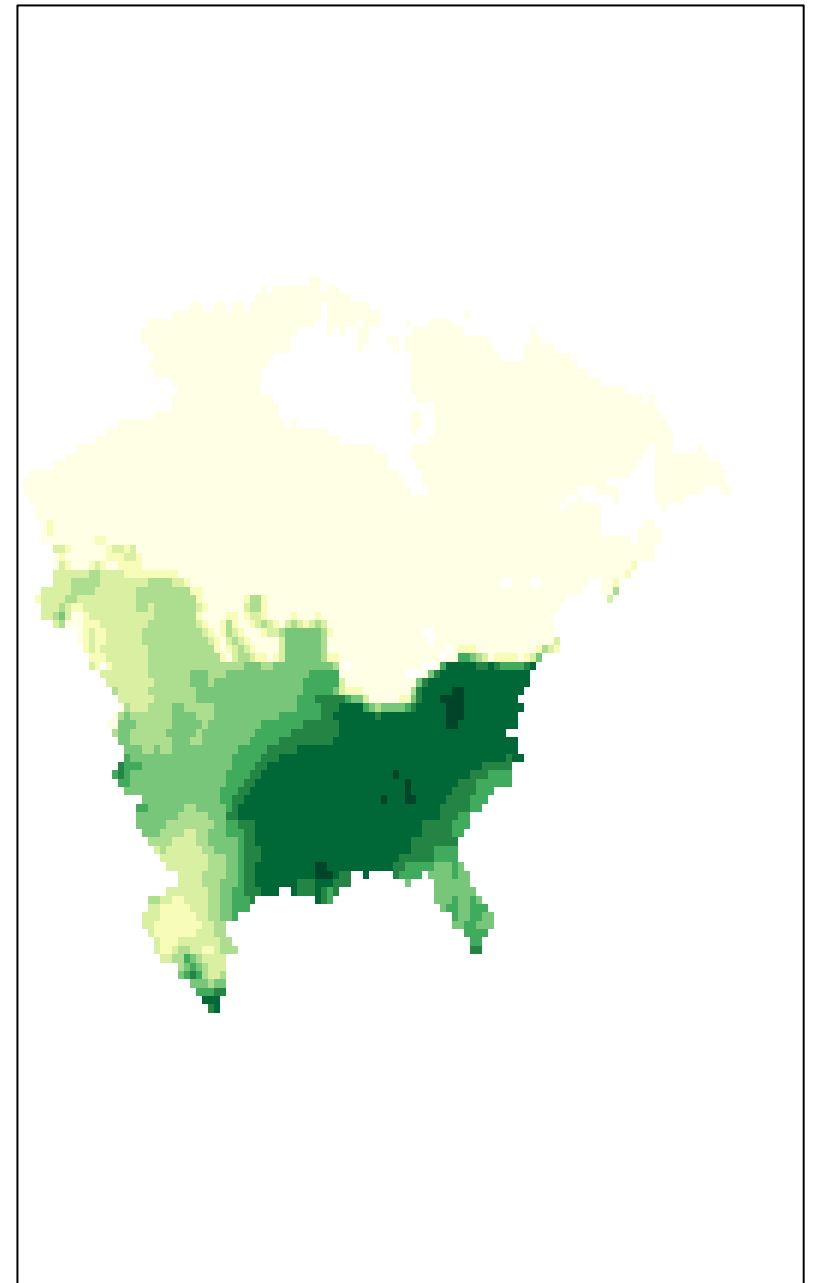


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X2000.ybp

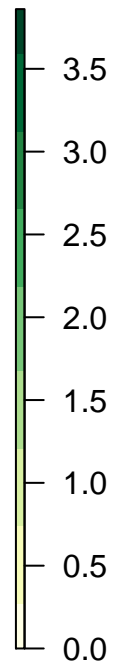
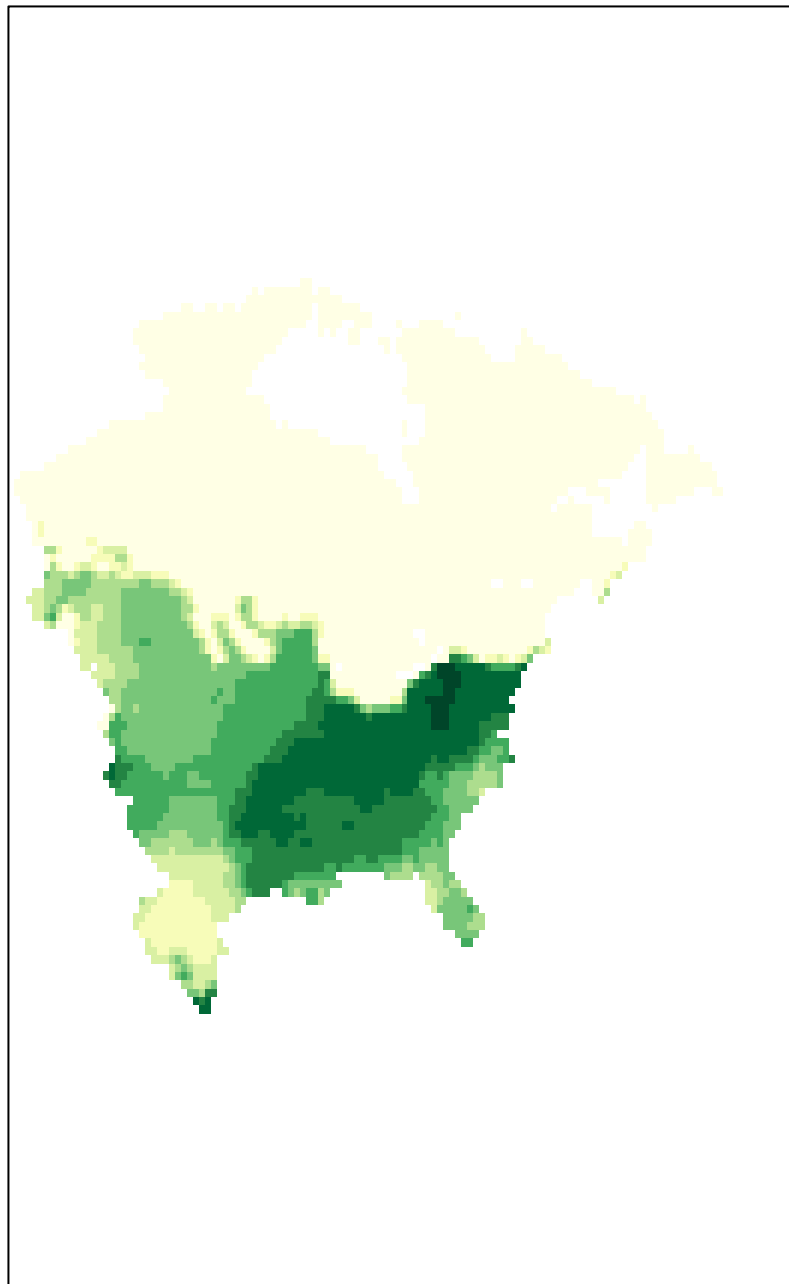


SUM, X2000.ybp

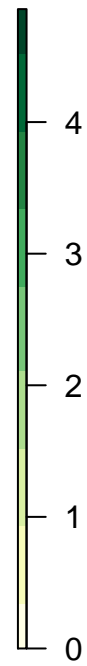
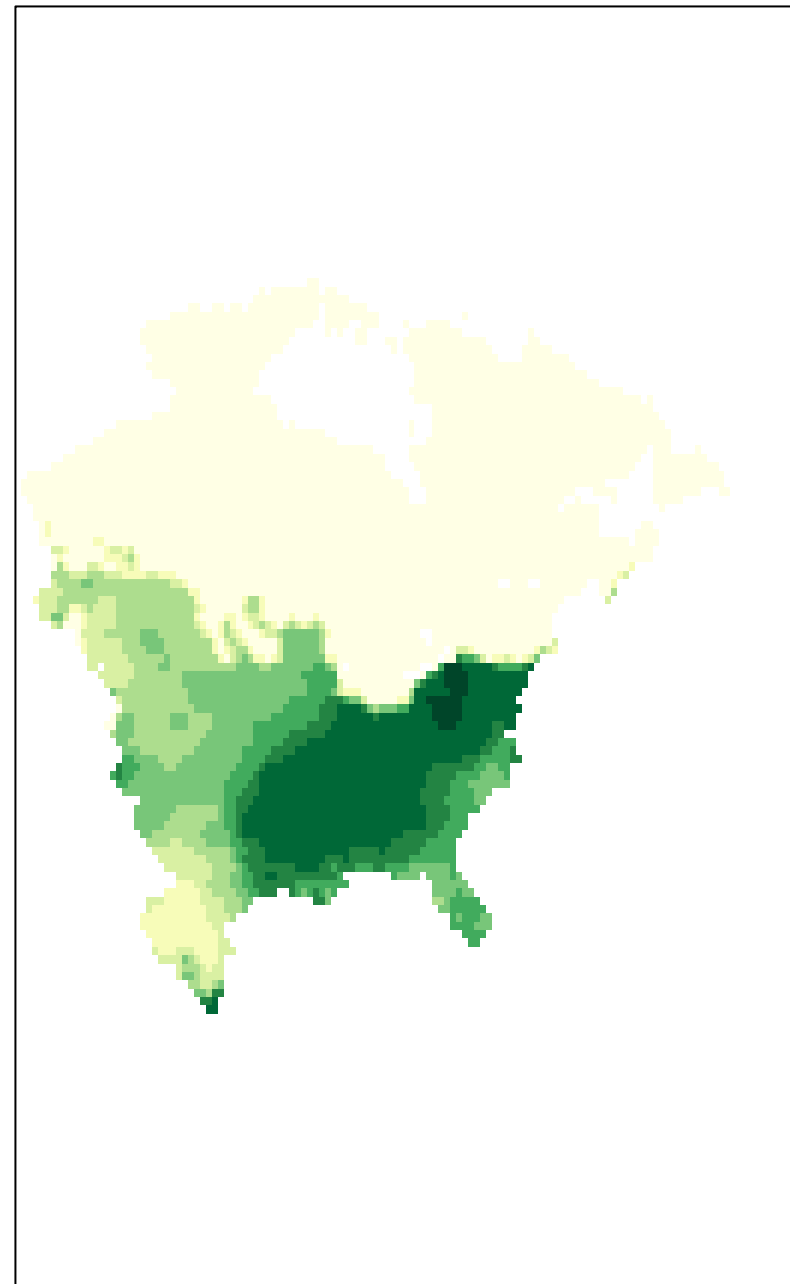


Species skipped = *Fraxinus profunda*, GCM = Beyer

SUM, X1000.ybp

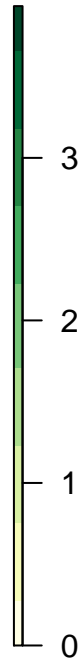
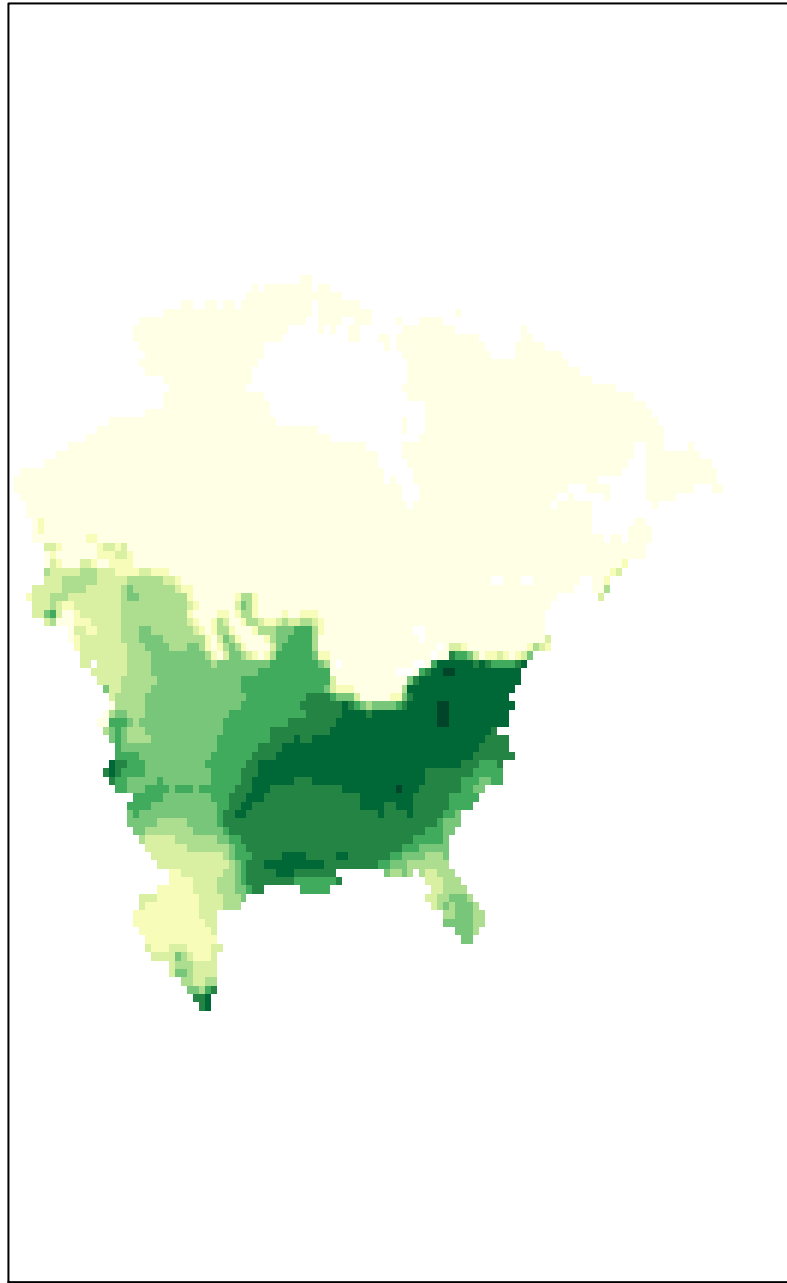


SUM, X1000.ybp

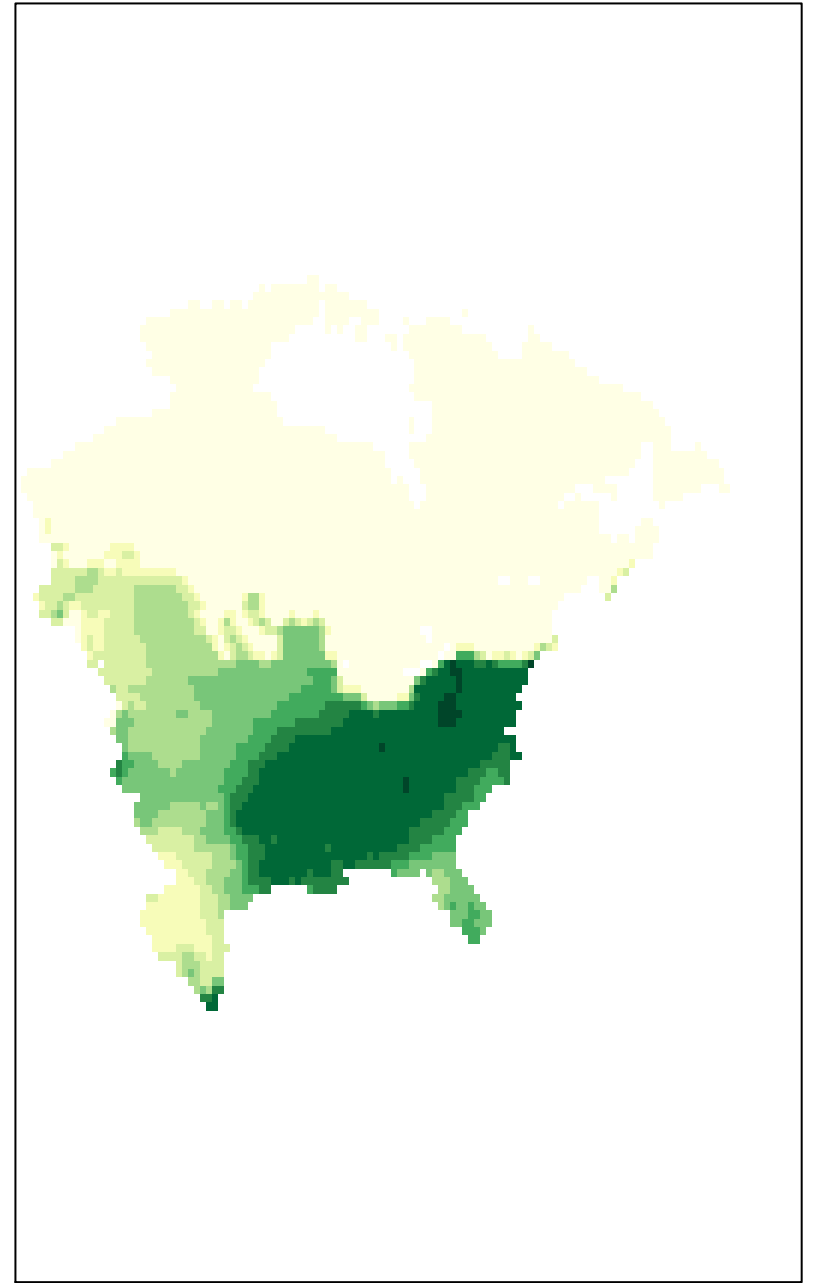


Species skipped = Fraxinus profunda, GCM = Beyer

SUM, X0.ybp



SUM, X0.ybp



Species skipped = Fraxinus profunda, GCM = Beyer