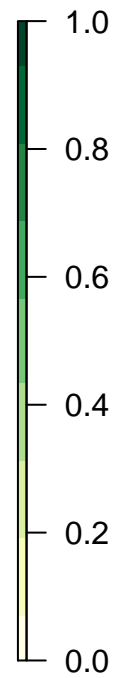
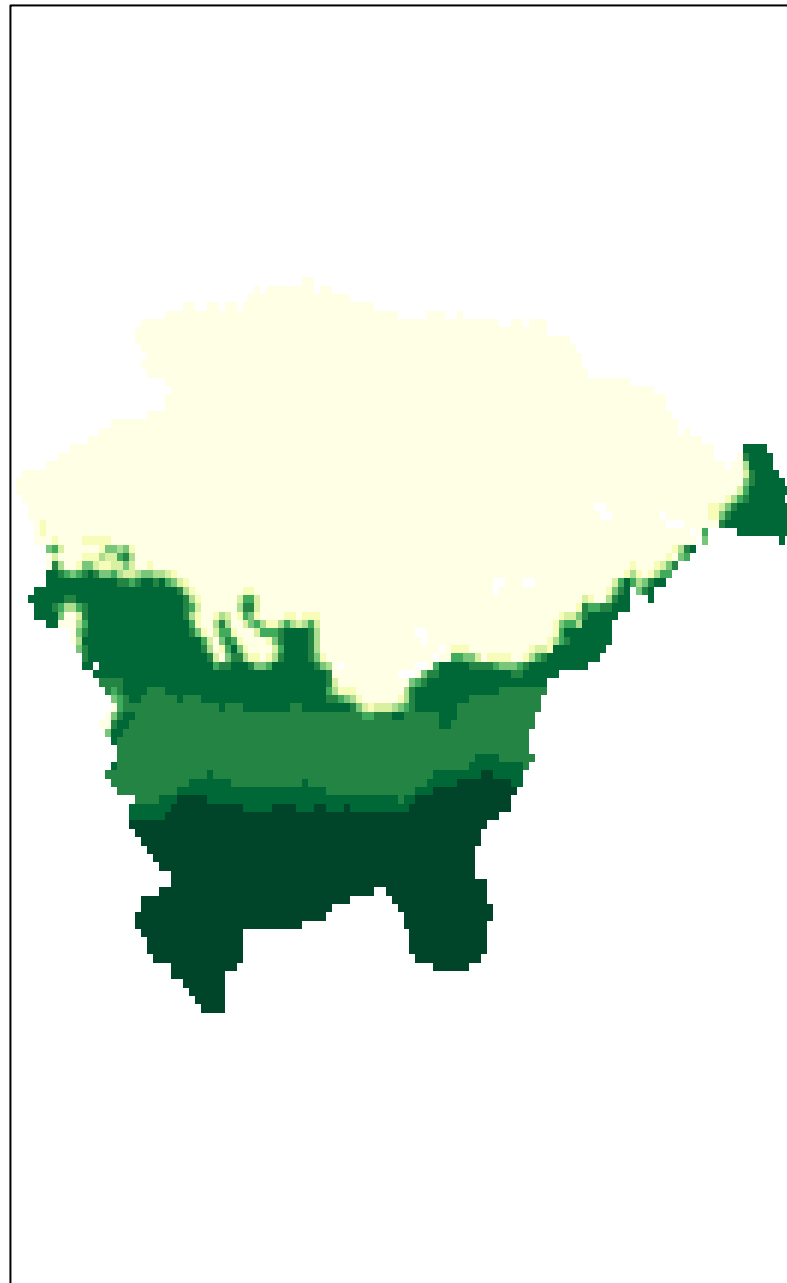
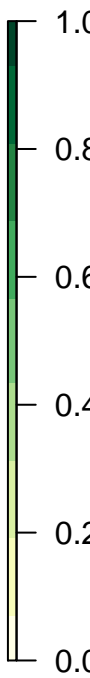
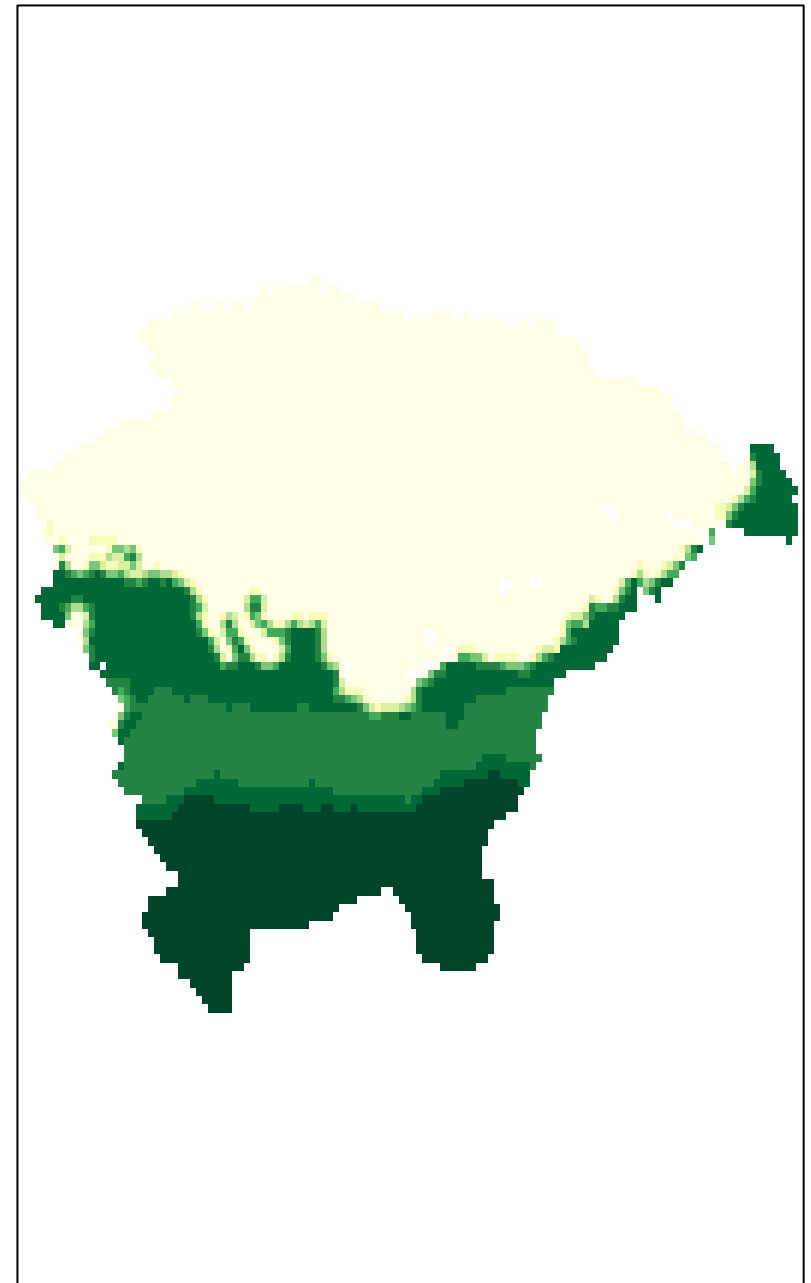


MAX, X21000.ybp

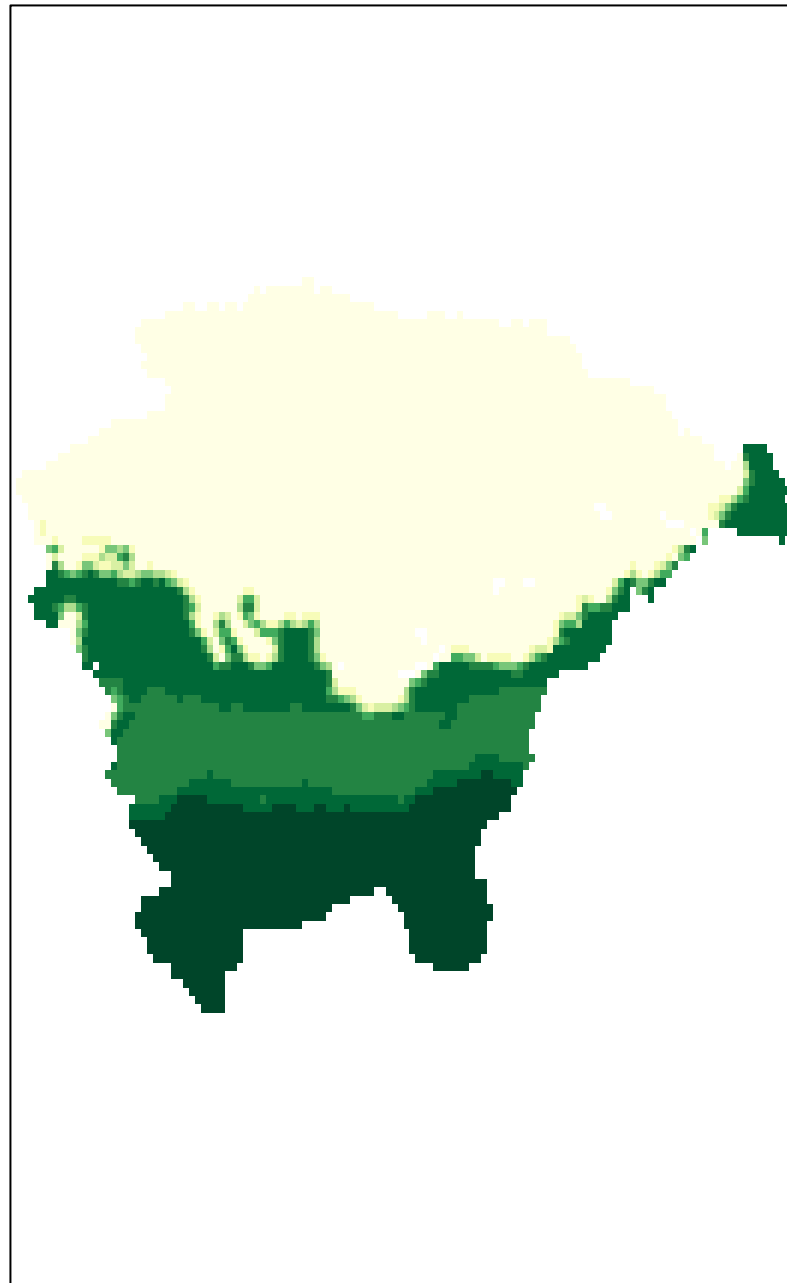


MAX, X21000.ybp

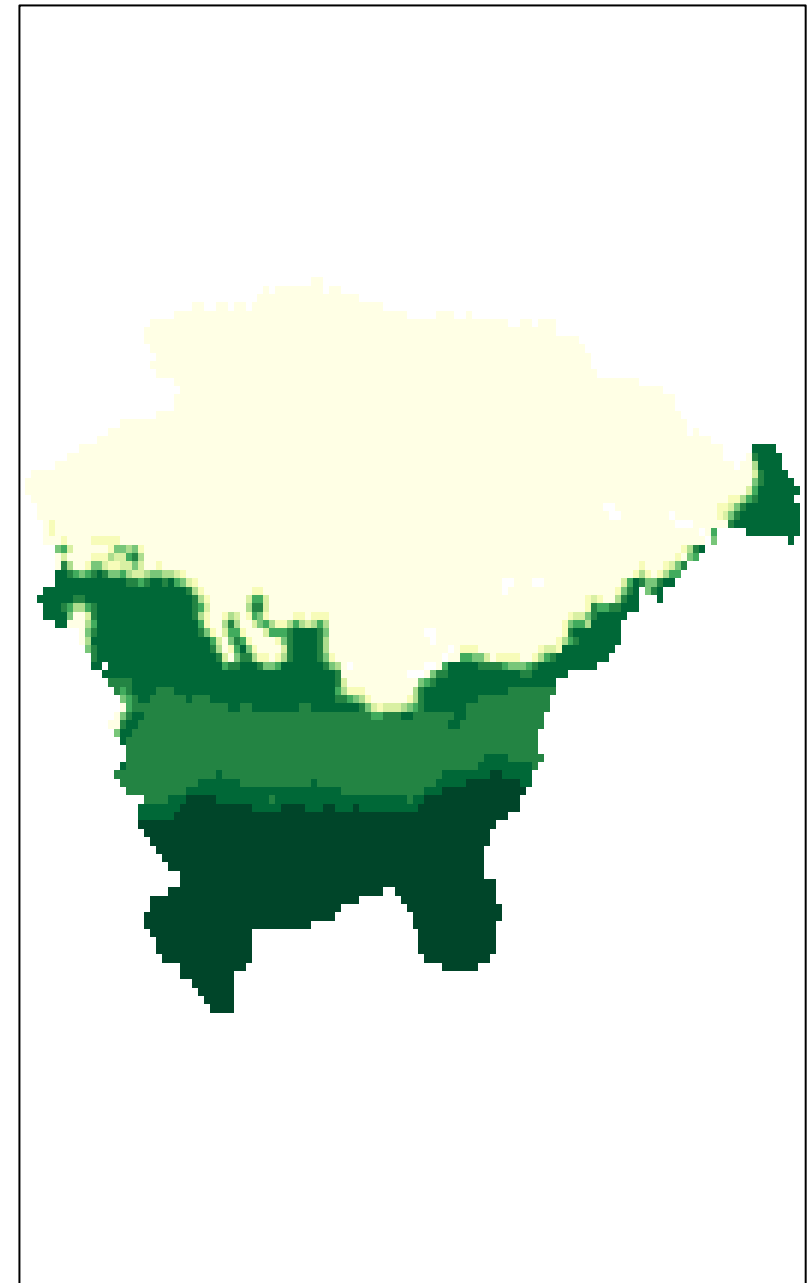


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X20000.ybp

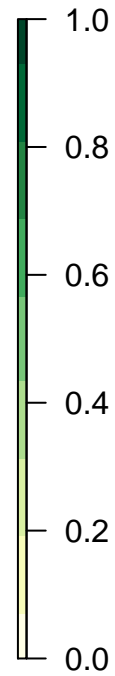
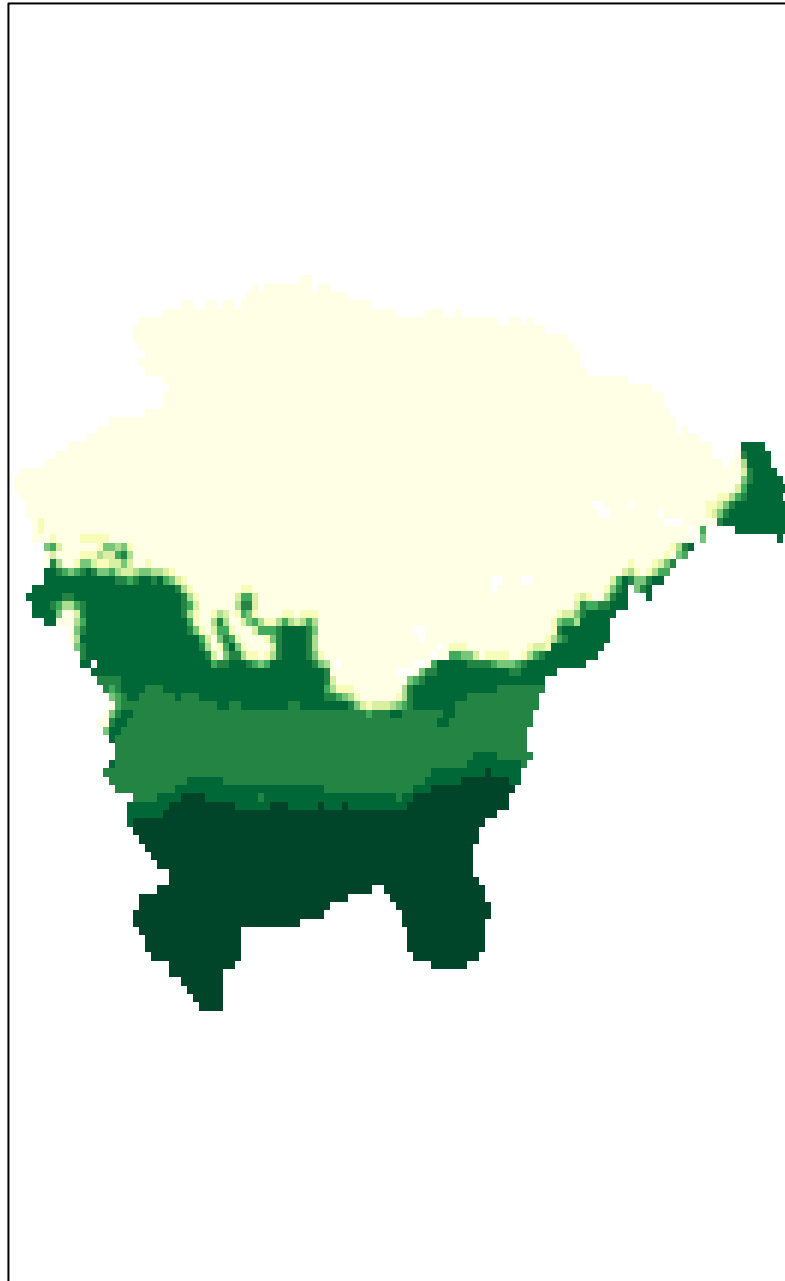


MAX, X20000.ybp

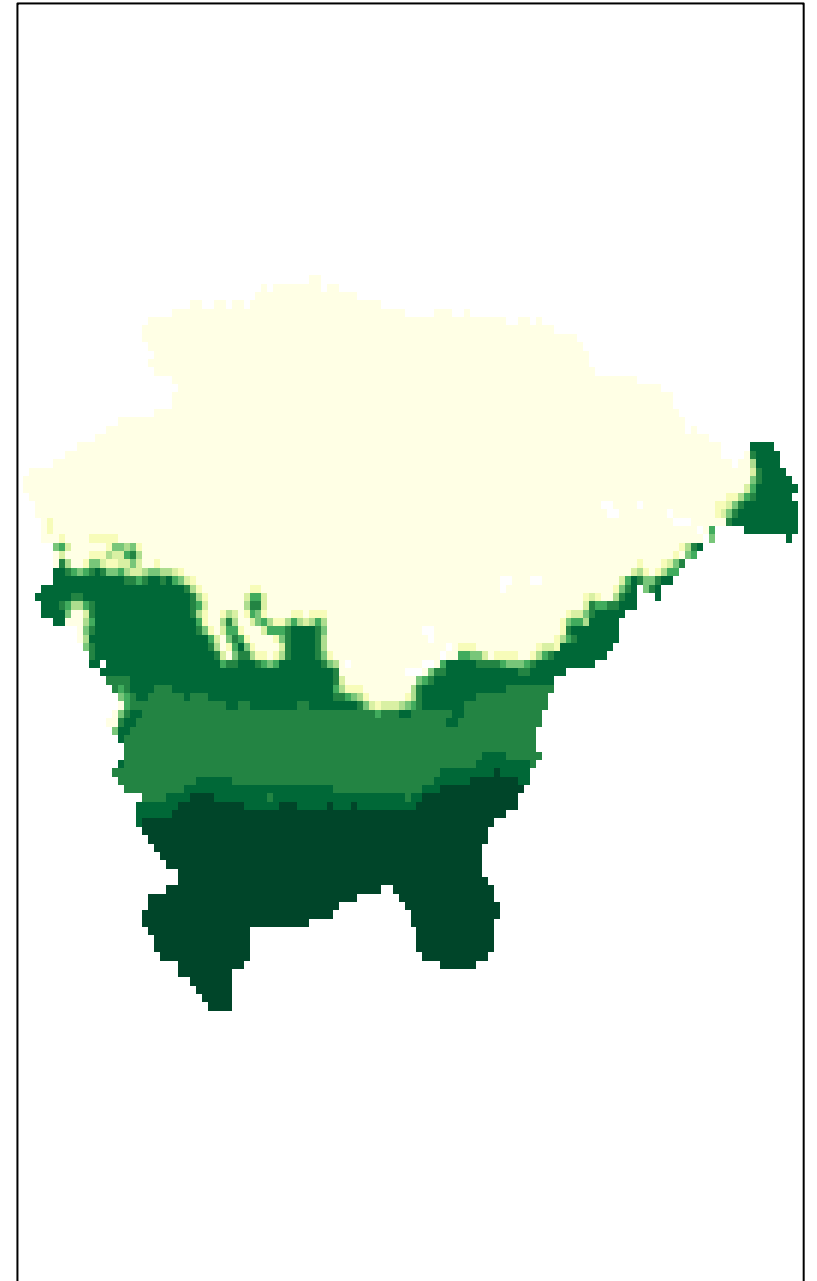


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MAX, X19000.ybp

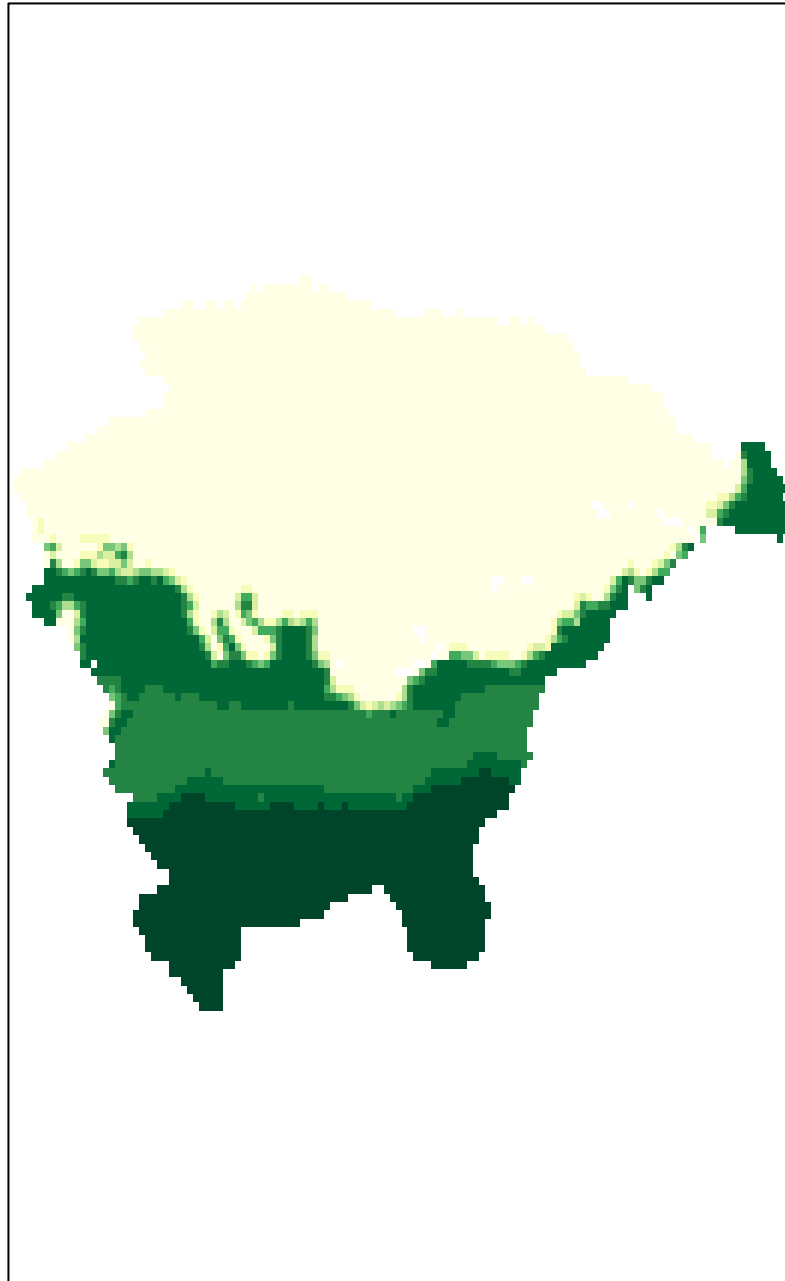


MAX, X19000.ybp

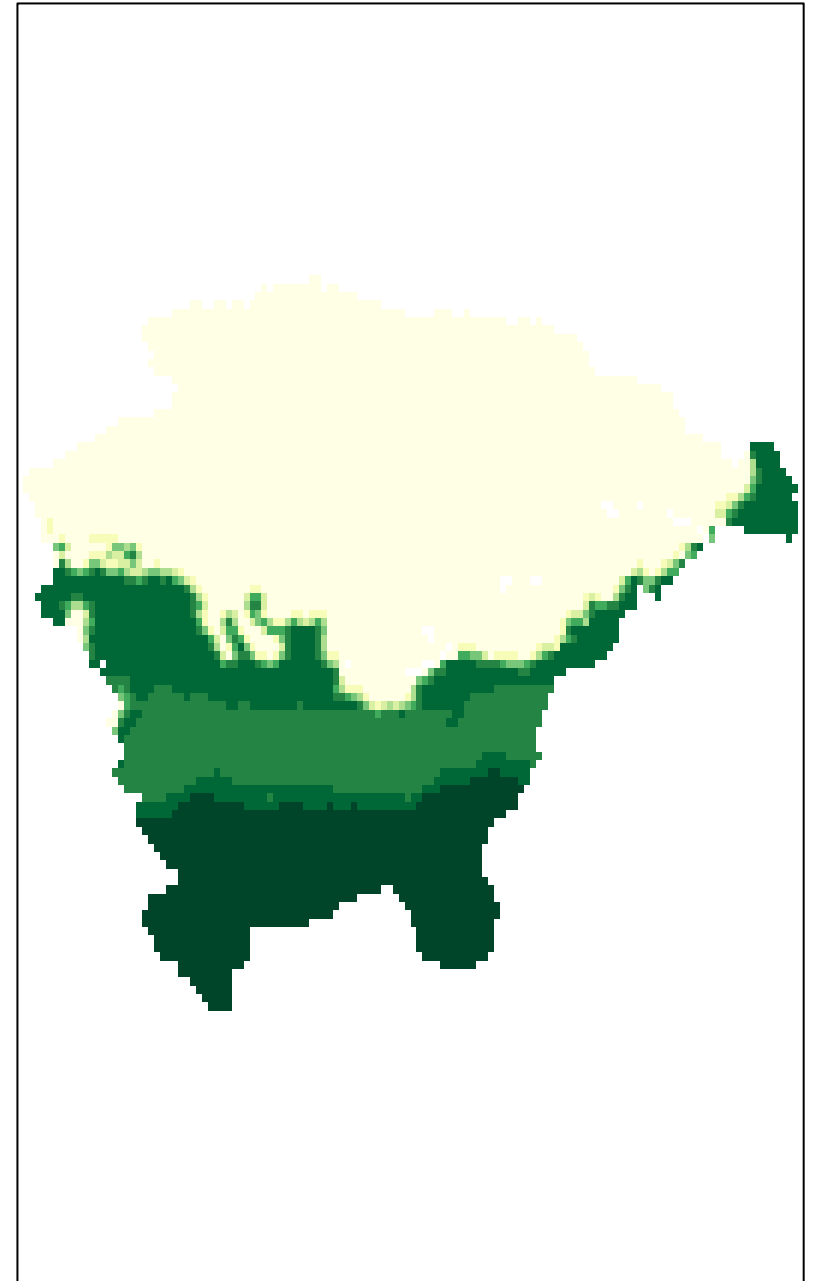


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MAX, X18000.ybp

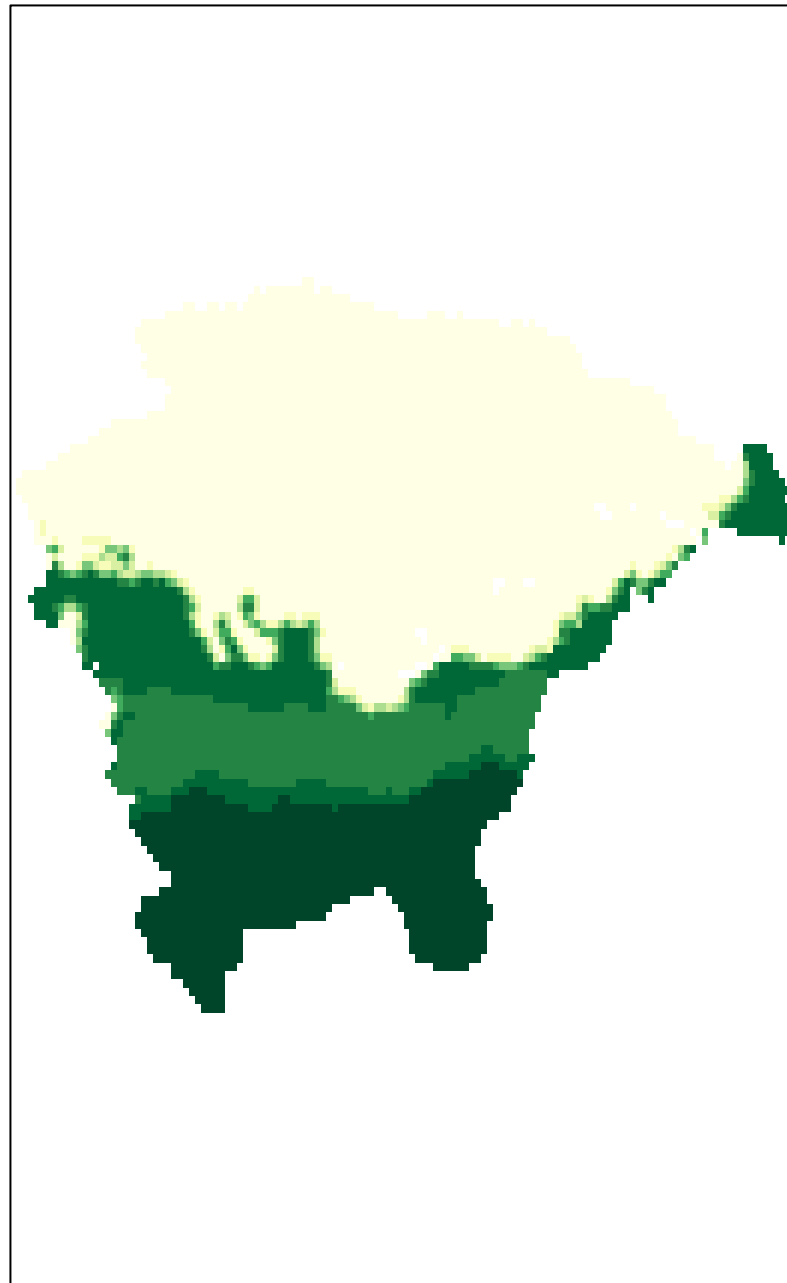


MAX, X18000.ybp

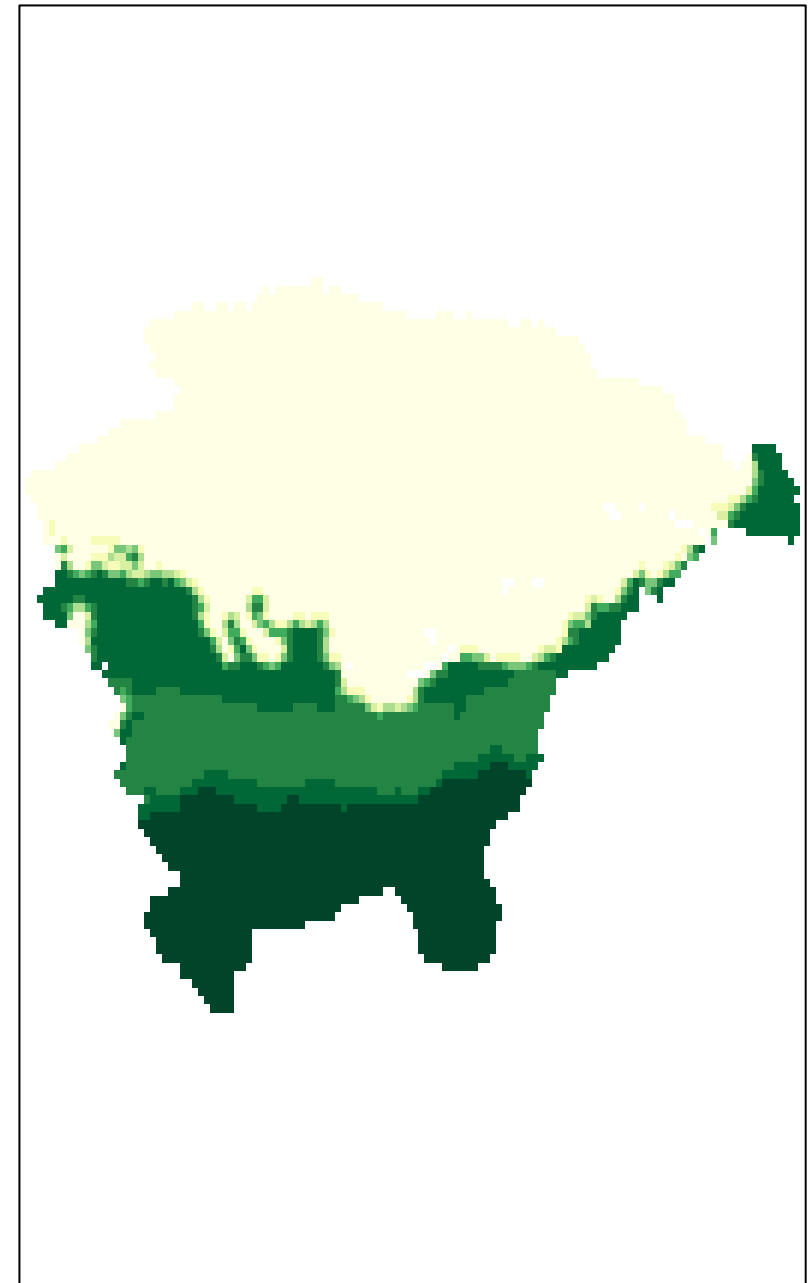


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X17000.ybp

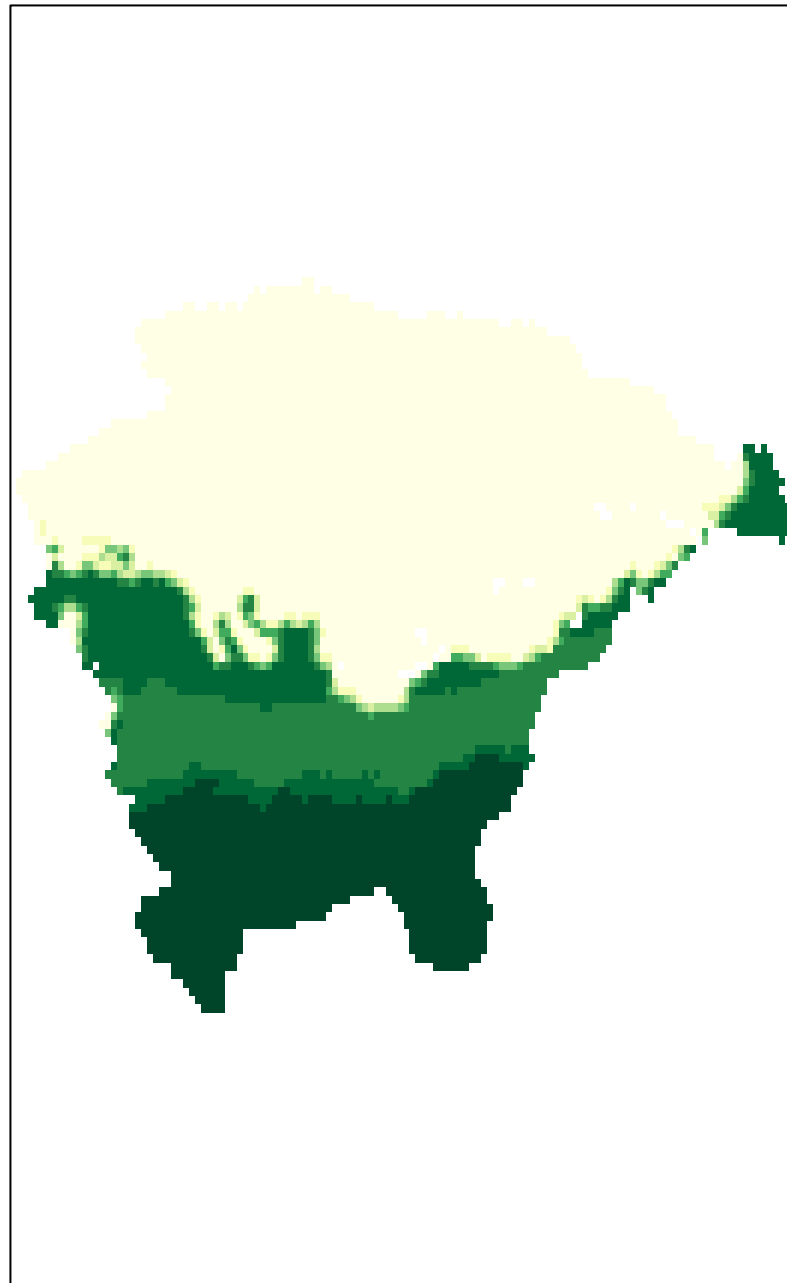


MAX, X17000.ybp

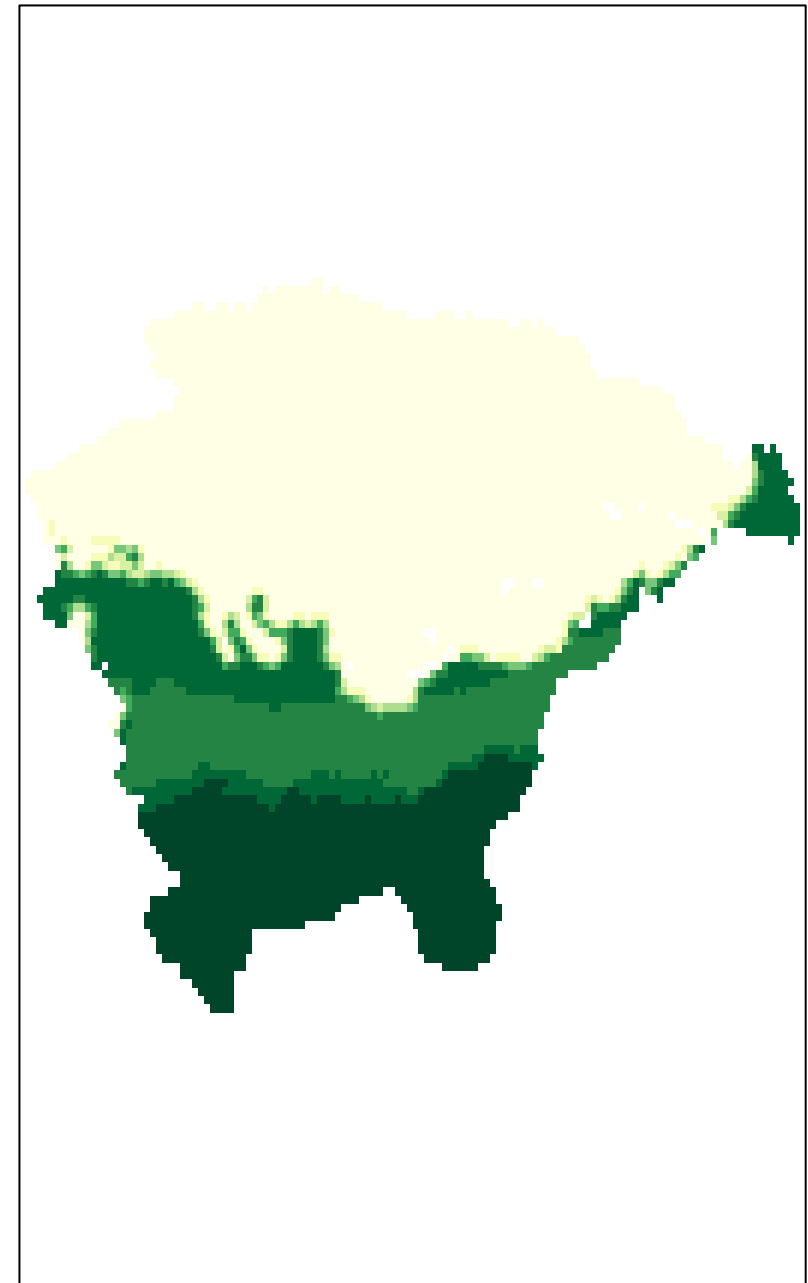


Species skipped = *Fraxinus americana*, GCM = ecbitl

MAX, X16000.ybp

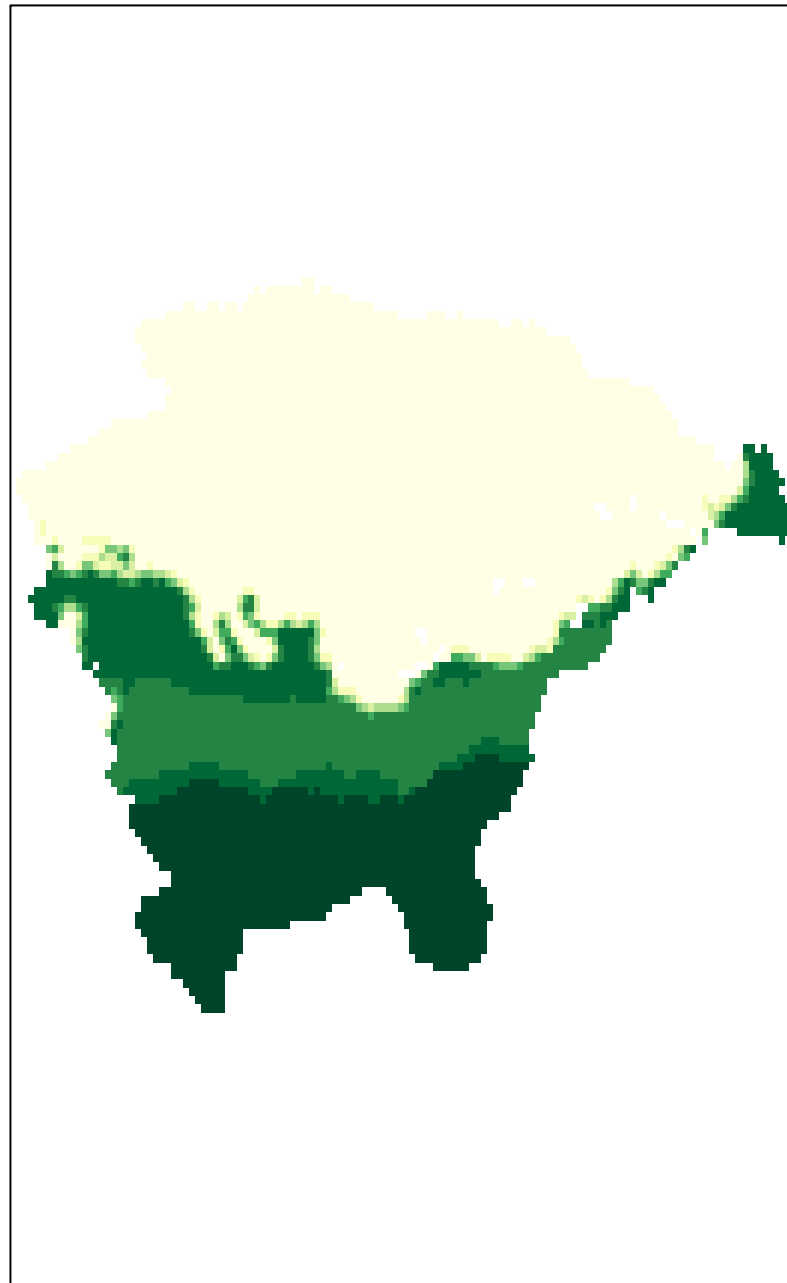


MAX, X16000.ybp

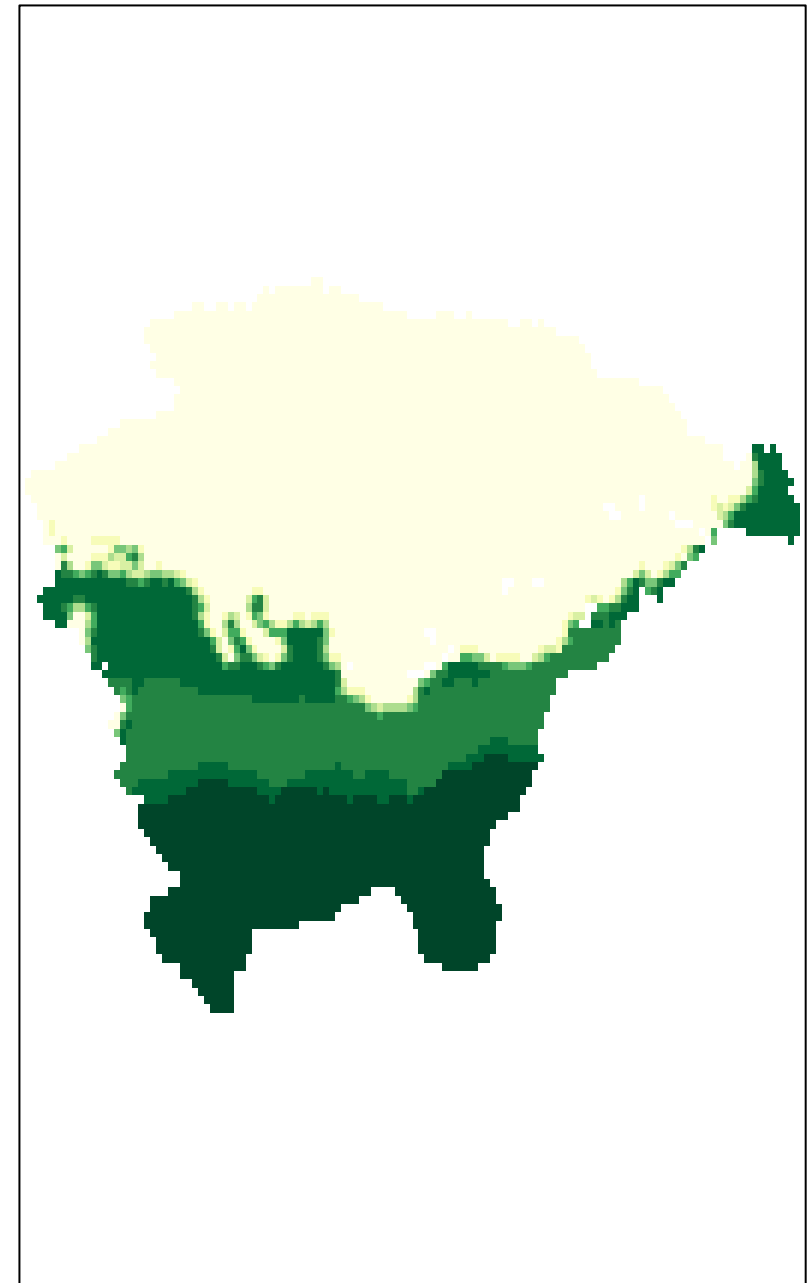


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X15000.ybp

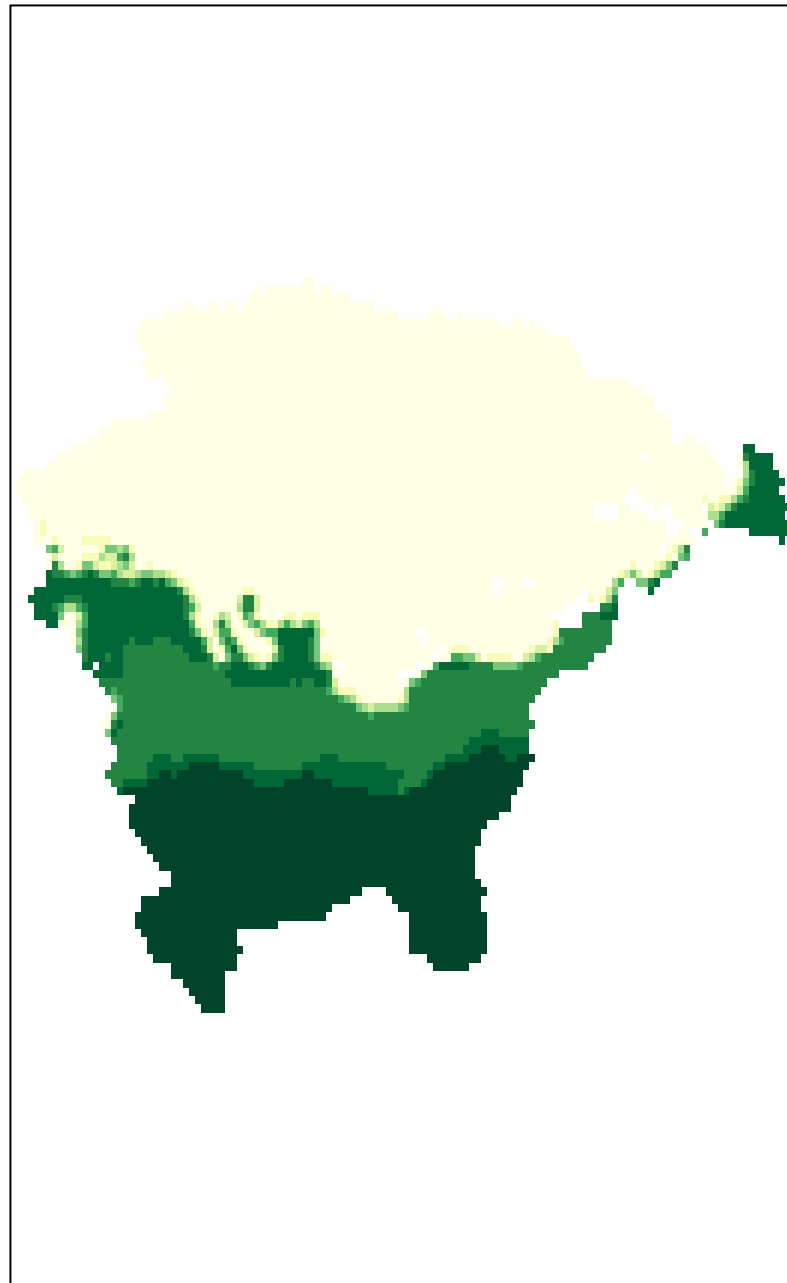


MAX, X15000.ybp

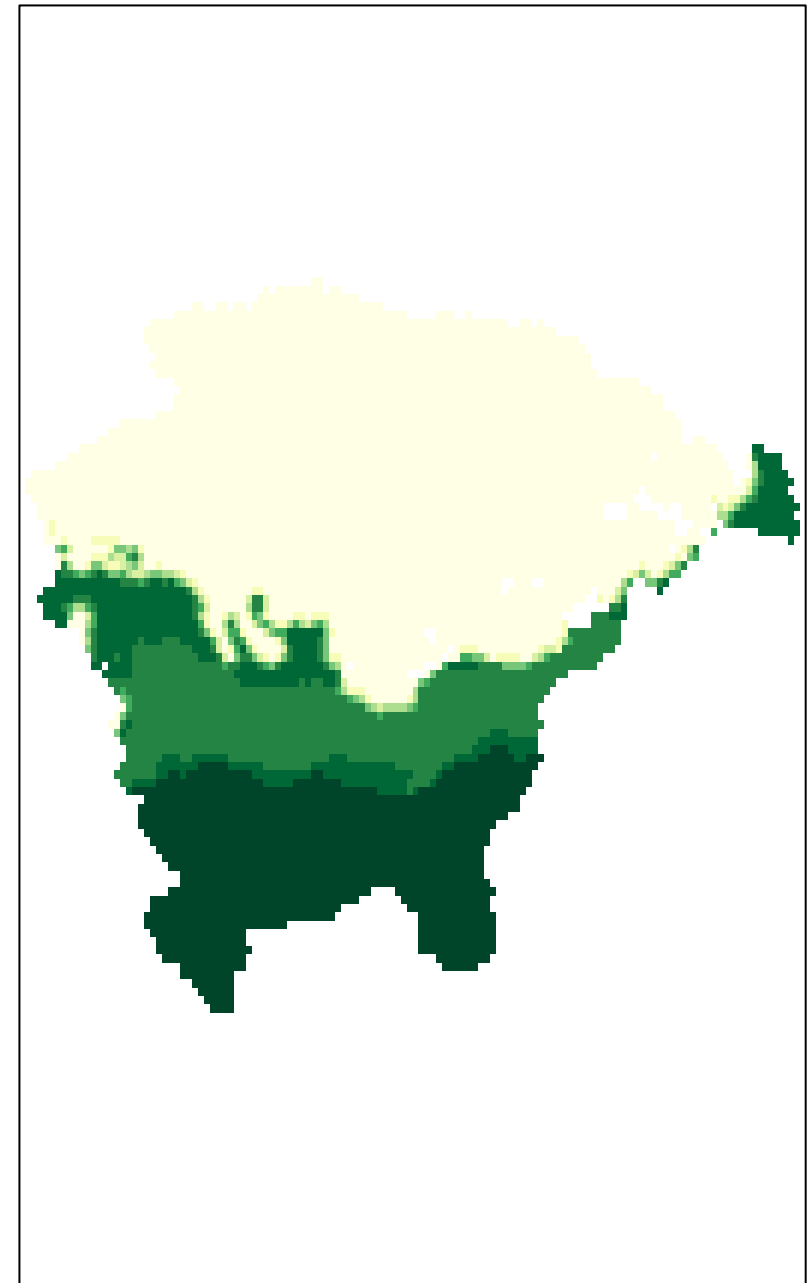


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X14000.ybp



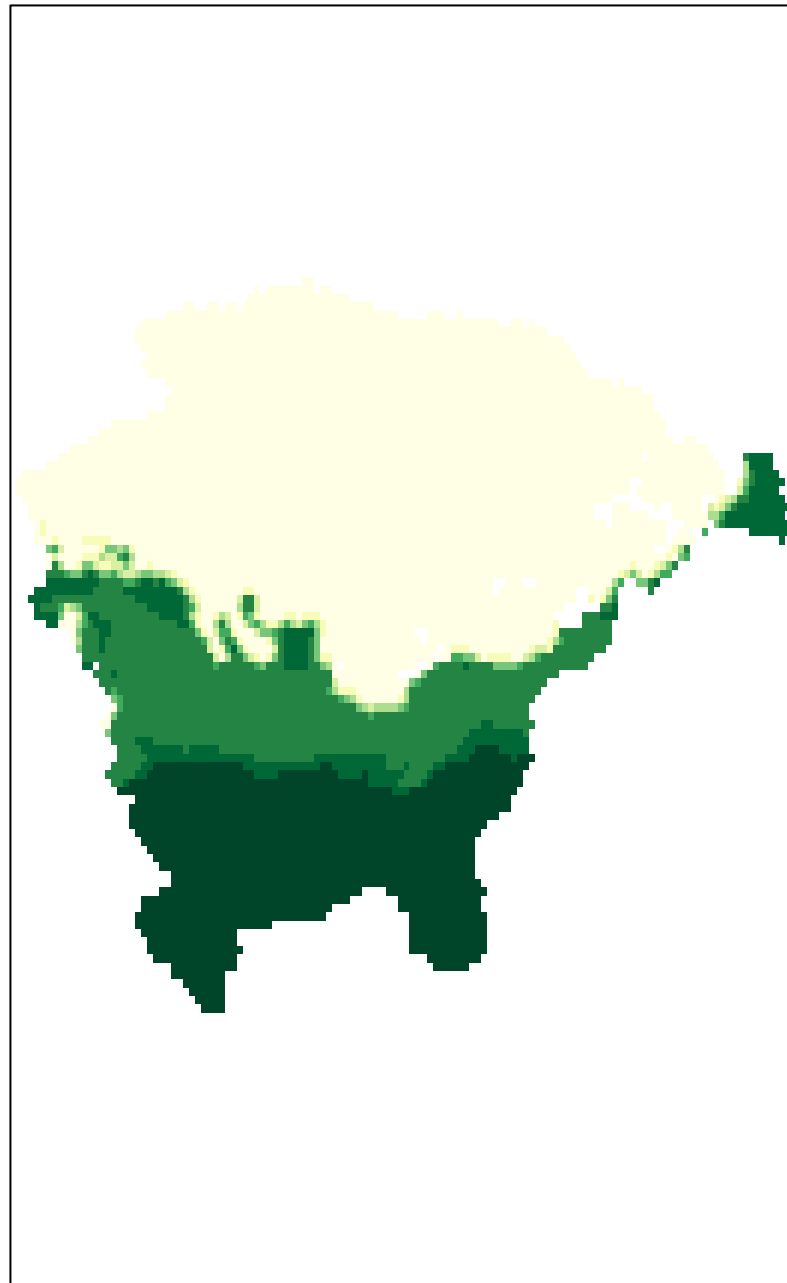
MAX, X14000.ybp



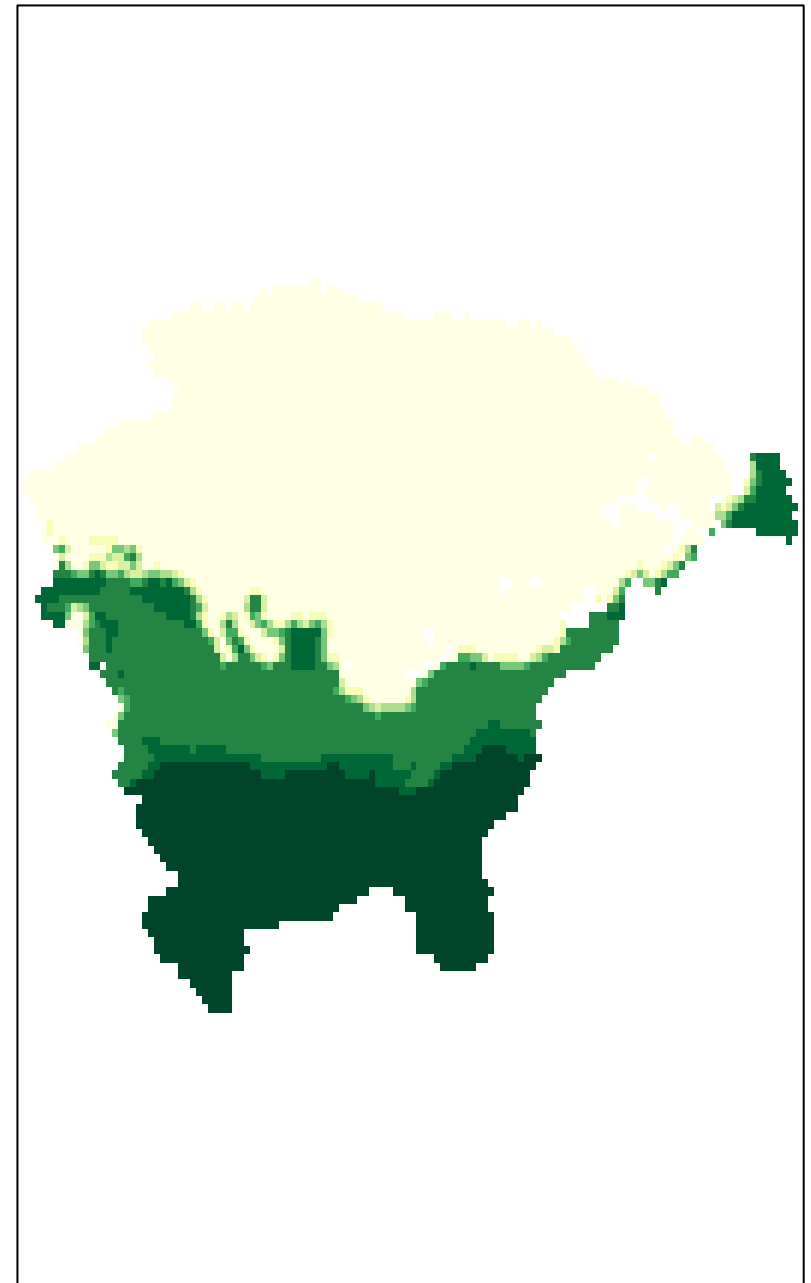
Species skipped = Fraxinus americana, GCM = ecbilt



MAX, X13000.ybp

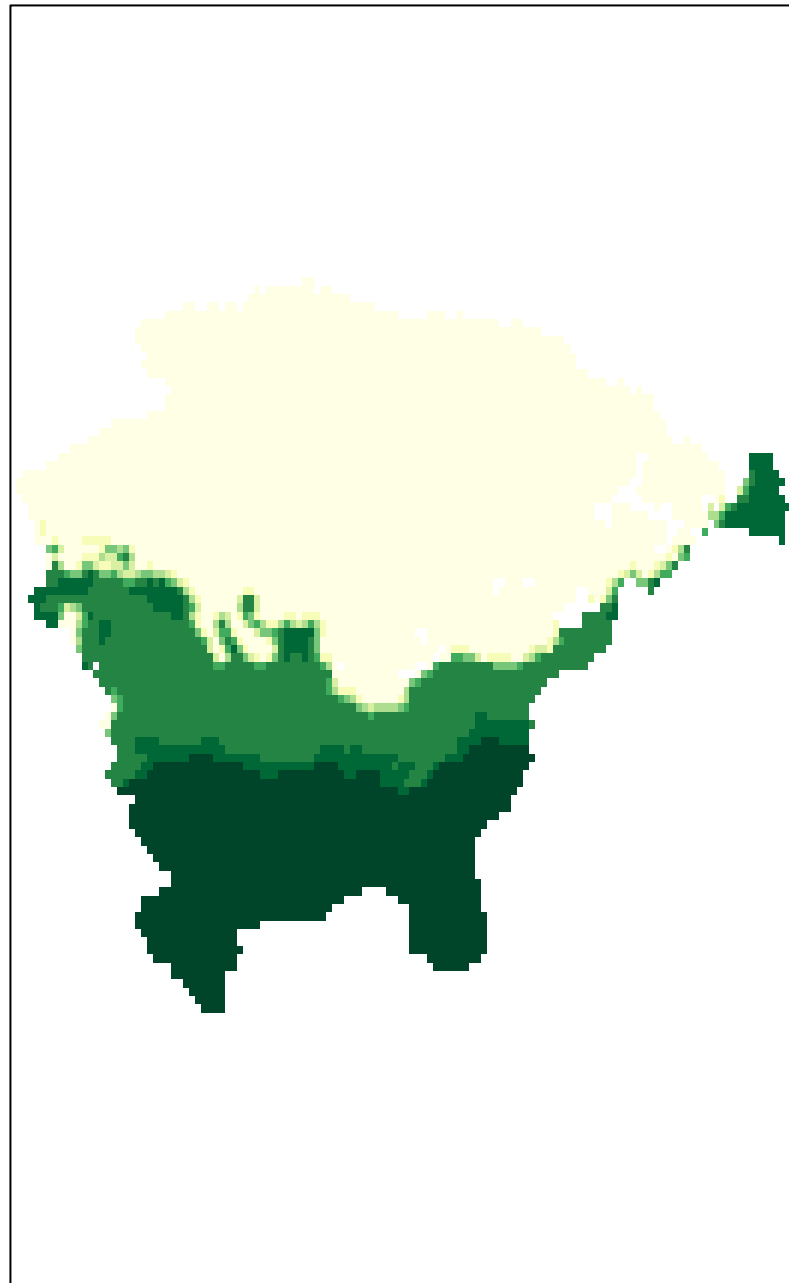


MAX, X13000.ybp

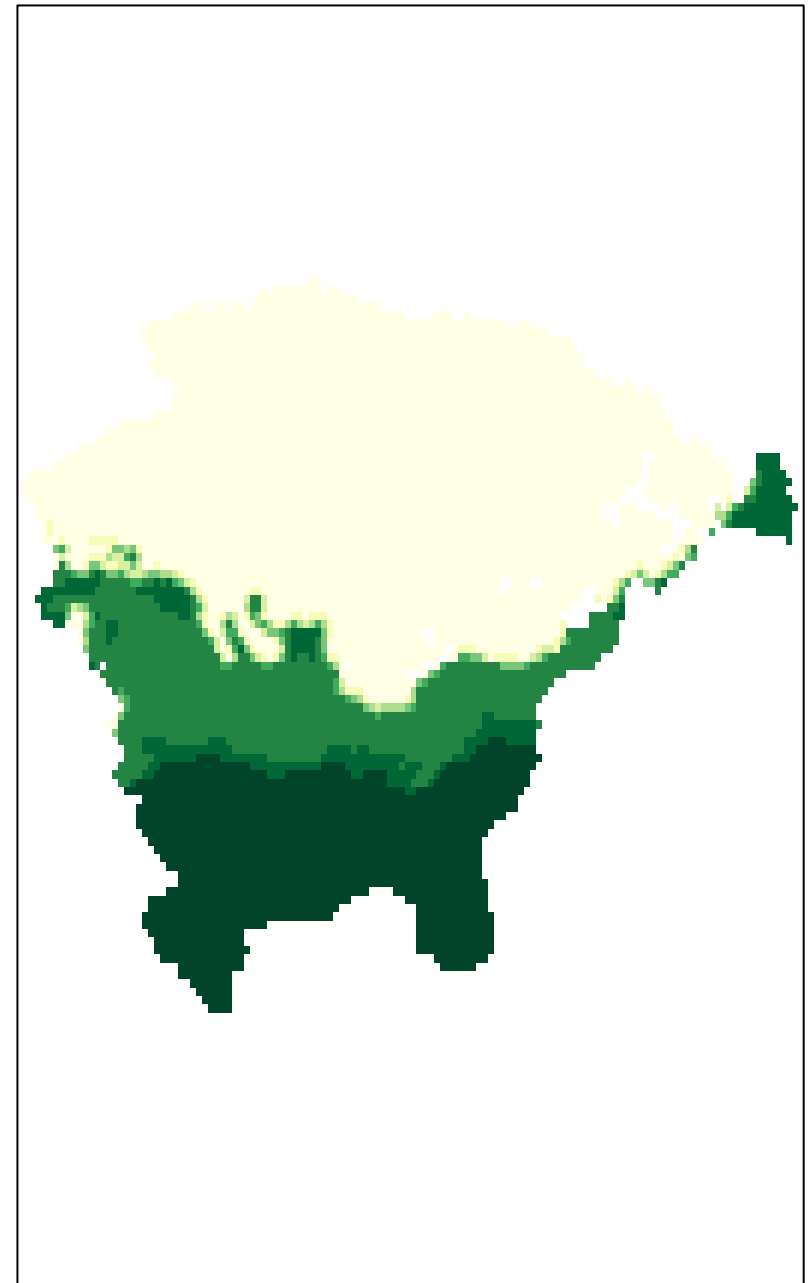


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X12000.ybp

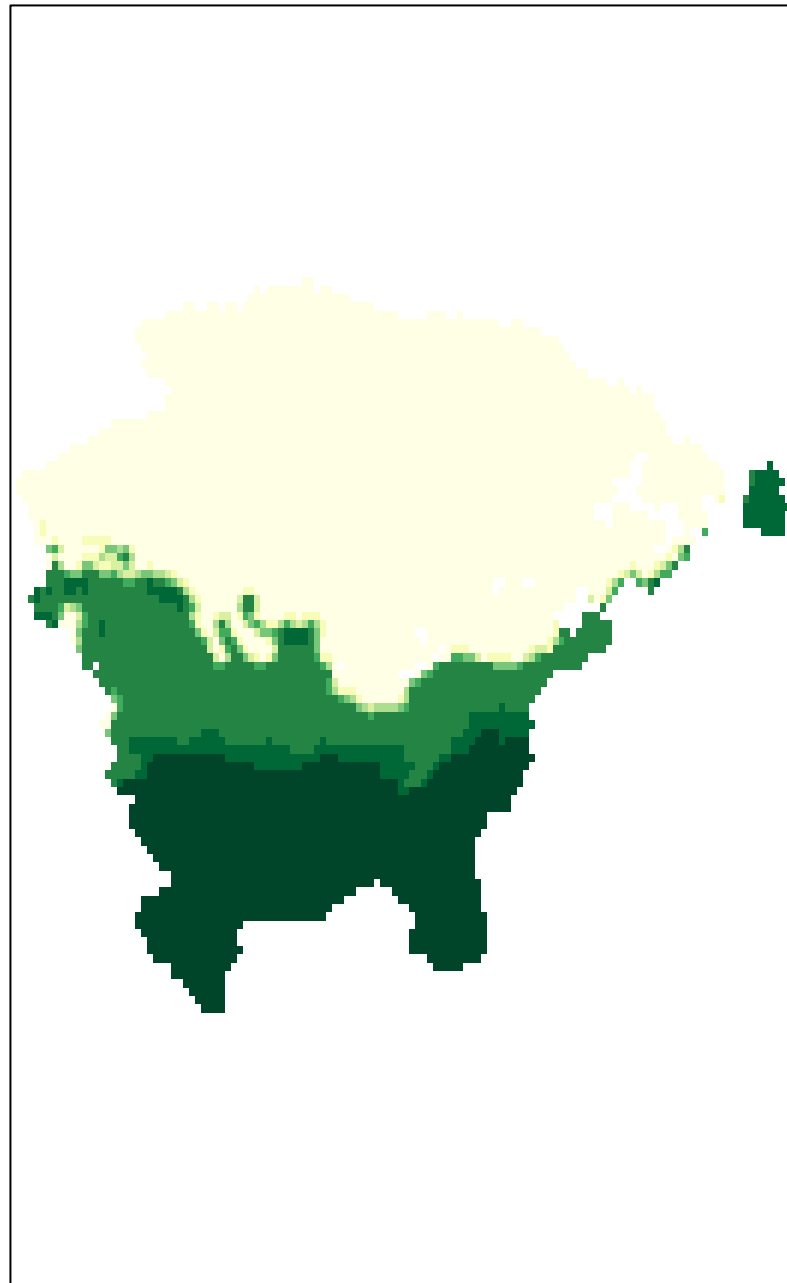


MAX, X12000.ybp

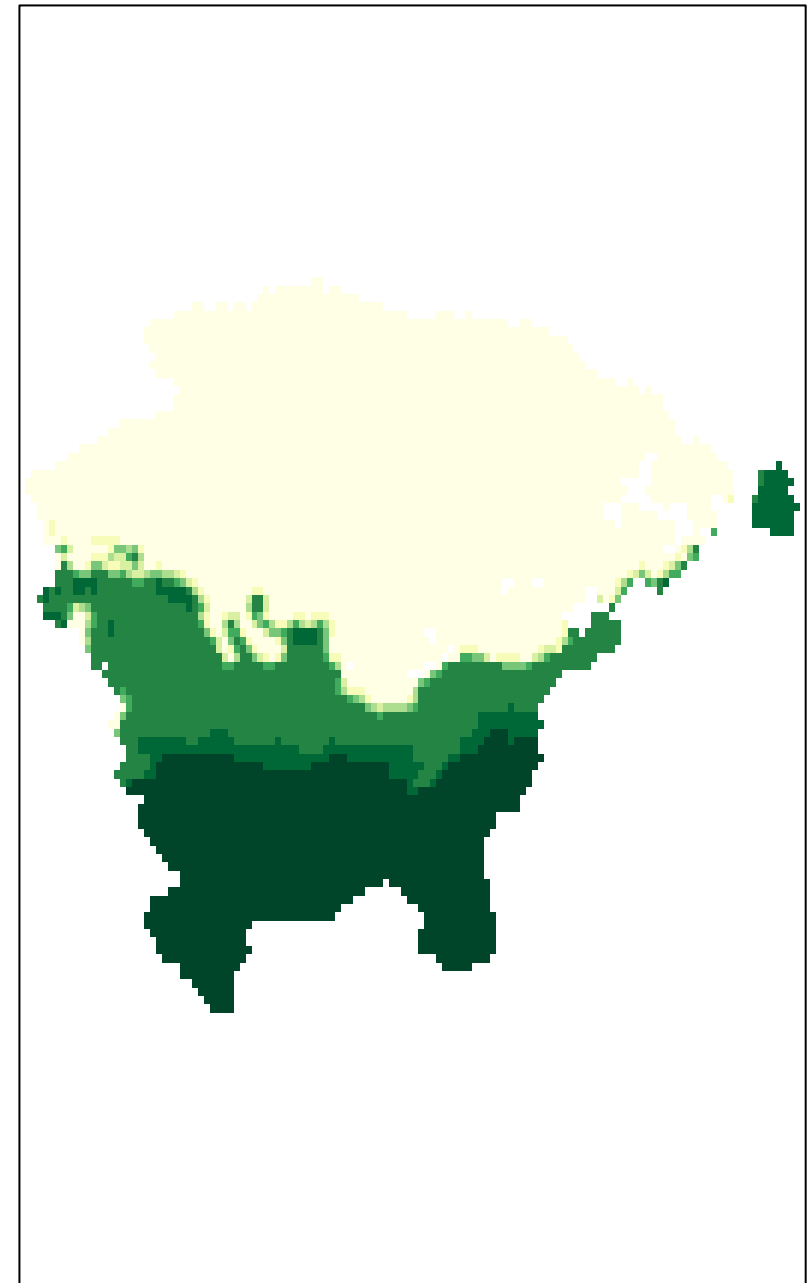


Species skipped = Fraxinus americana, GCM = ecbilt

MAX, X11000.ybp

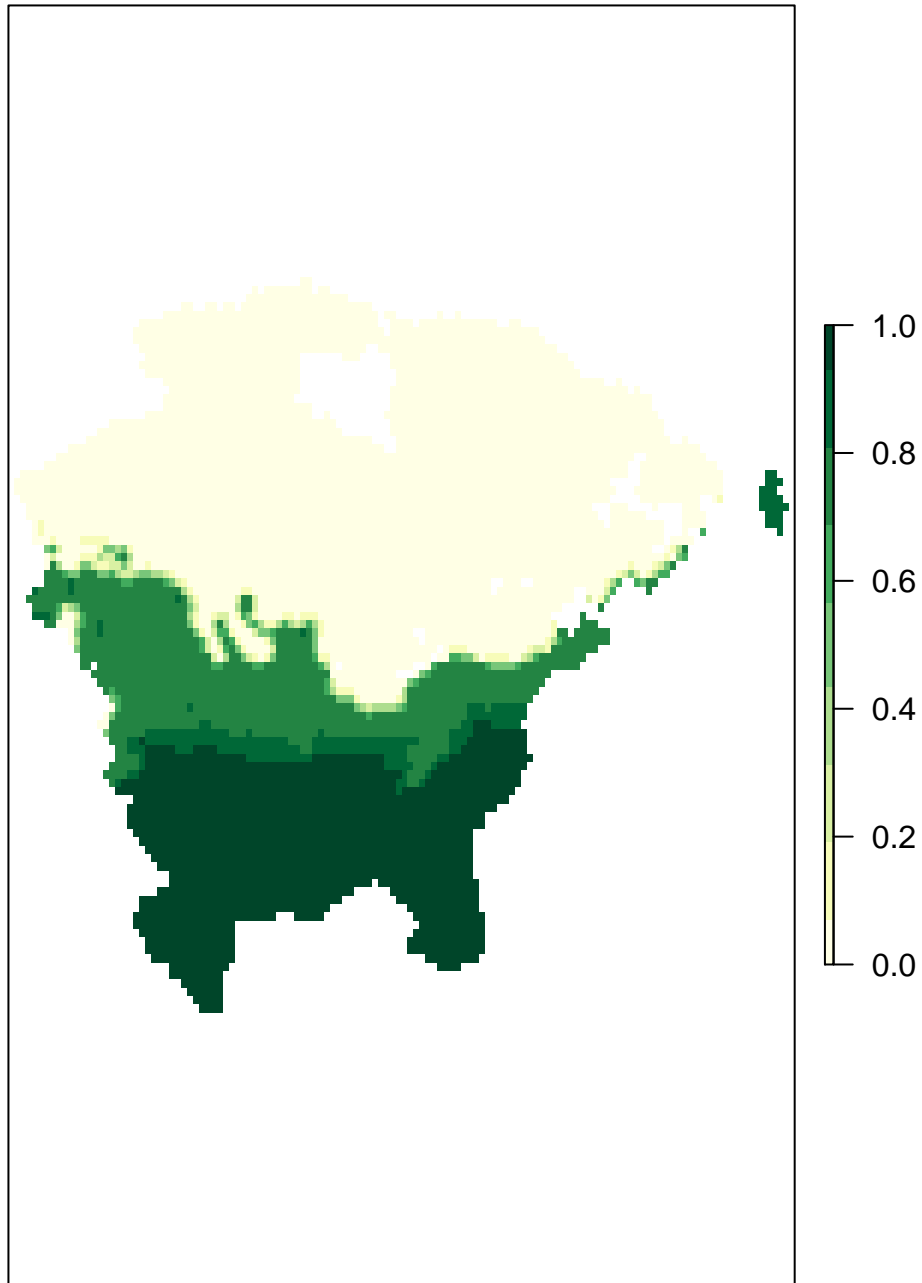


MAX, X11000.ybp

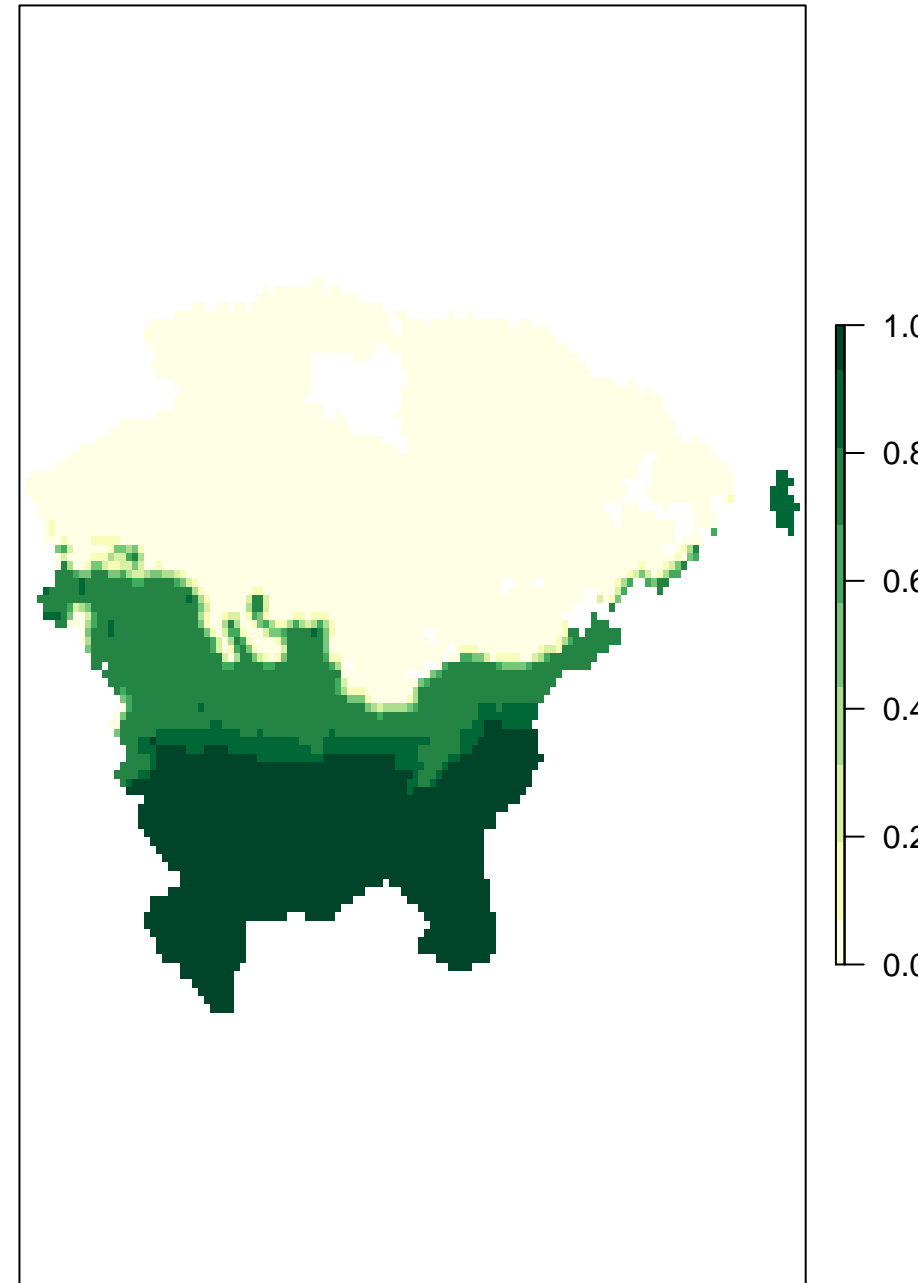


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X10000.ybp

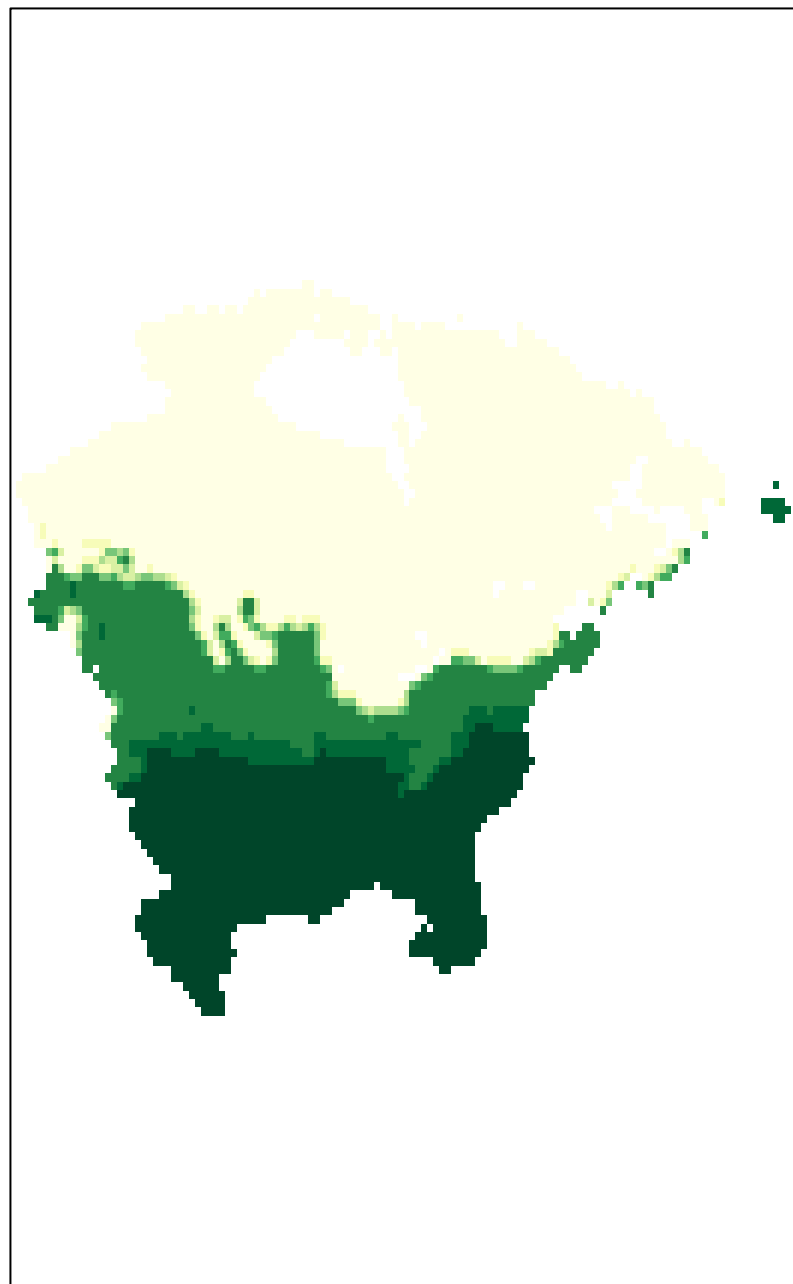


MAX, X10000.ybp

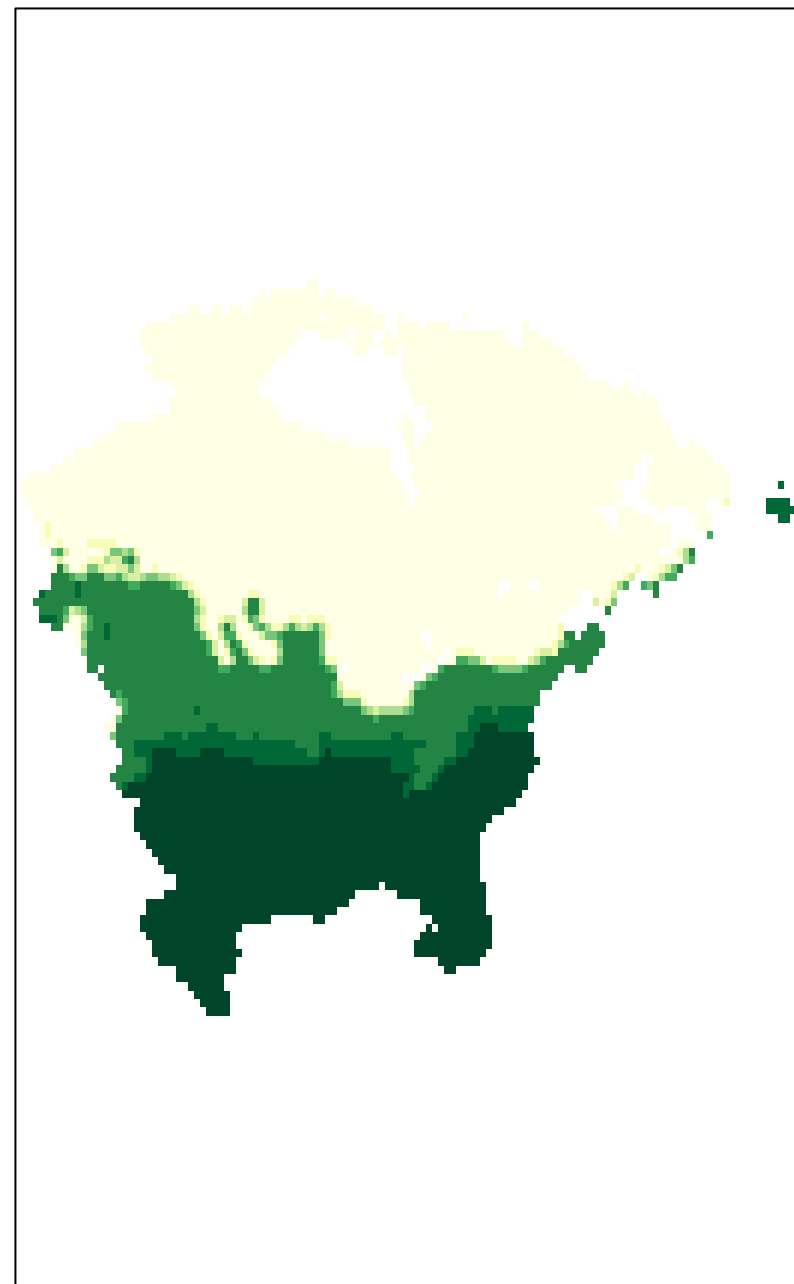


Species skipped = Fraxinus americana, GCM = ecbilt

MAX, X9000.ybp

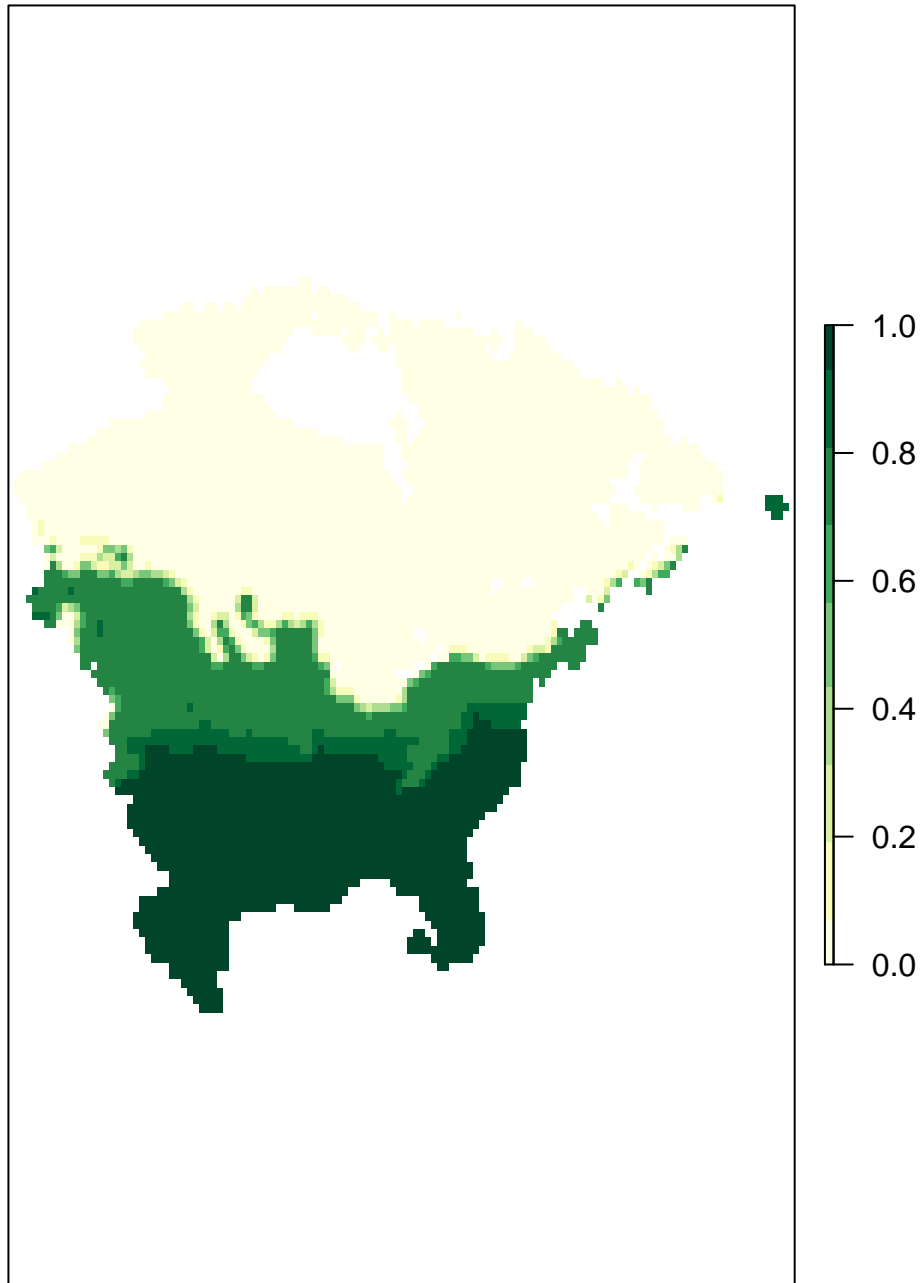


MAX, X9000.ybp

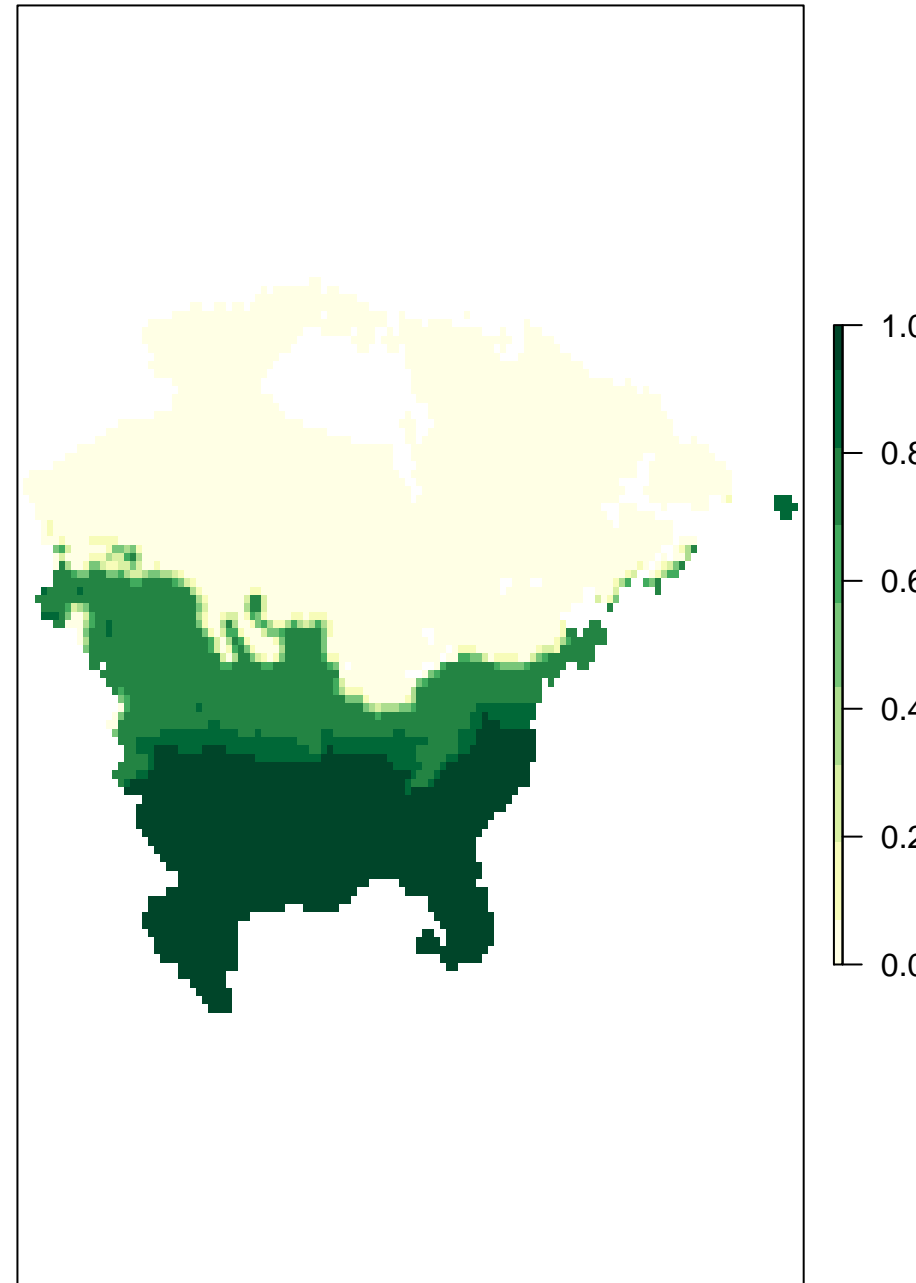


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X8000.ybp

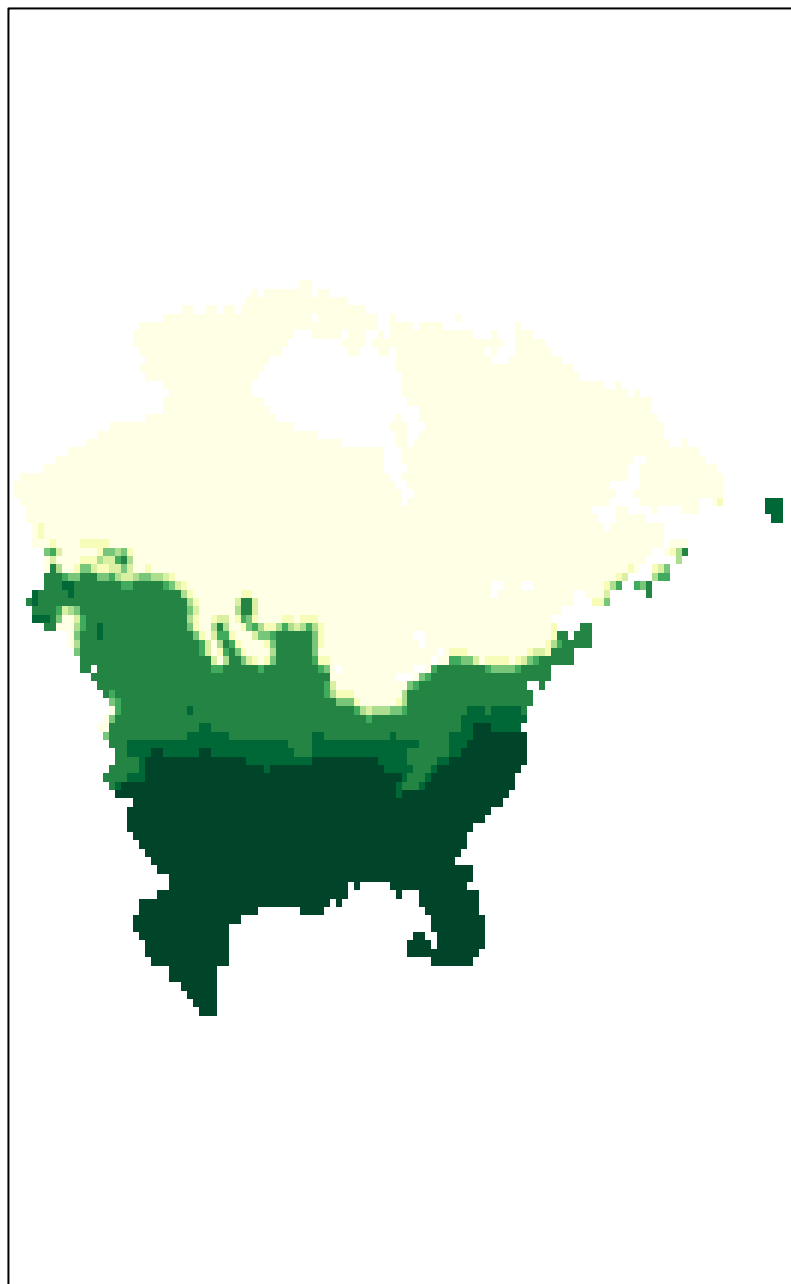


MAX, X8000.ybp

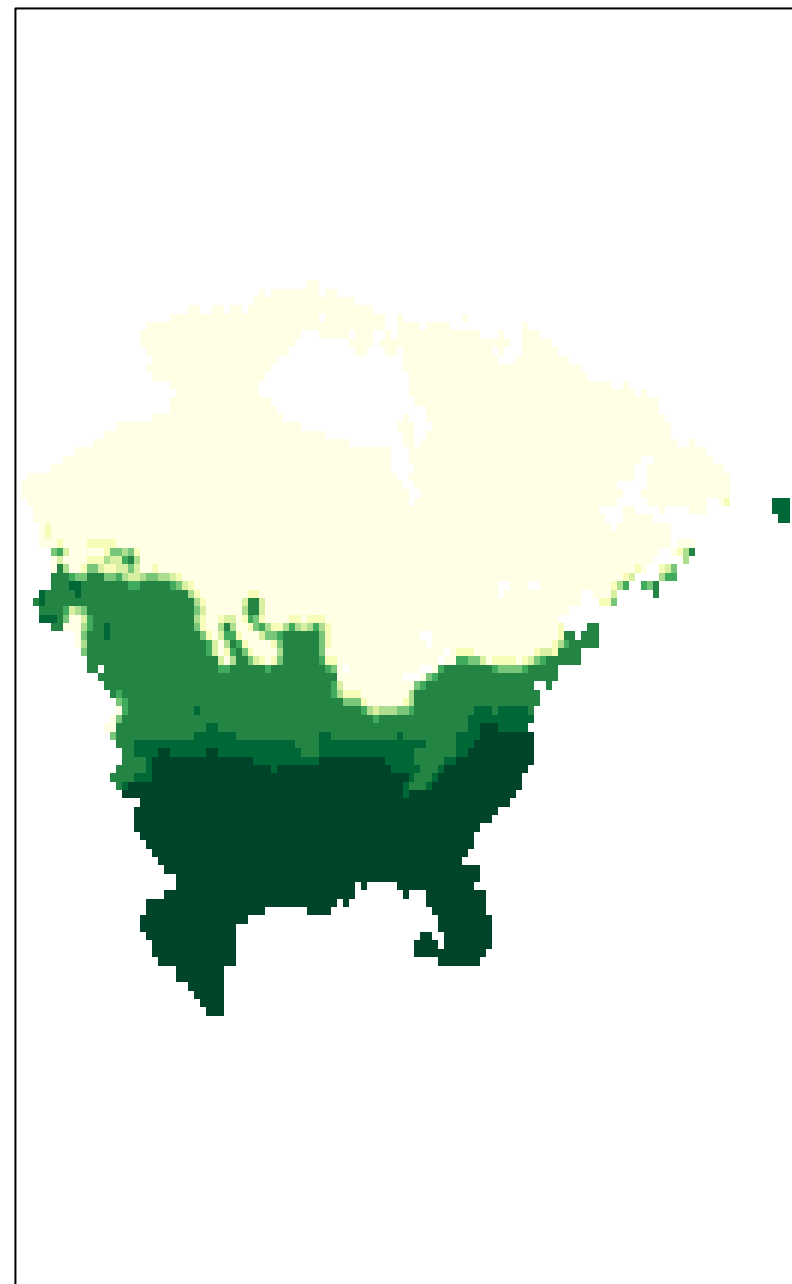


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X7000.ybp

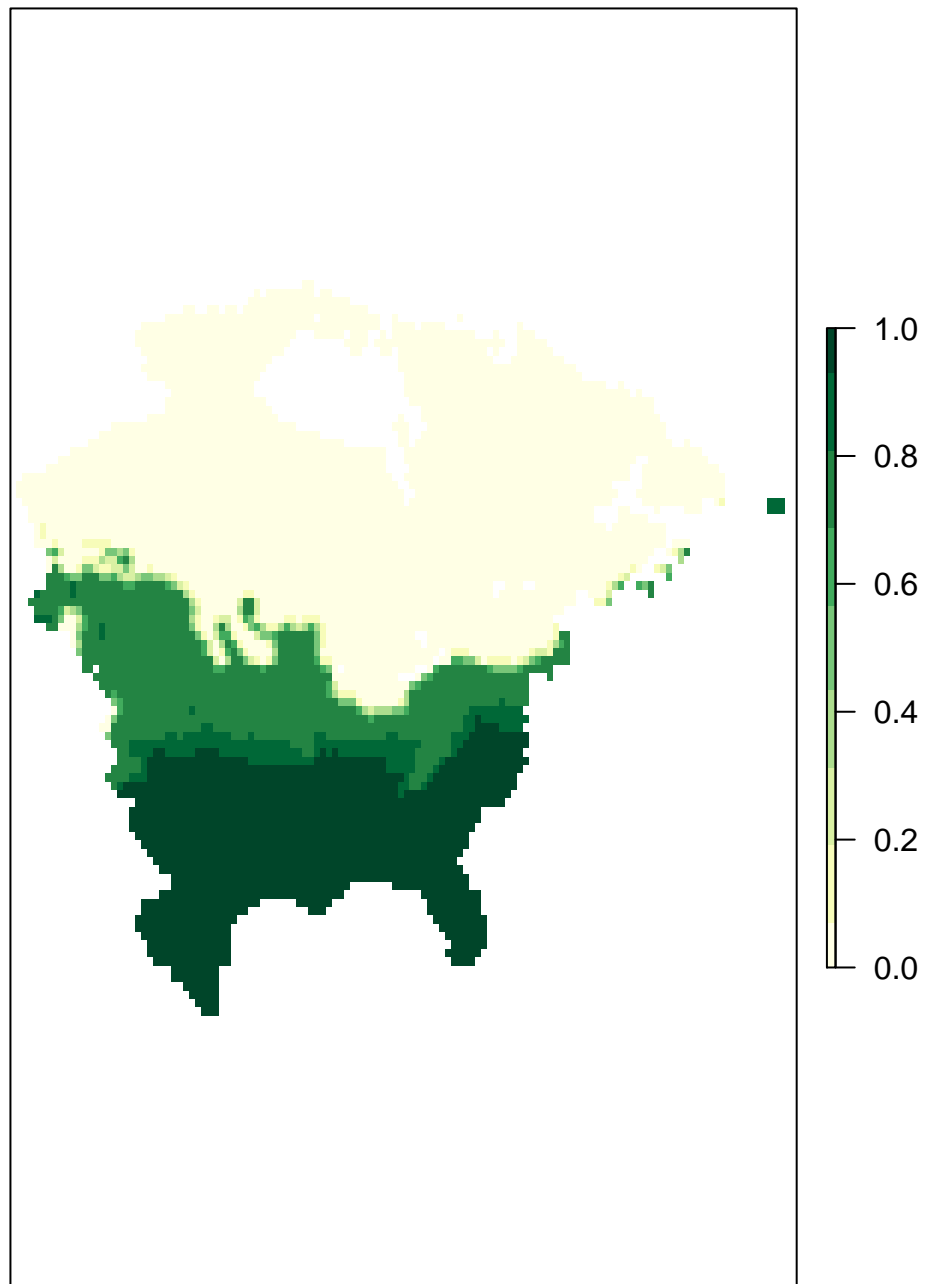


MAX, X7000.ybp

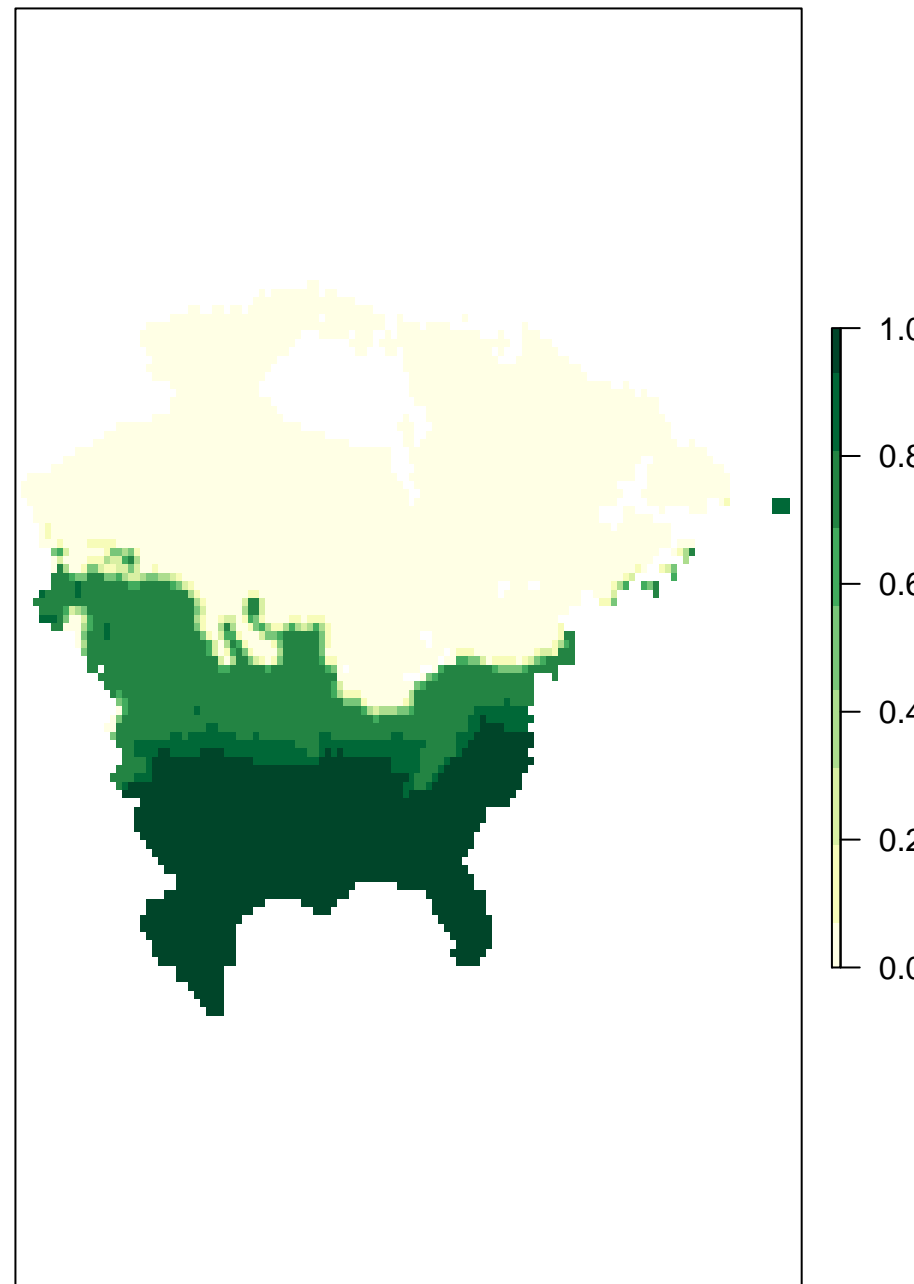


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X6000.ybp



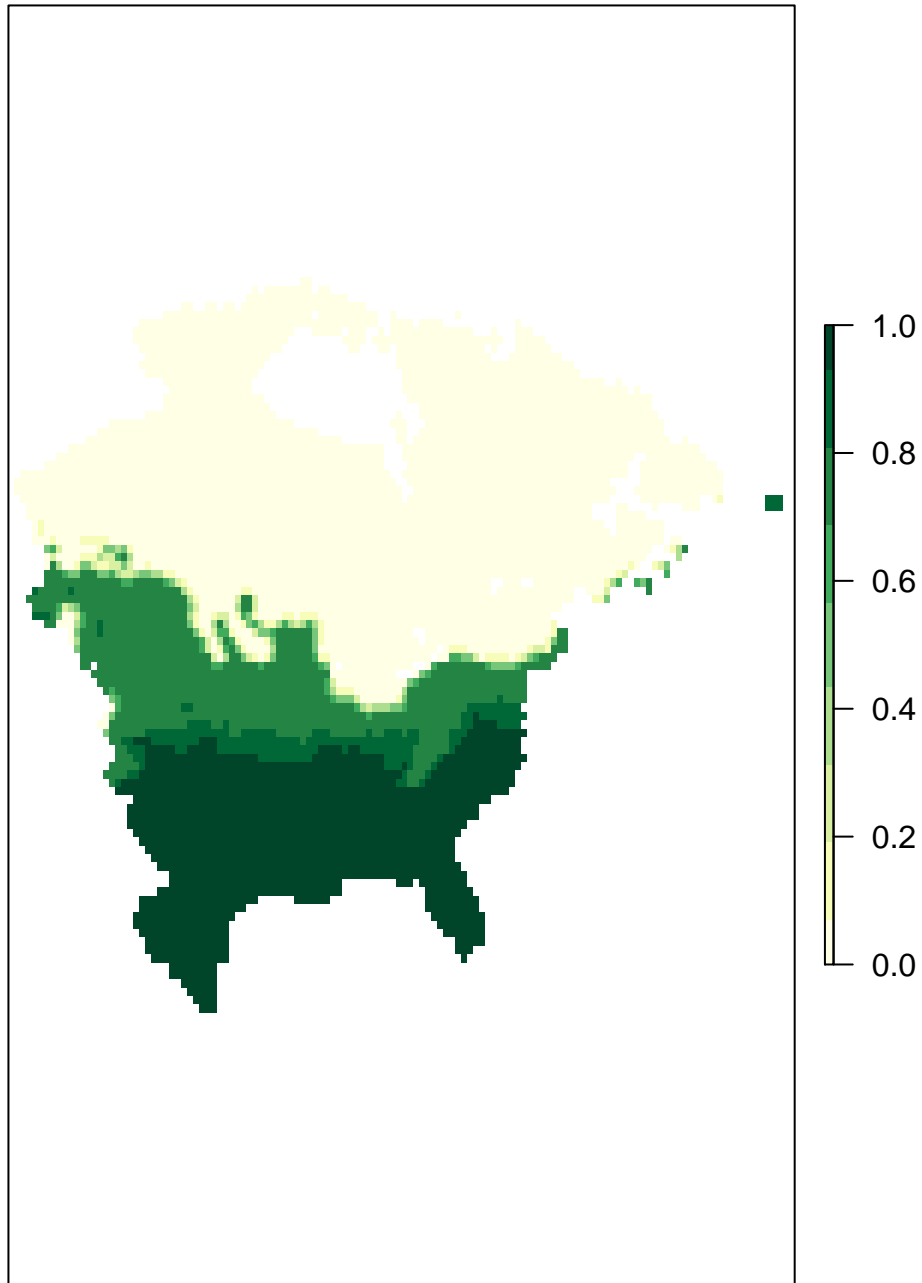
MAX, X6000.ybp



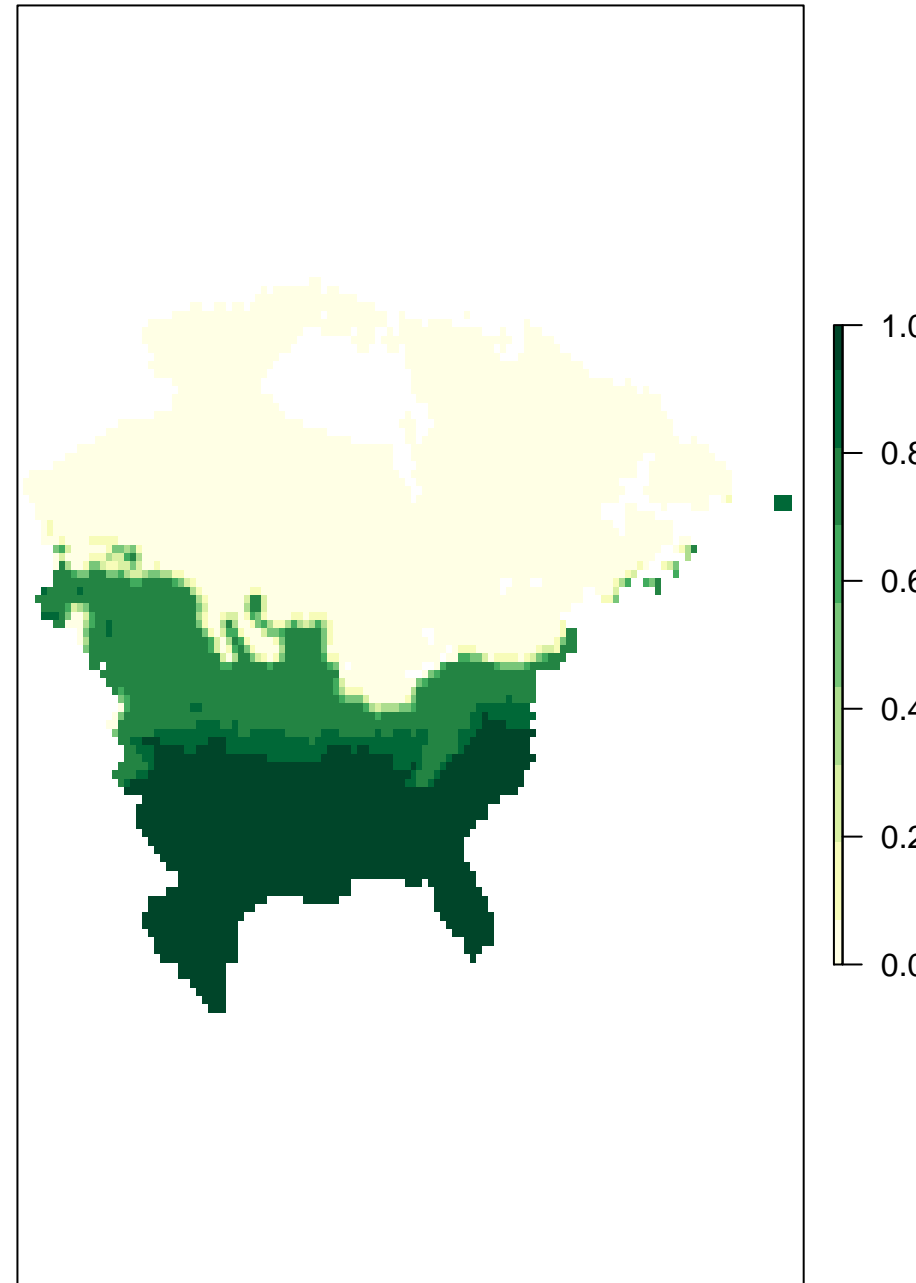
Species skipped = Fraxinus americana, GCM = ecbilt



MAX, X5000.ybp

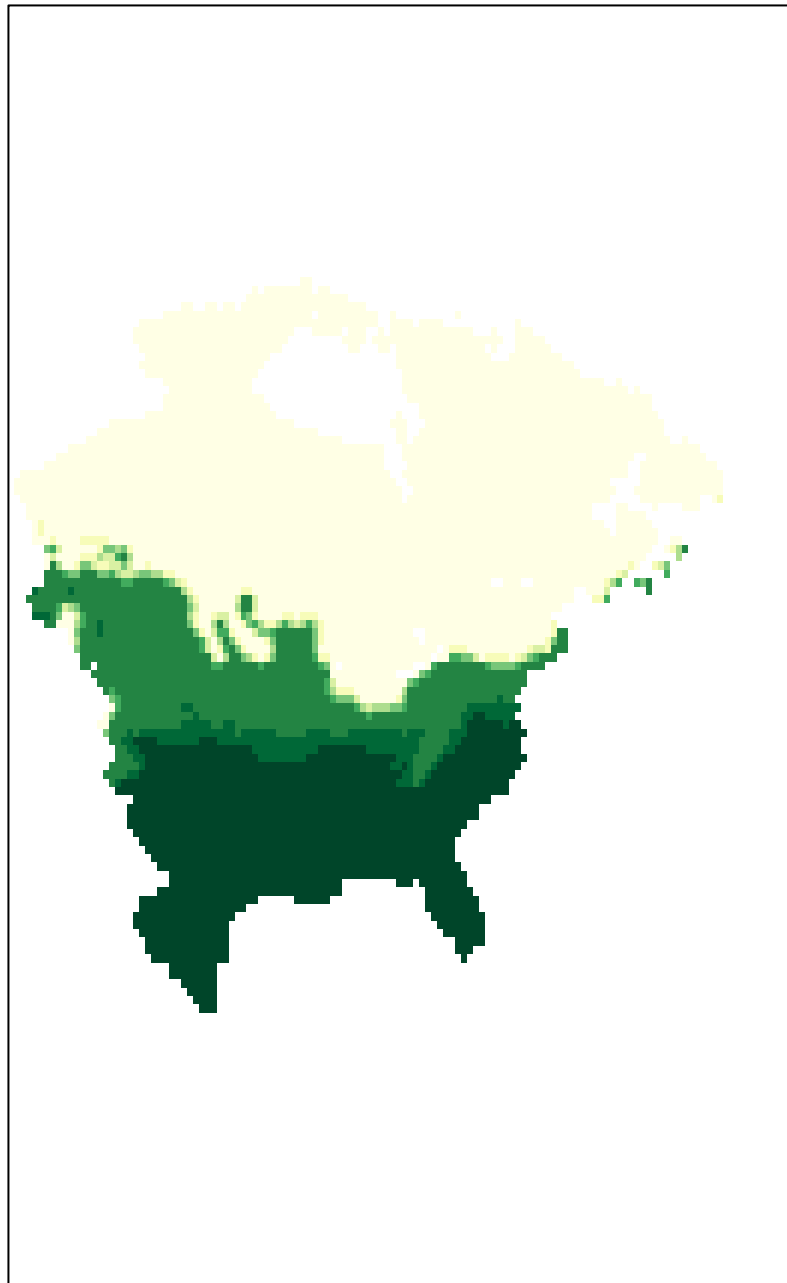


MAX, X5000.ybp

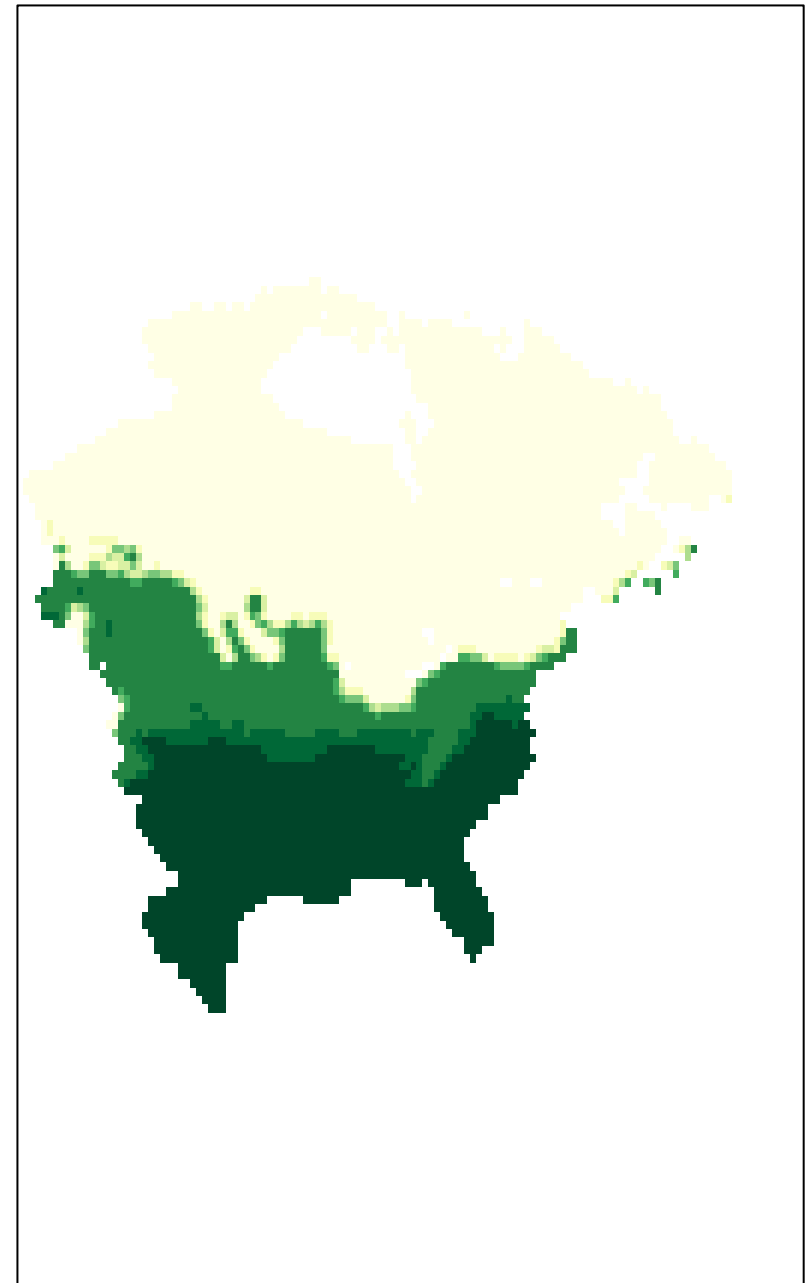


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X4000.ybp

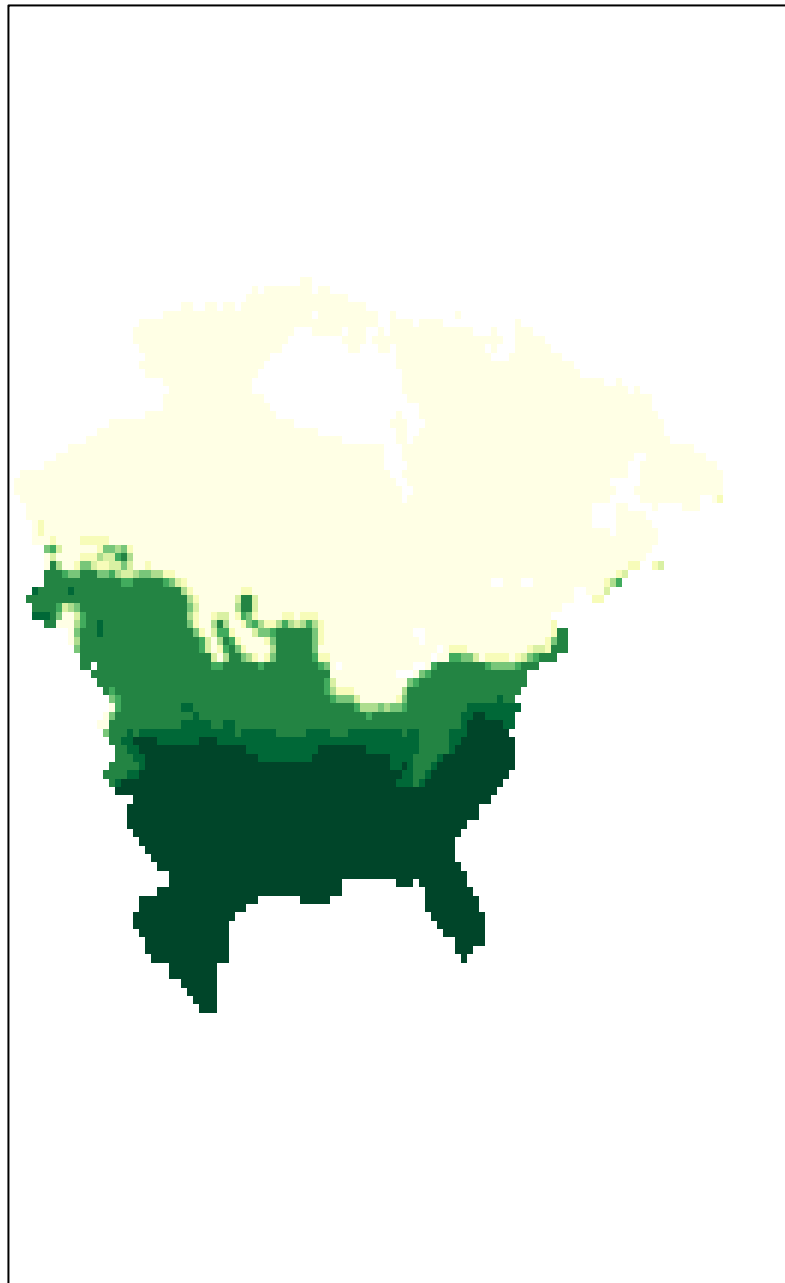


MAX, X4000.ybp

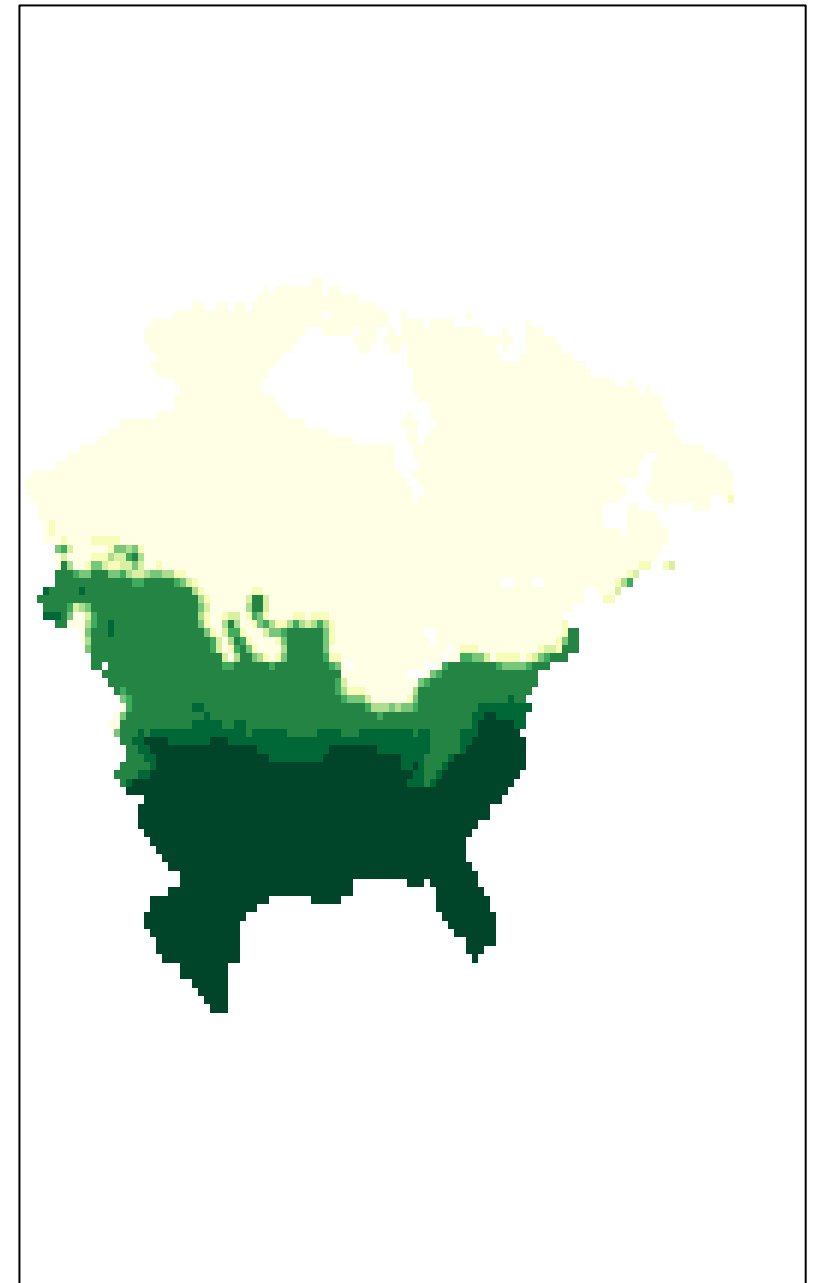


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X3000.ybp

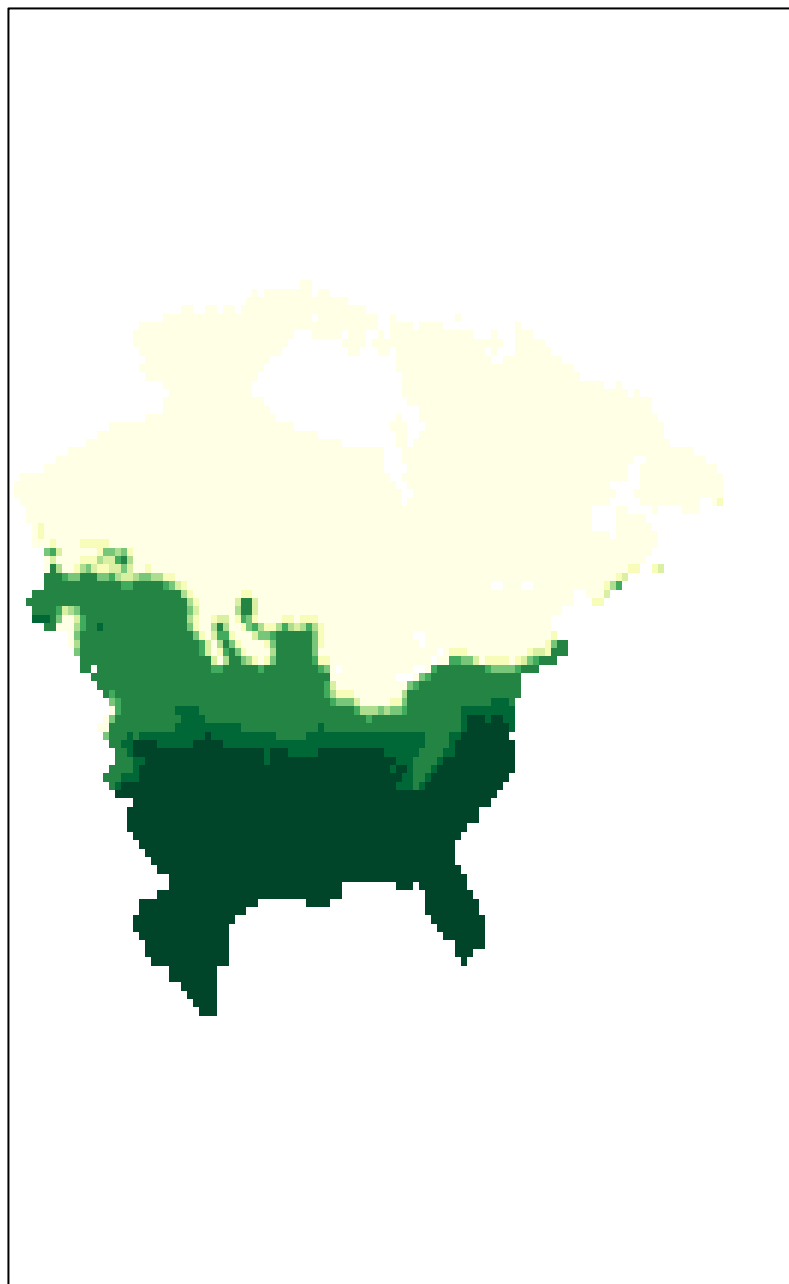


MAX, X3000.ybp

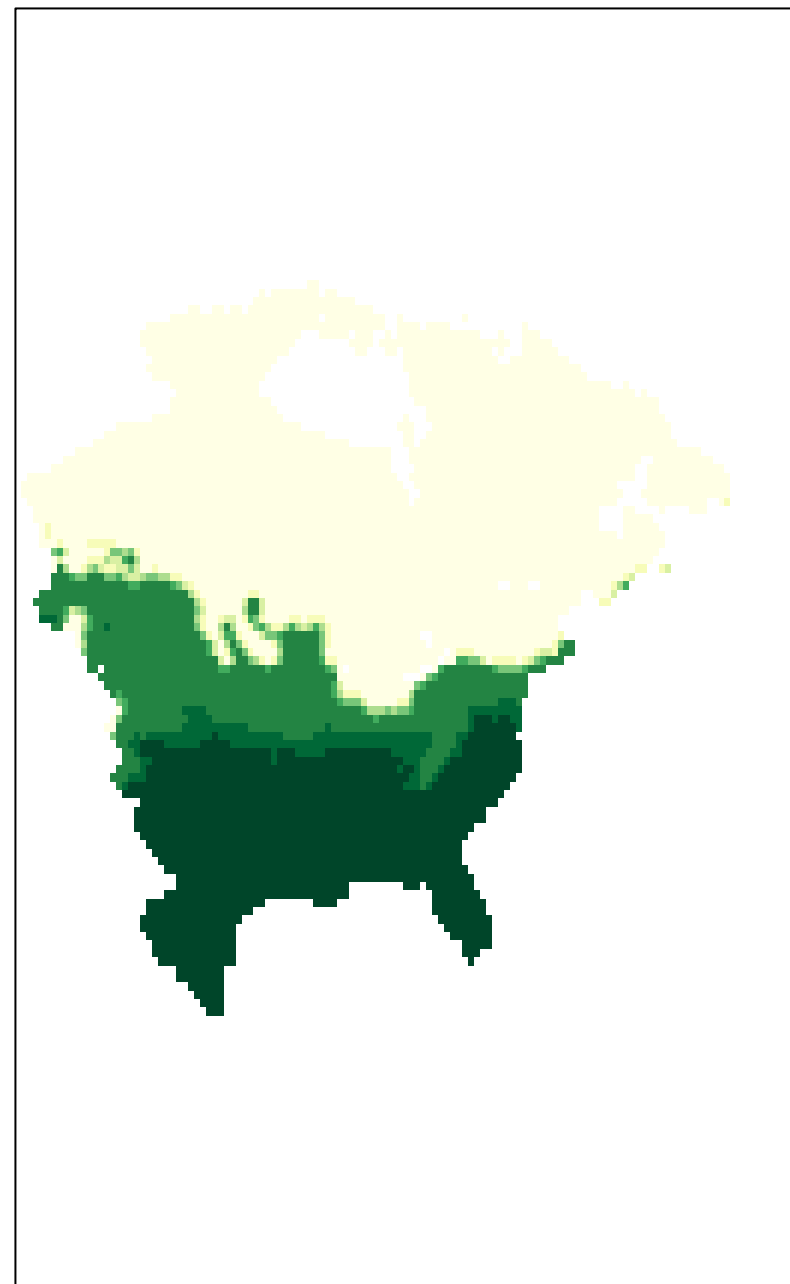


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X2000.ybp

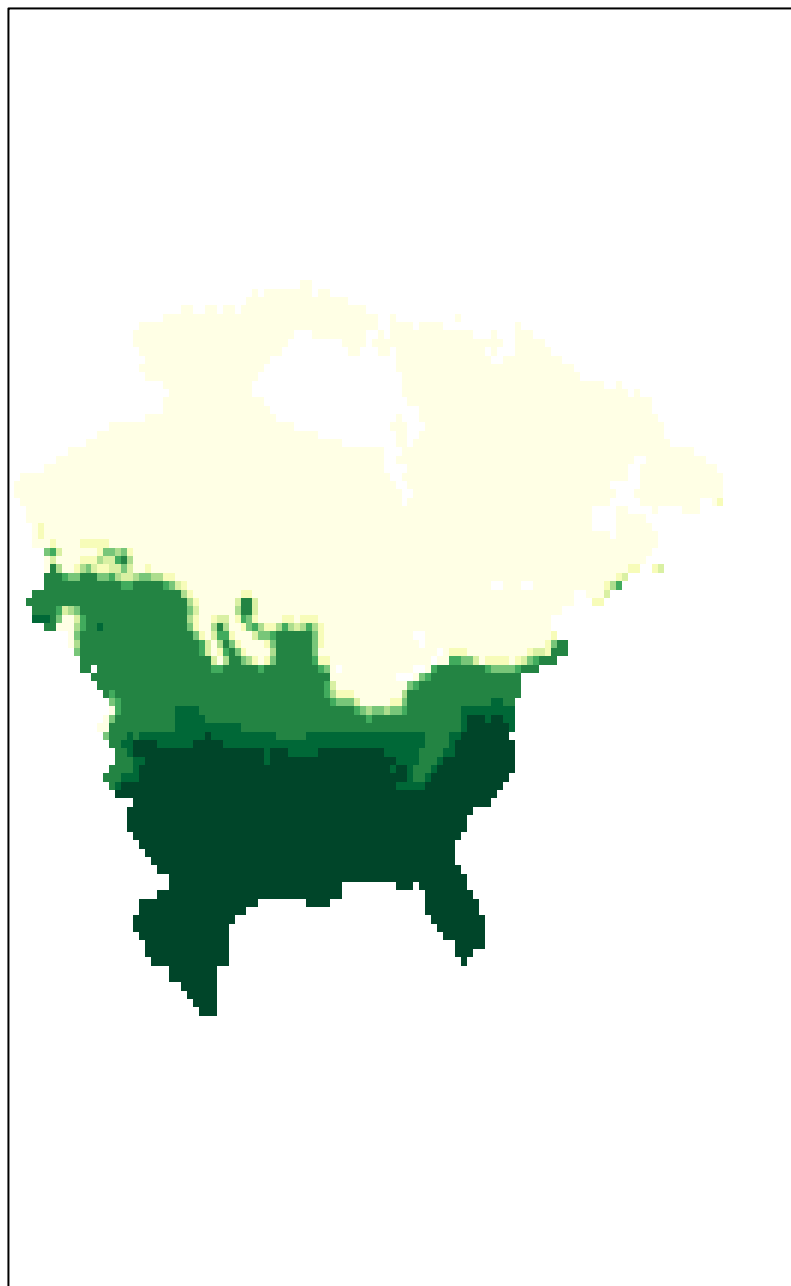


MAX, X2000.ybp

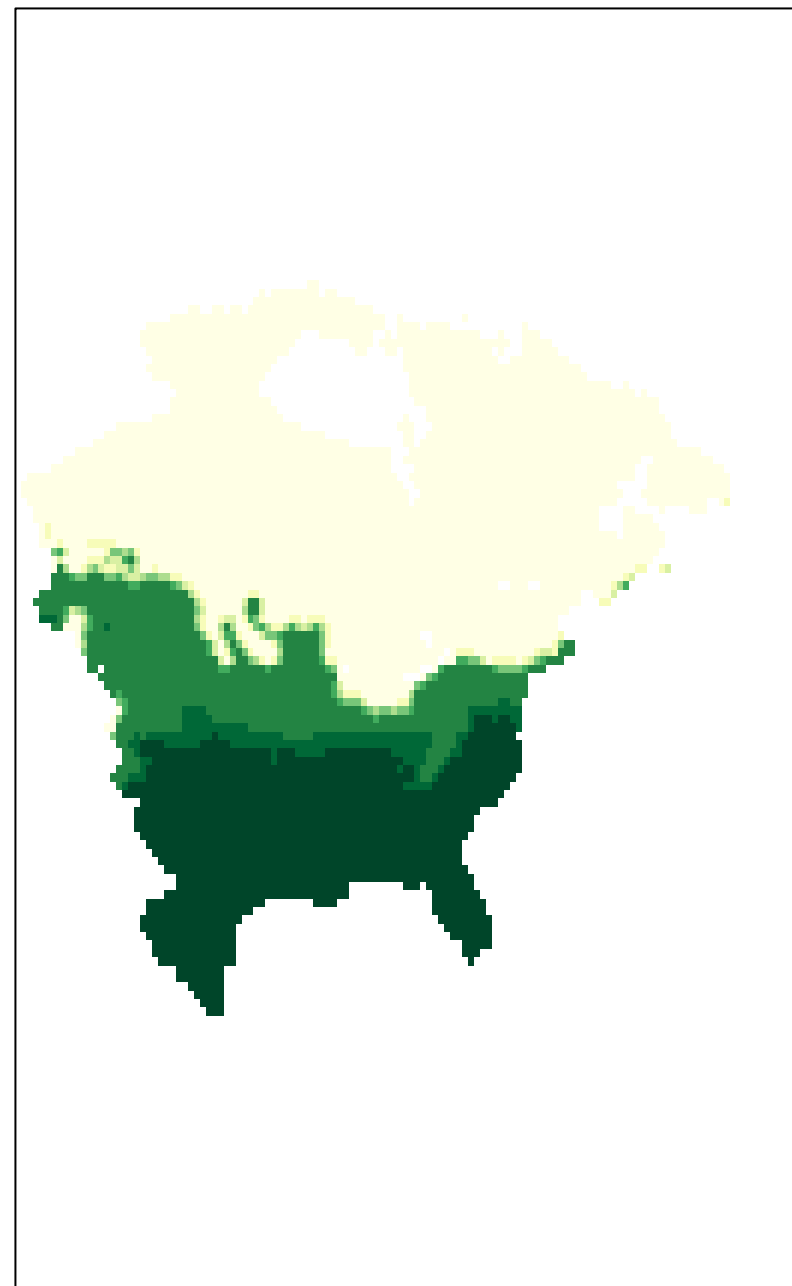


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X1000.ybp

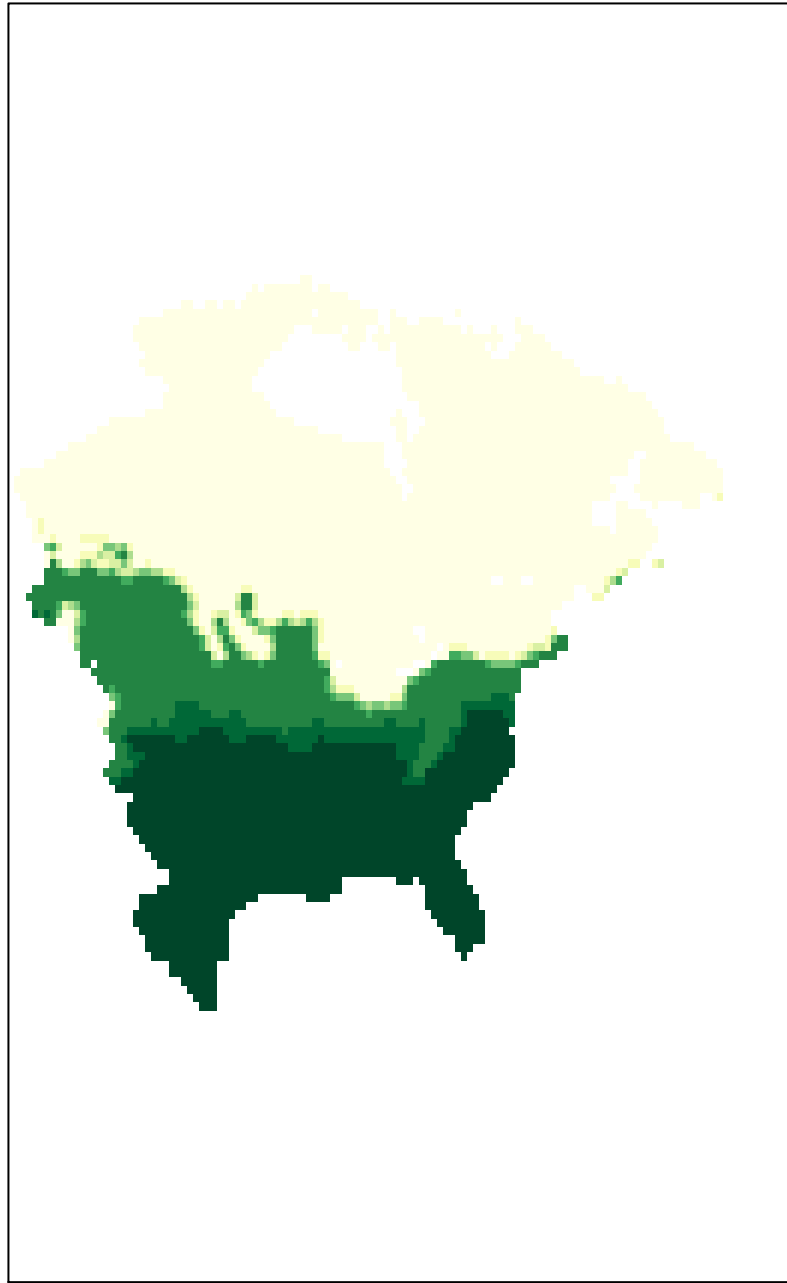


MAX, X1000.ybp

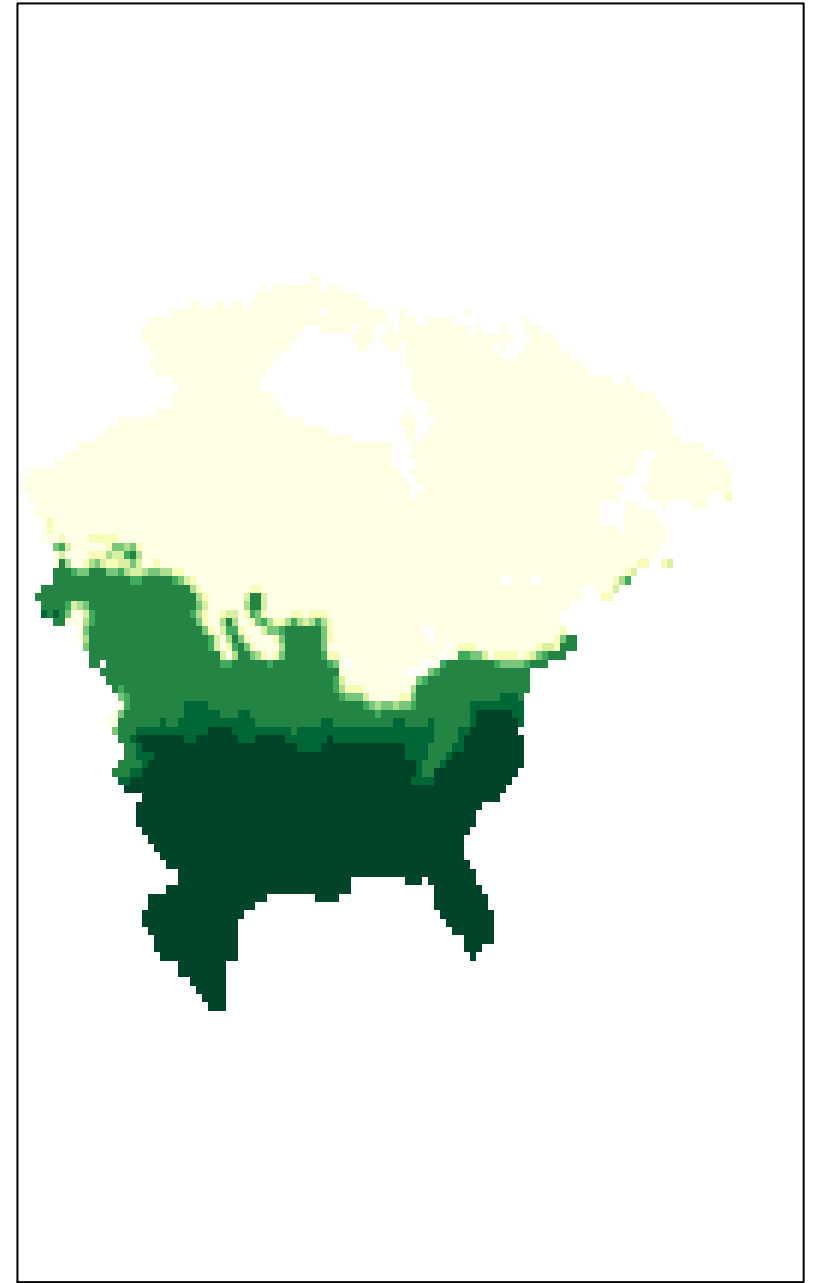


Species skipped = Fraxinus americana, GCM = ecbilt

MAX, X0.ybp



MAX, X0.ybp



Species skipped = *Fraxinus americana*, GCM = ecbilt