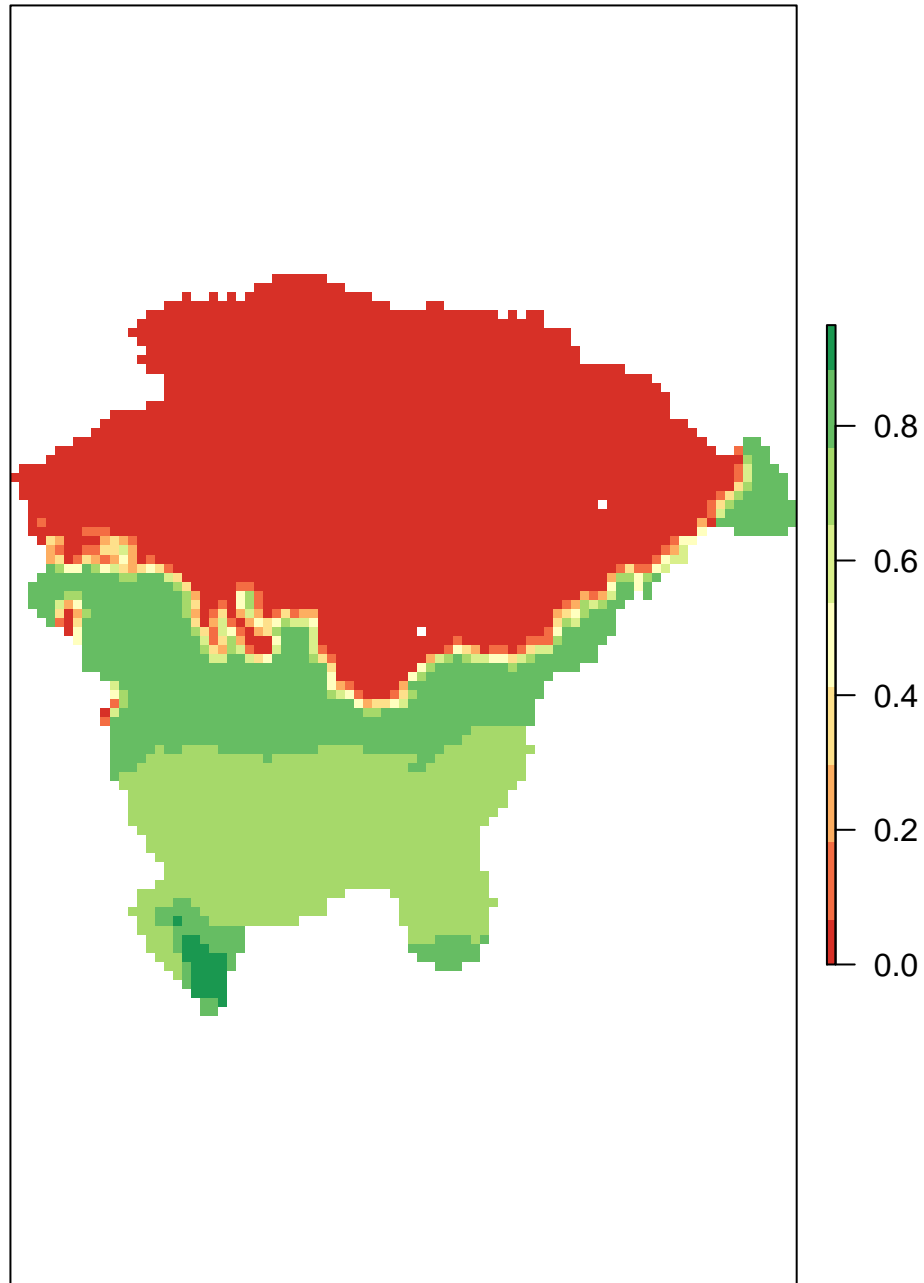
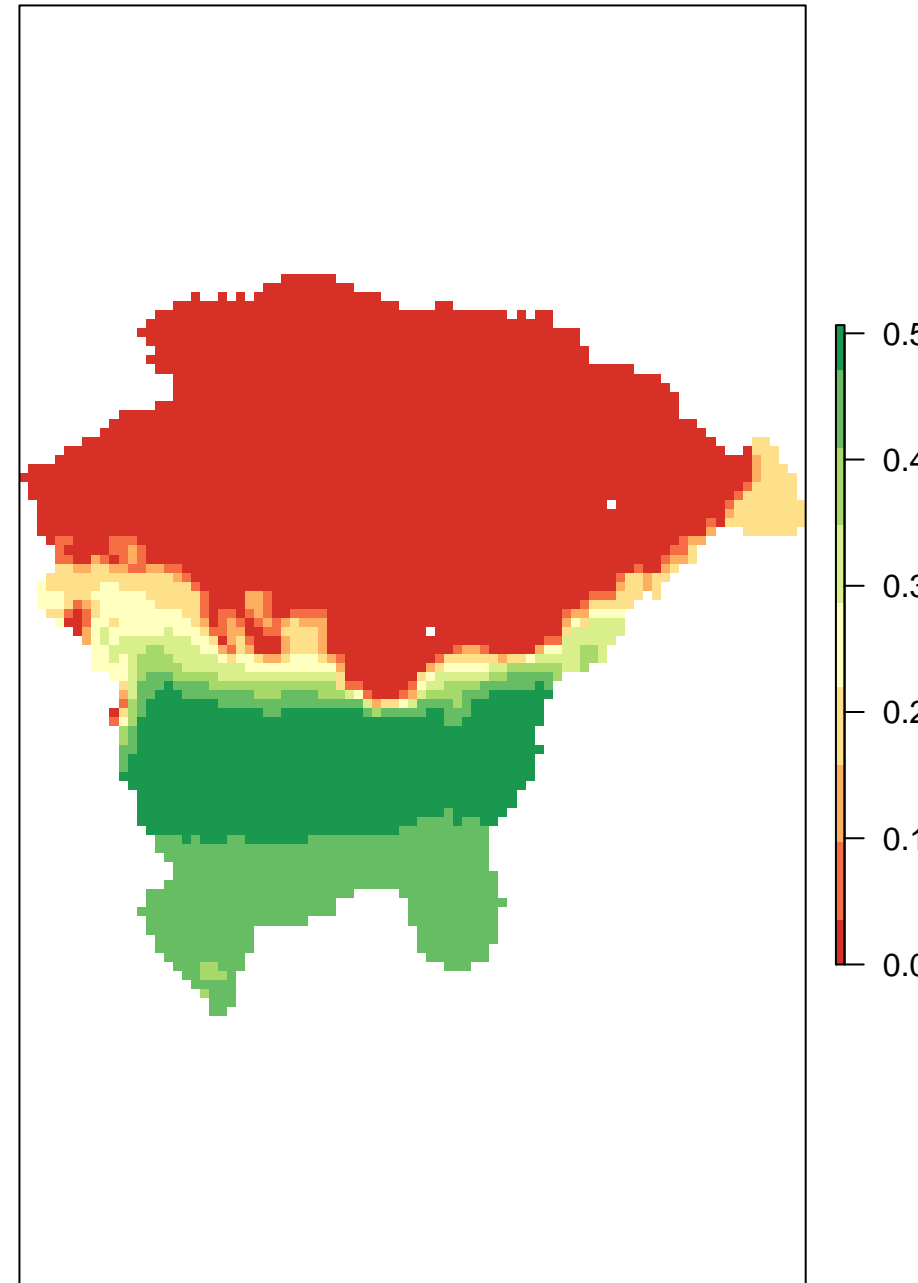


MAX, X21000.ybp

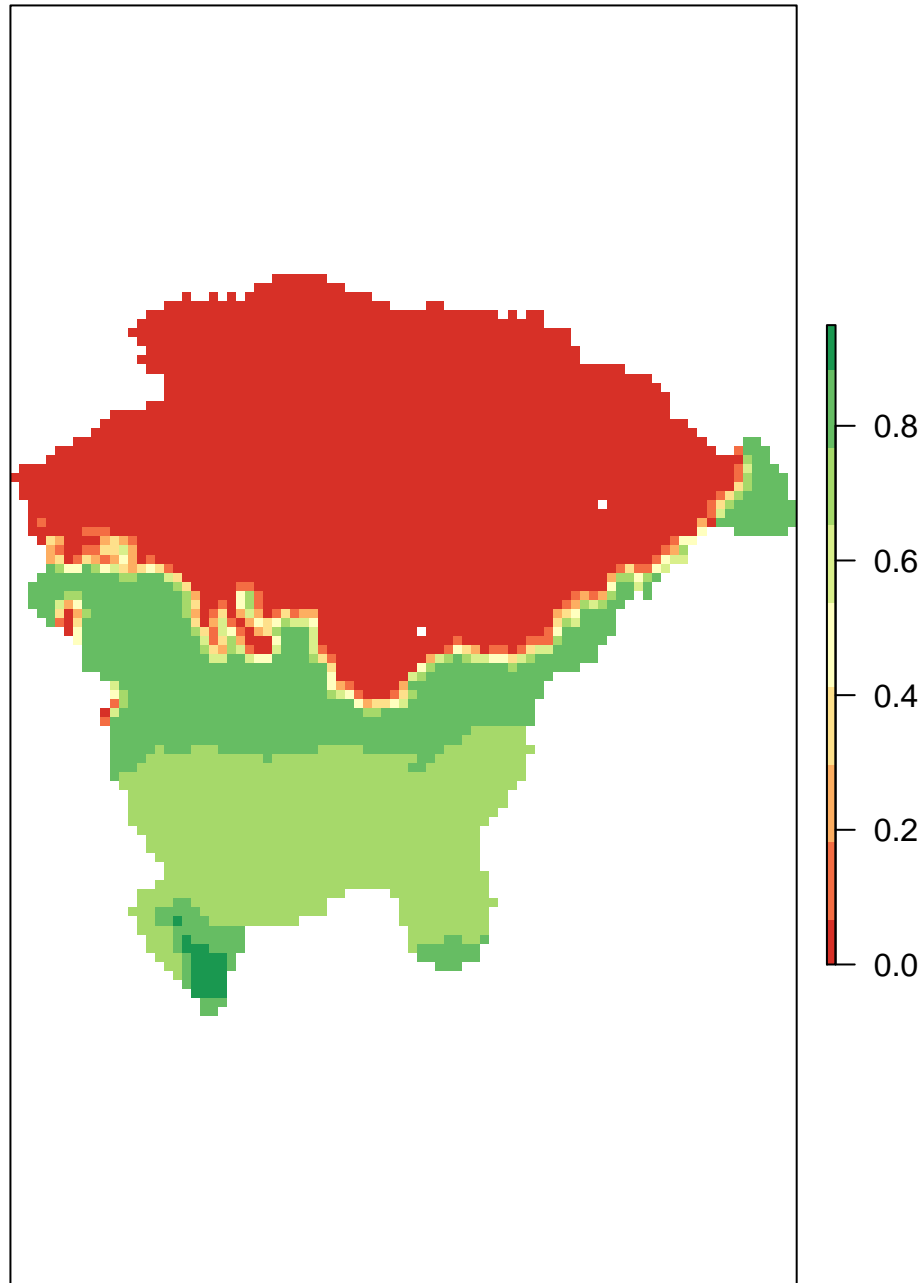


MAX without sp removed, X21000.ybp

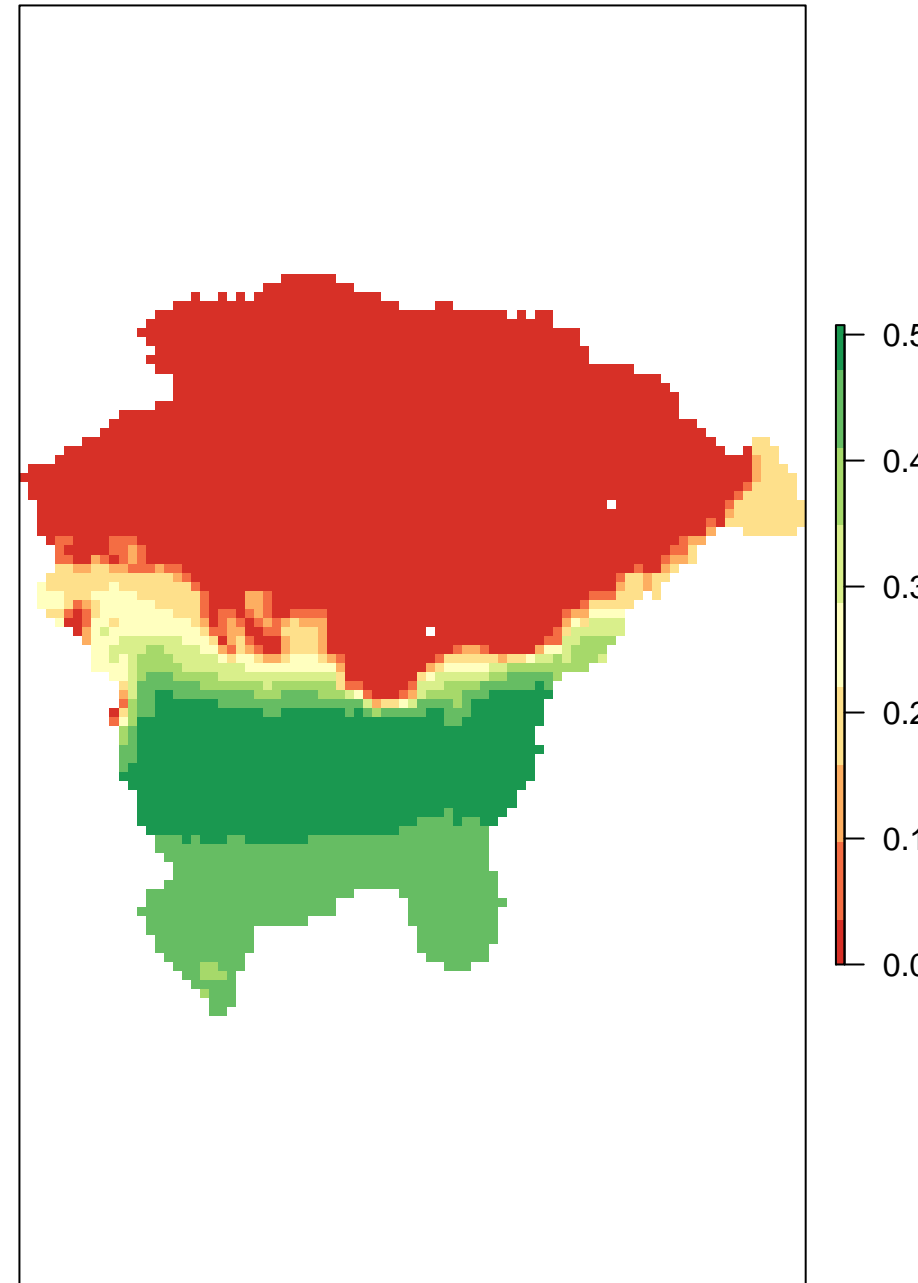


Species skipped = Fraxinus americana, GCM = ecbilt

MAX, X20000.ybp

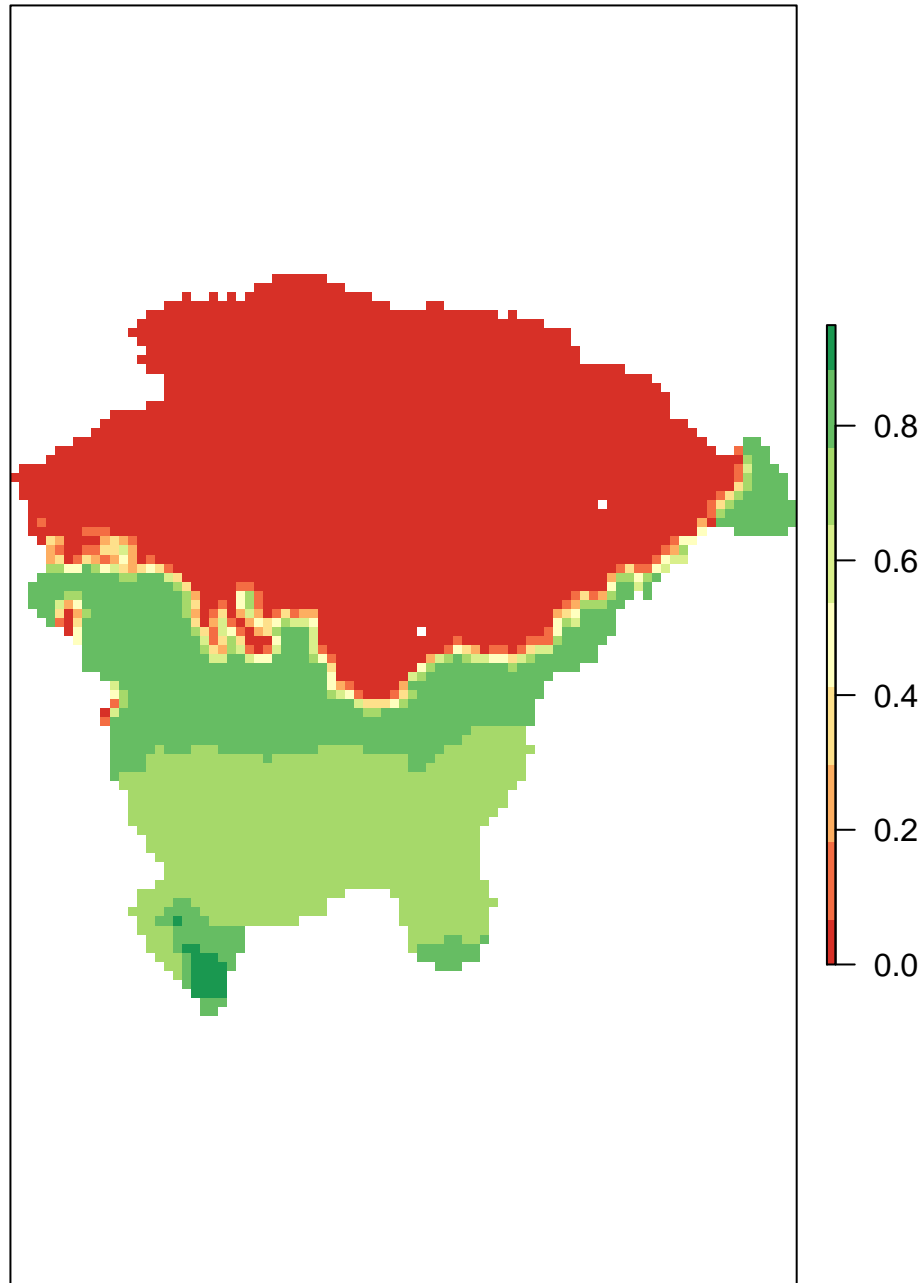


MAX without sp removed, X20000.ybp

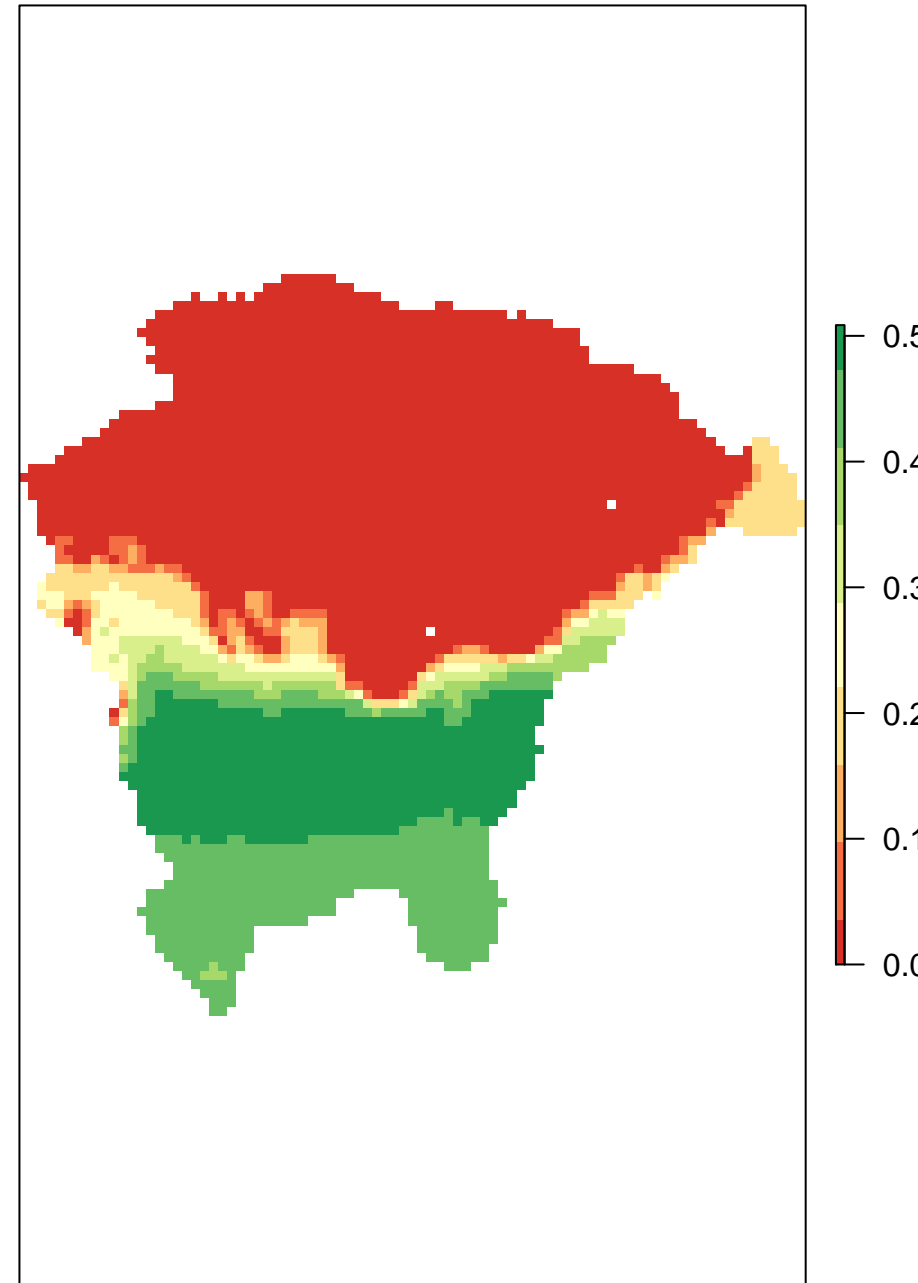


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X19000.ybp

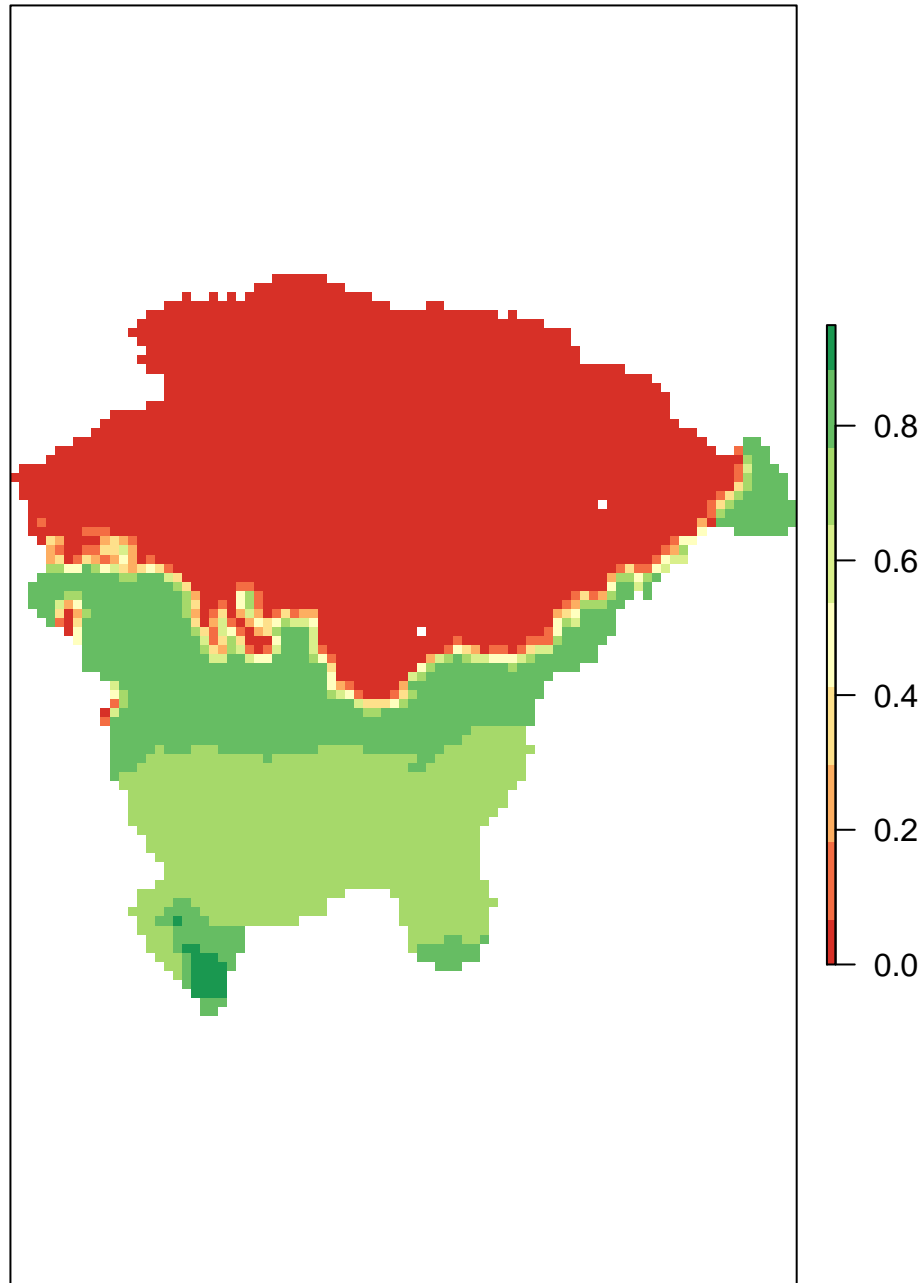


MAX without sp removed, X19000.ybp

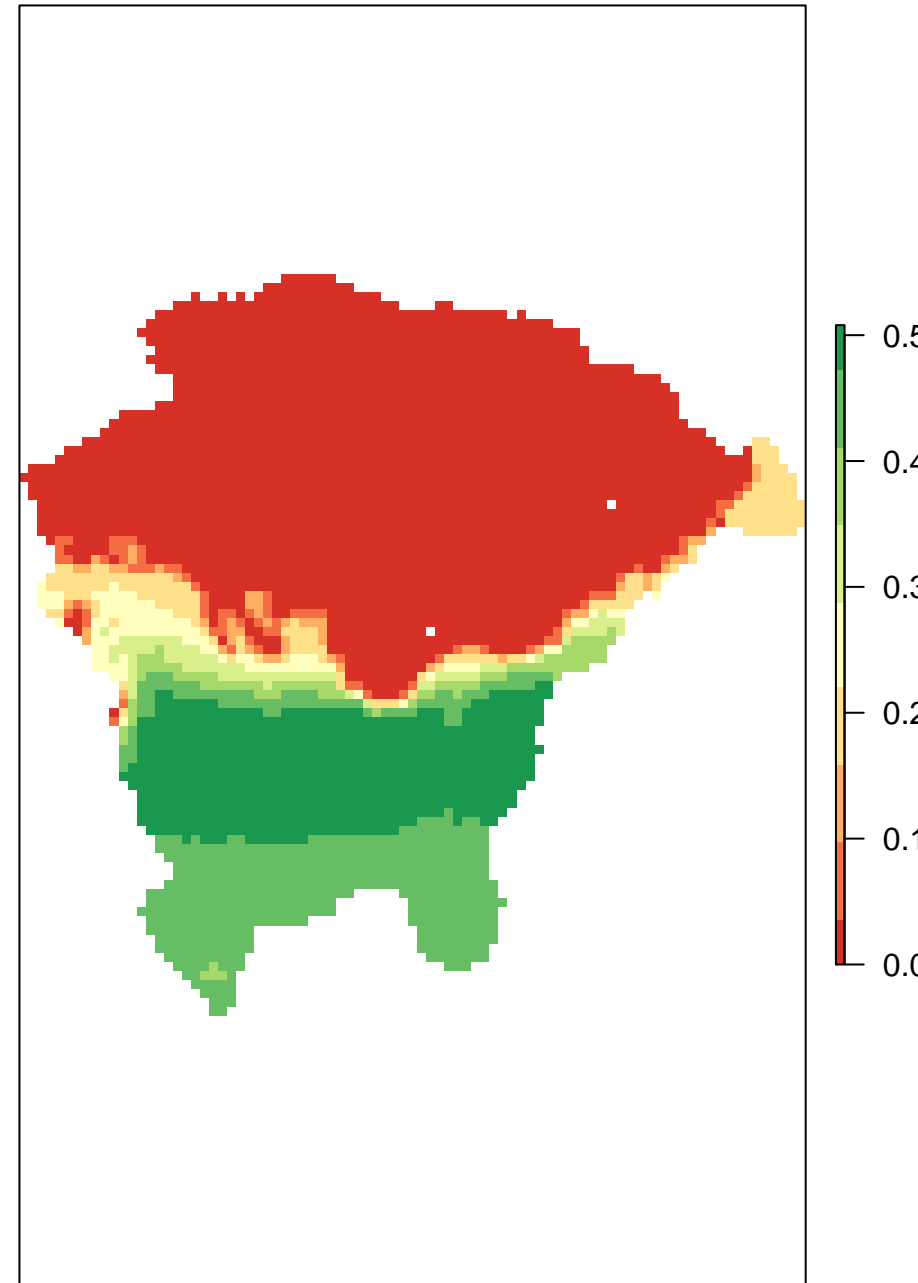


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X18000.ybp

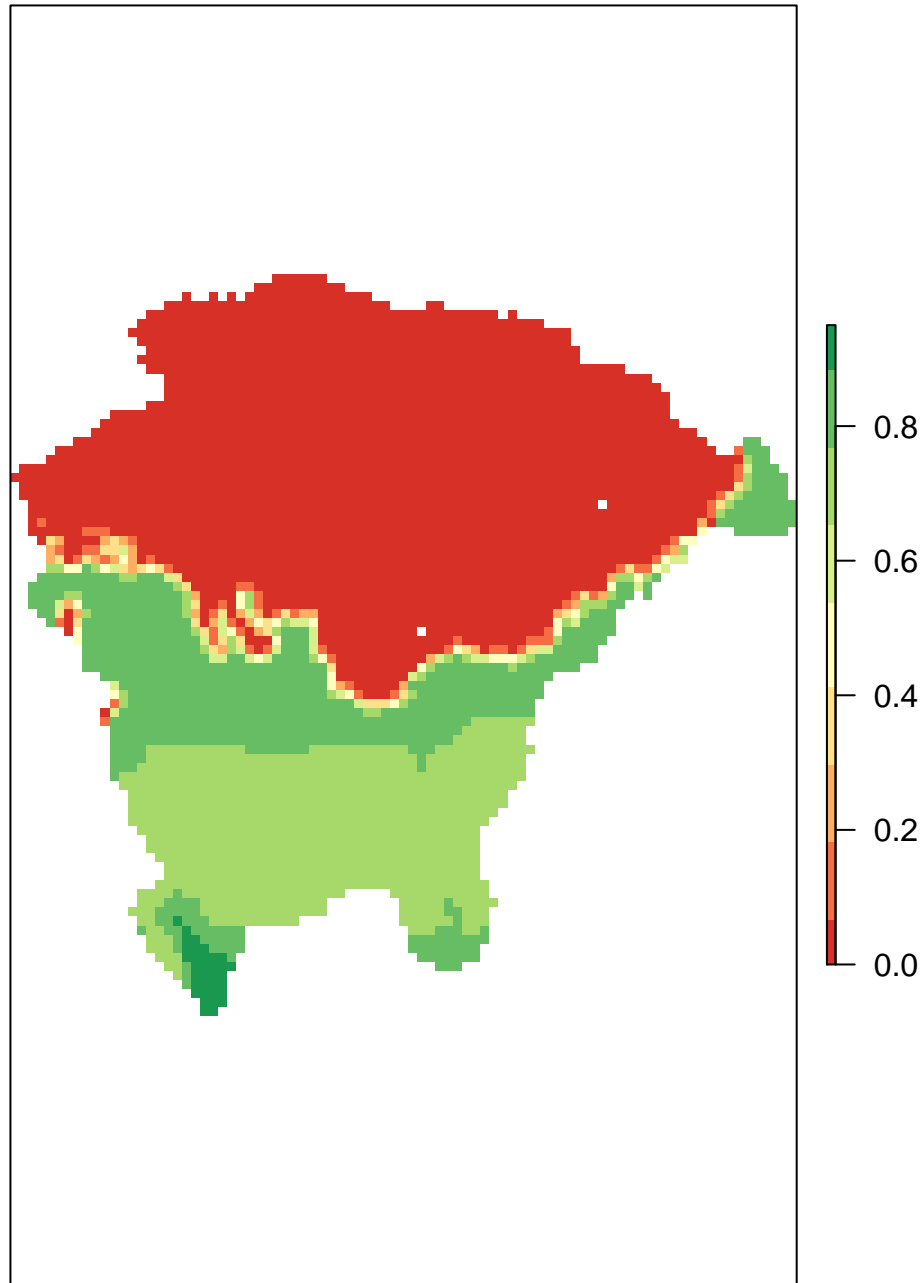


MAX without sp removed, X18000.ybp

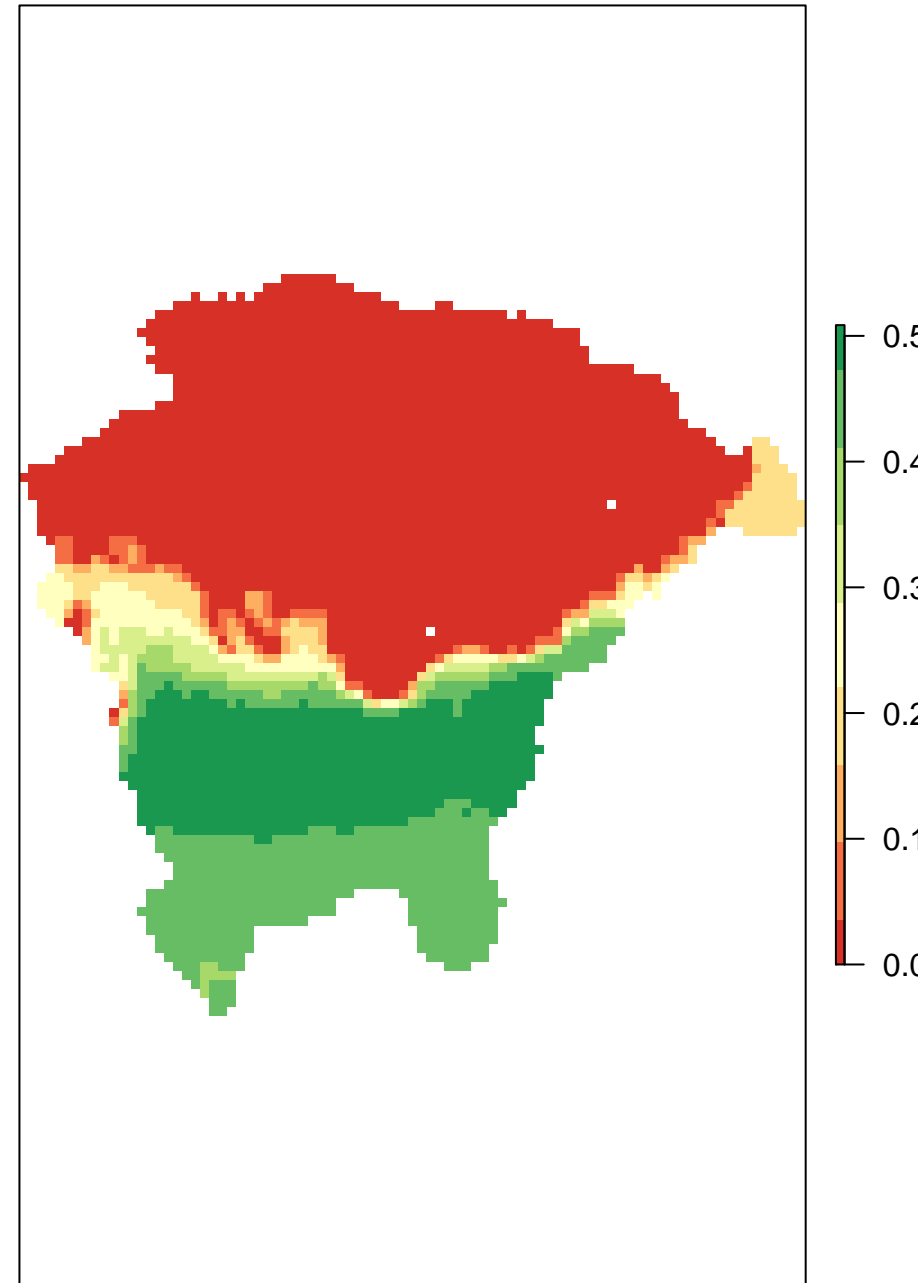


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X17000.ybp

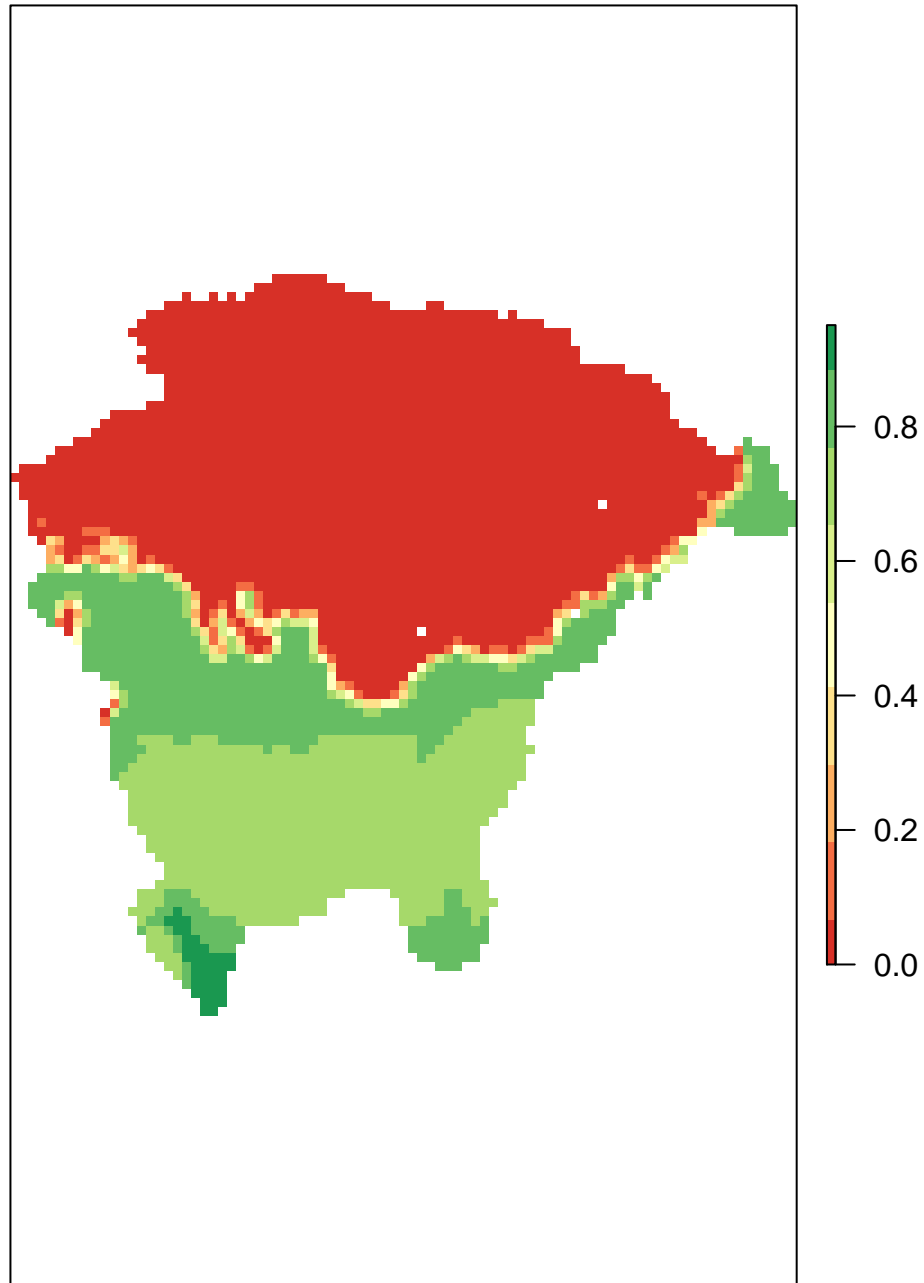


MAX without sp removed, X17000.ybp

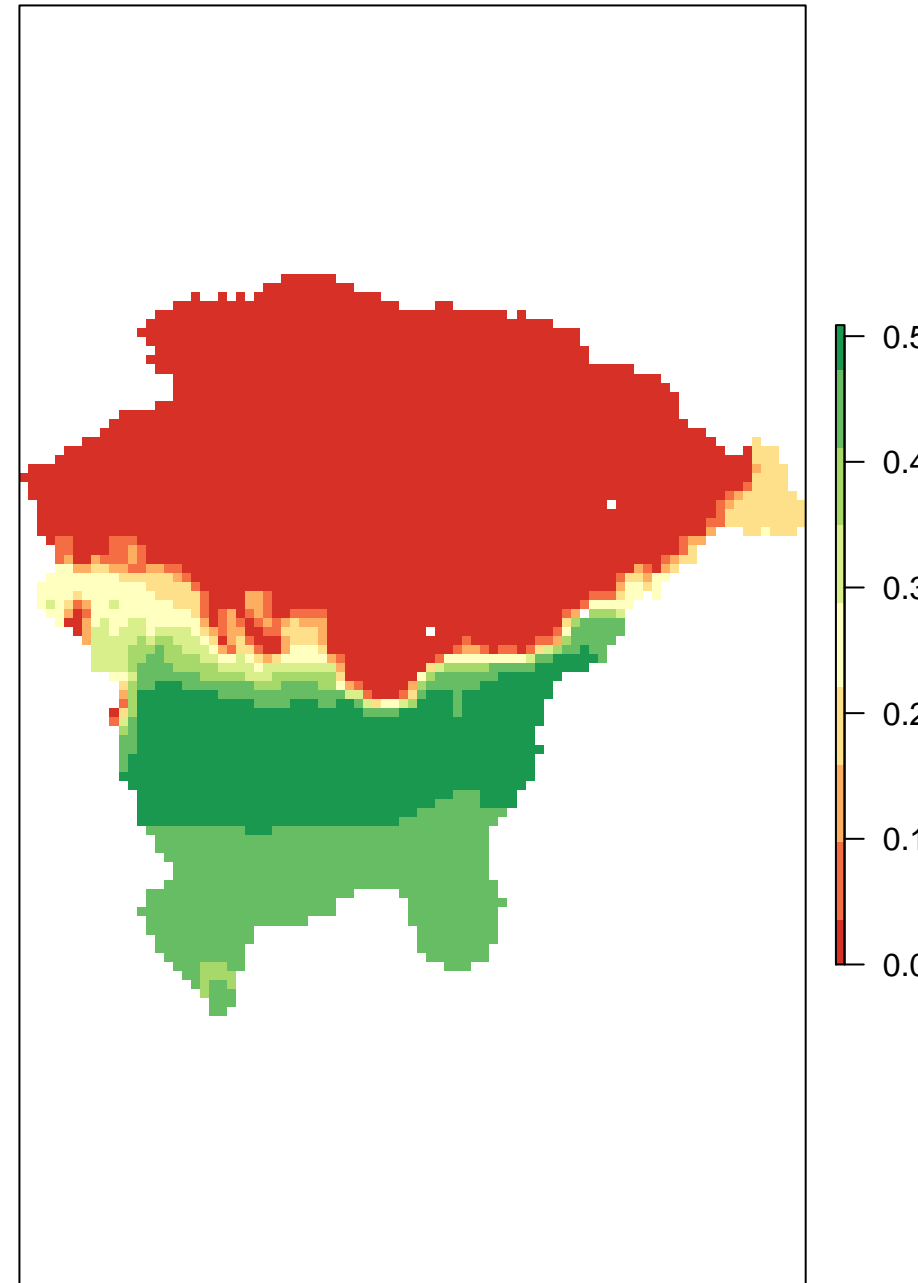


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X16000.ybp

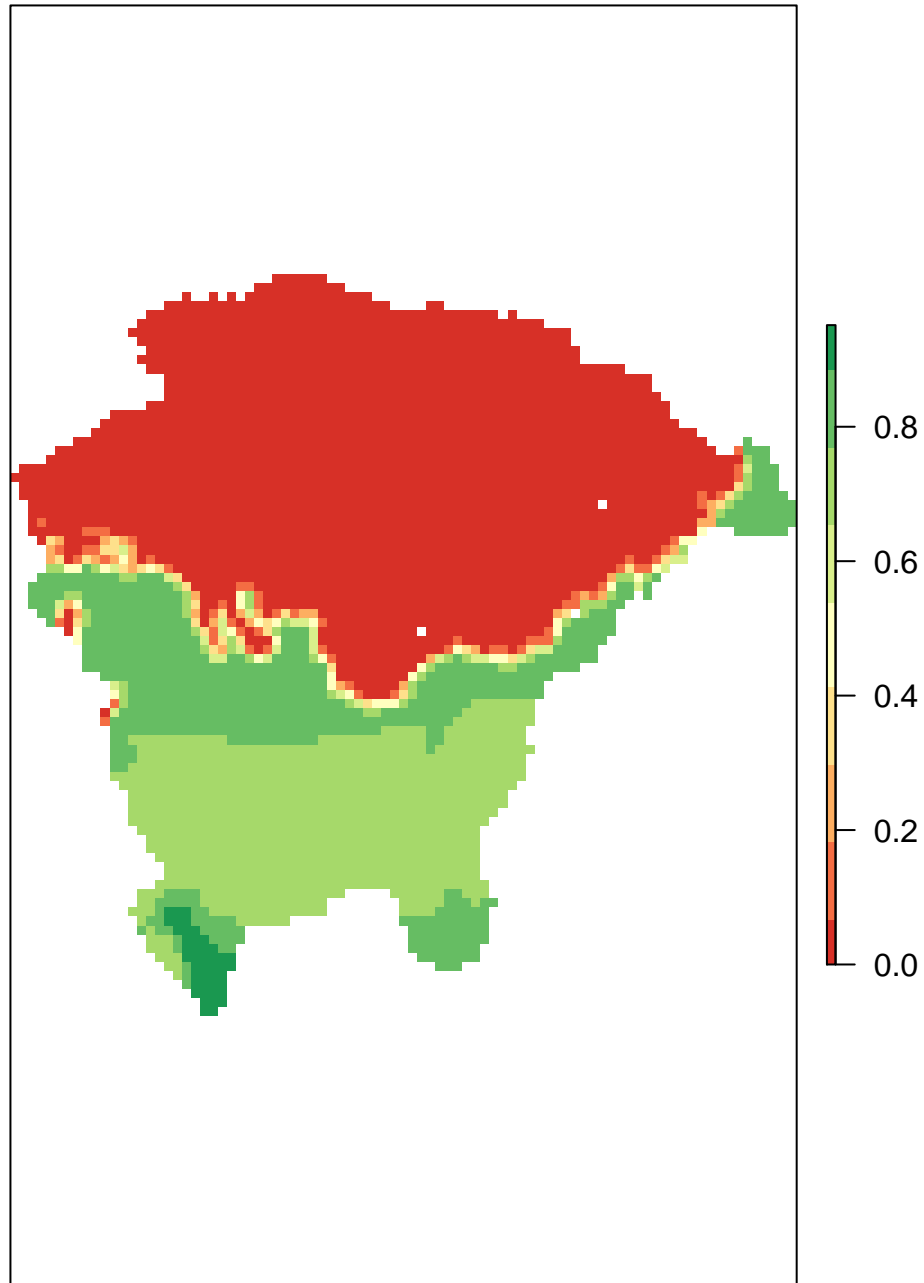


MAX without sp removed, X16000.ybp

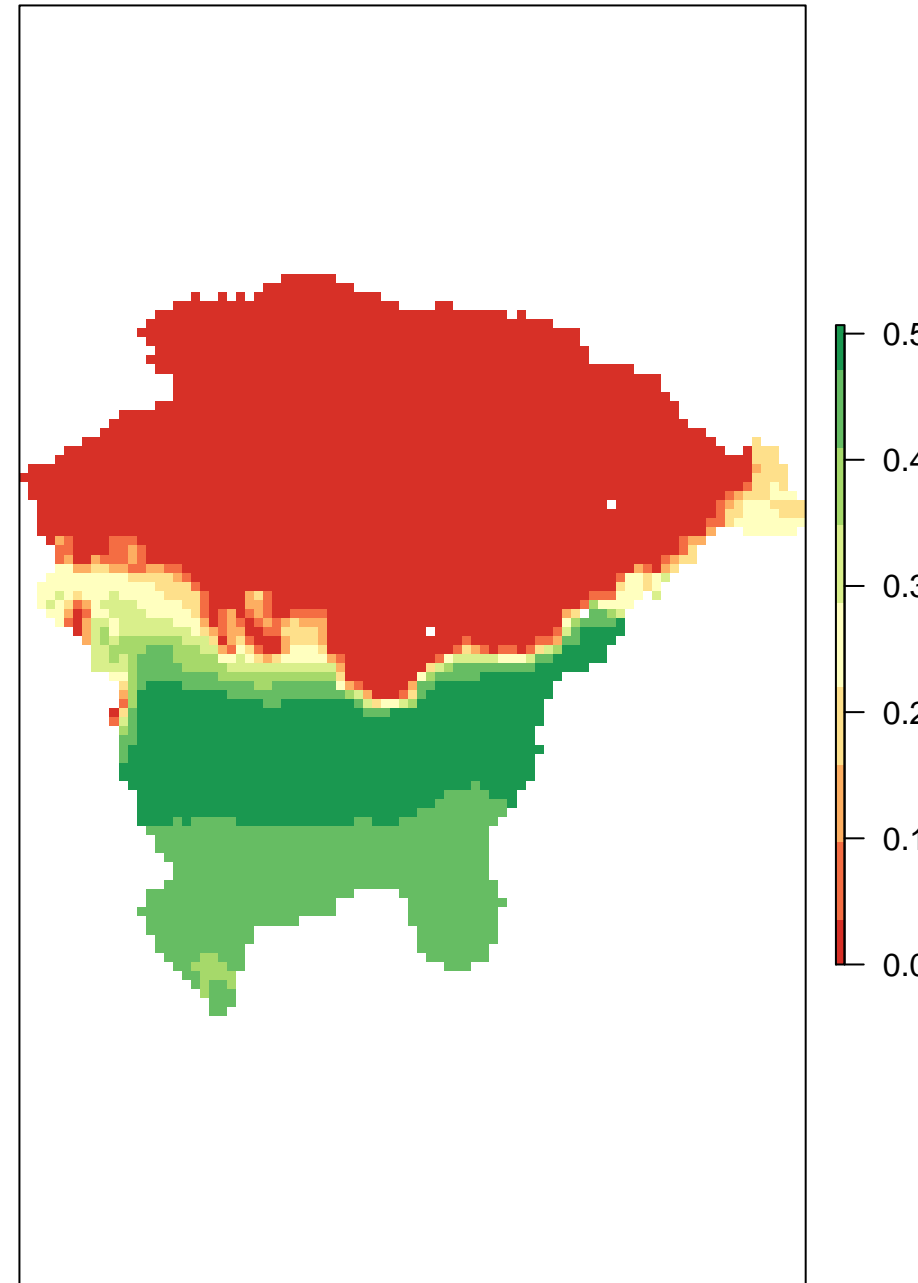


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X15000.ybp

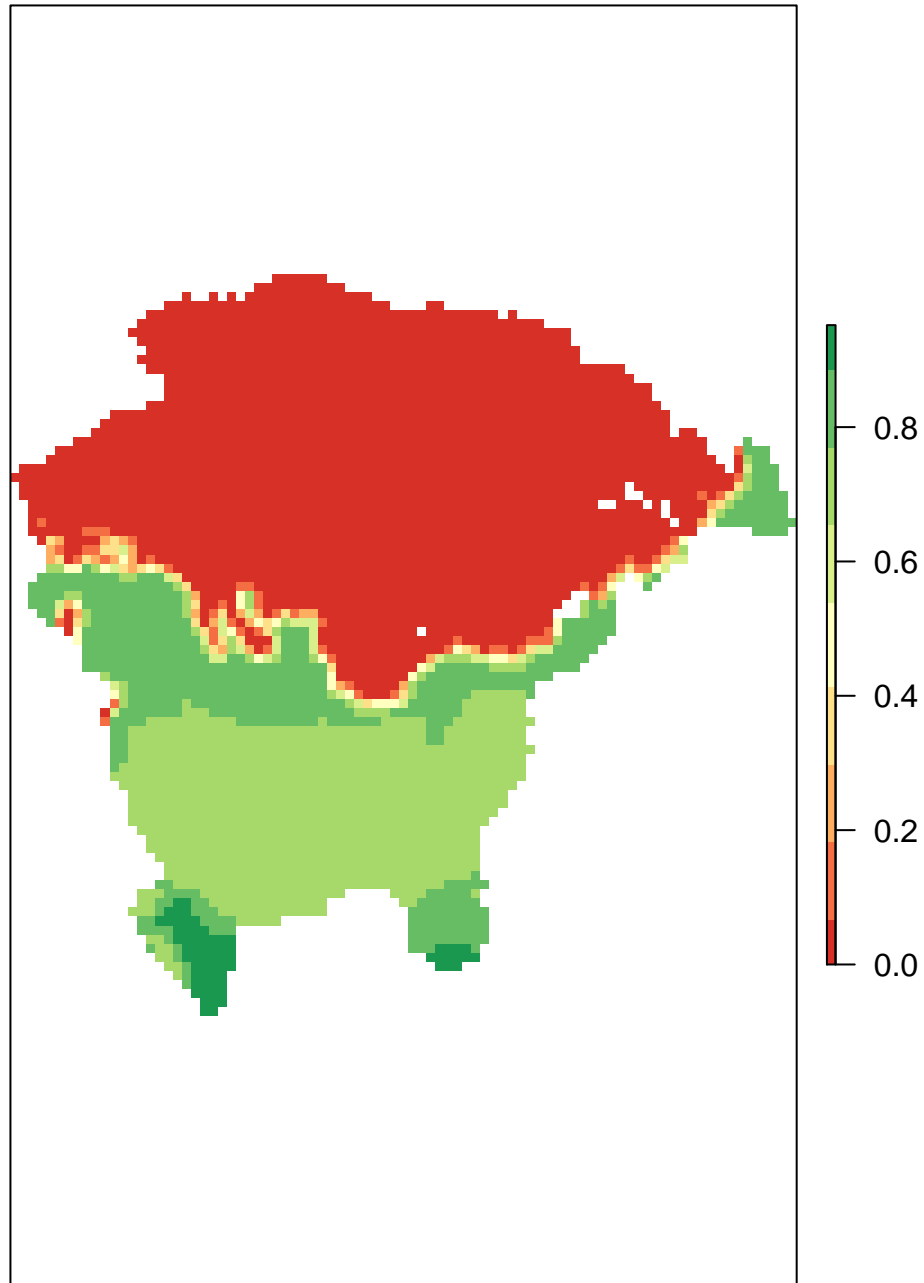


MAX without sp removed, X15000.ybp

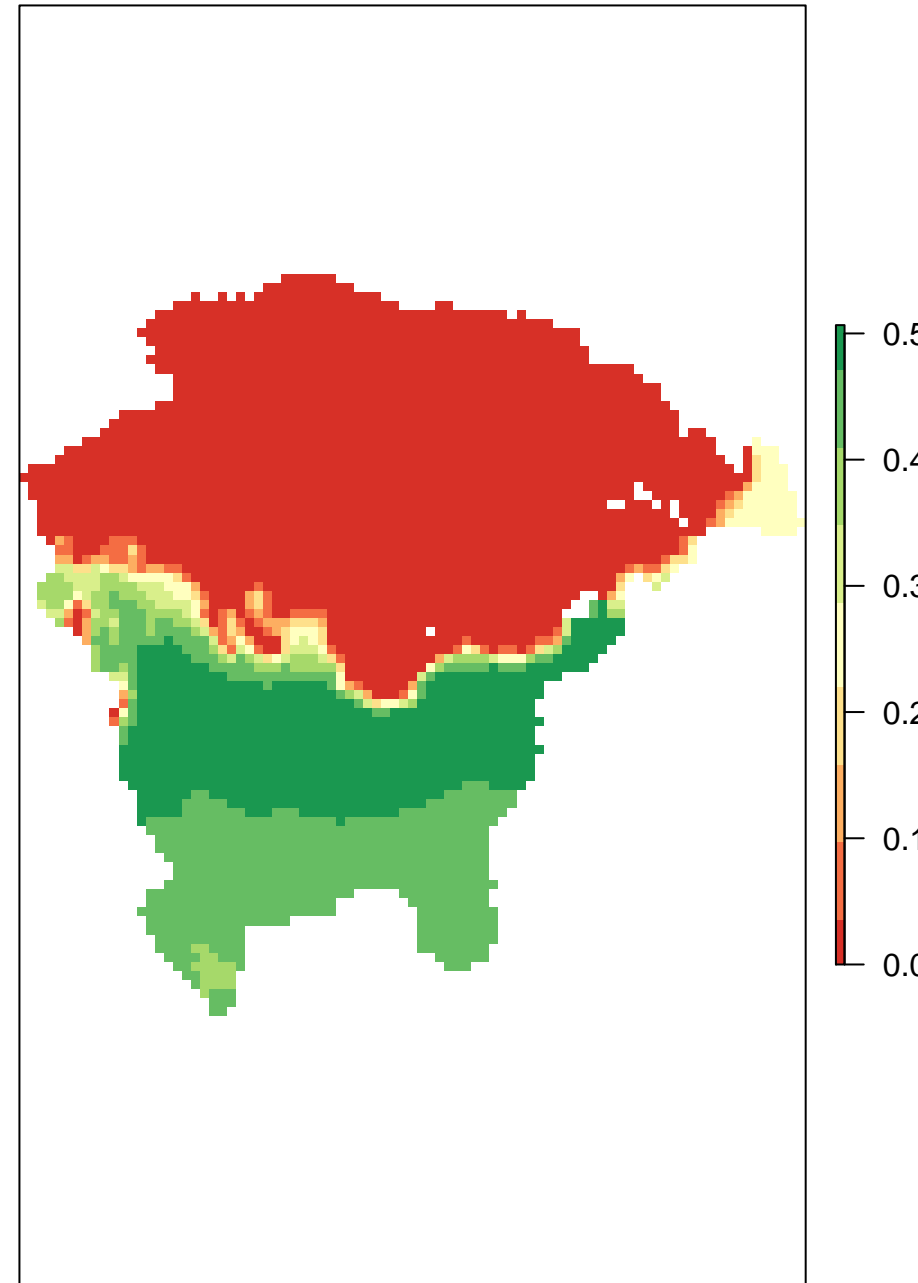


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X14000.ybp

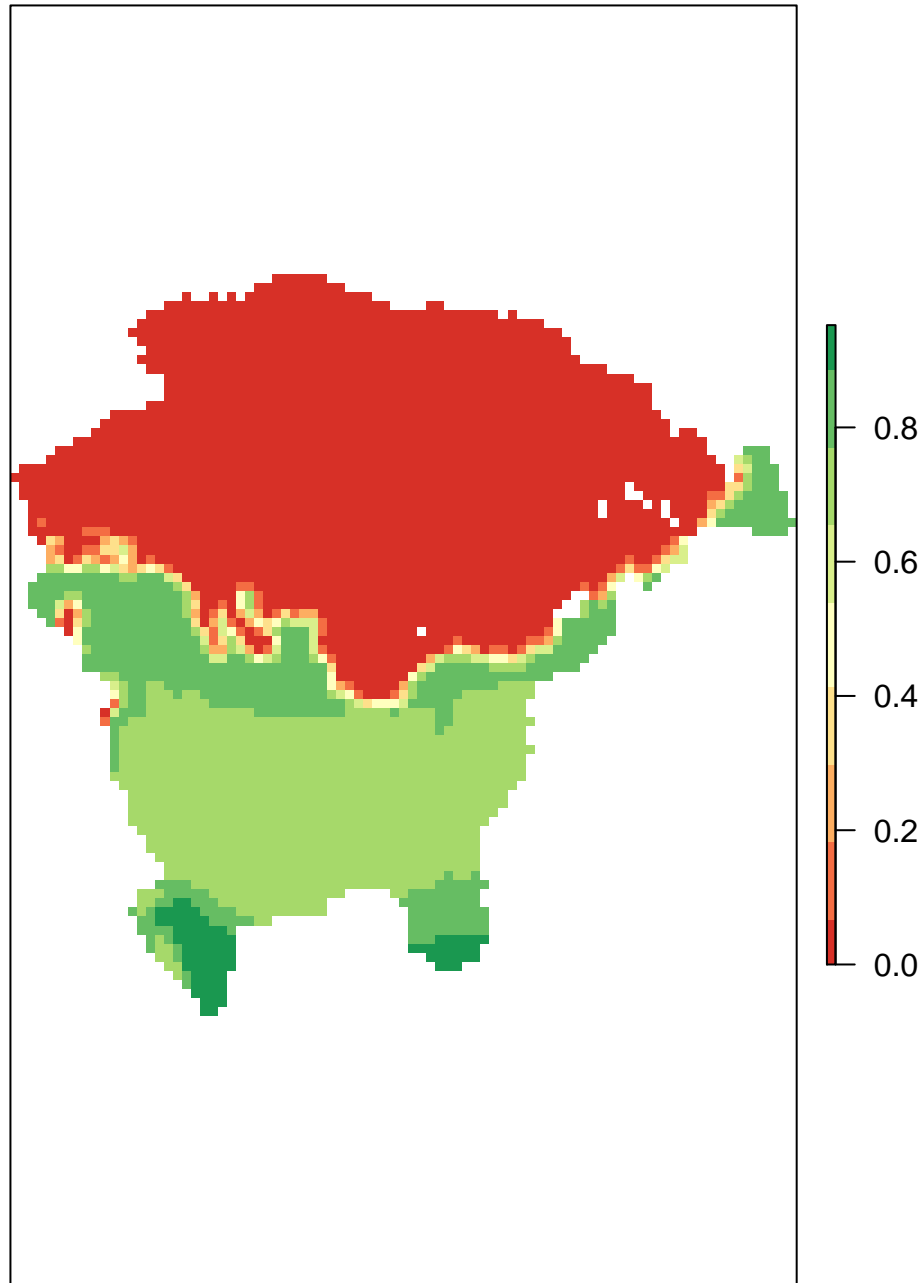


MAX without sp removed, X14000.ybp

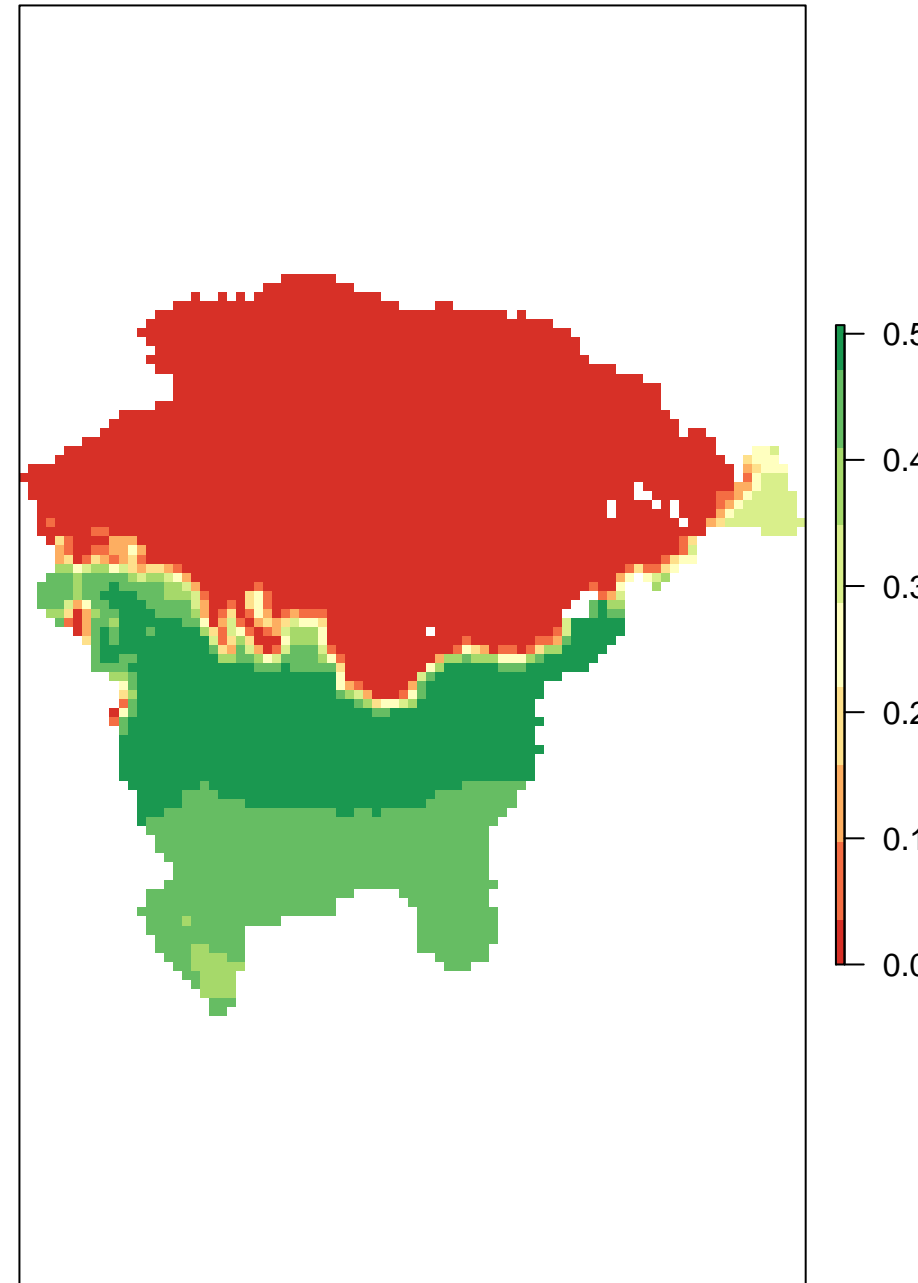


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X13000.ybp

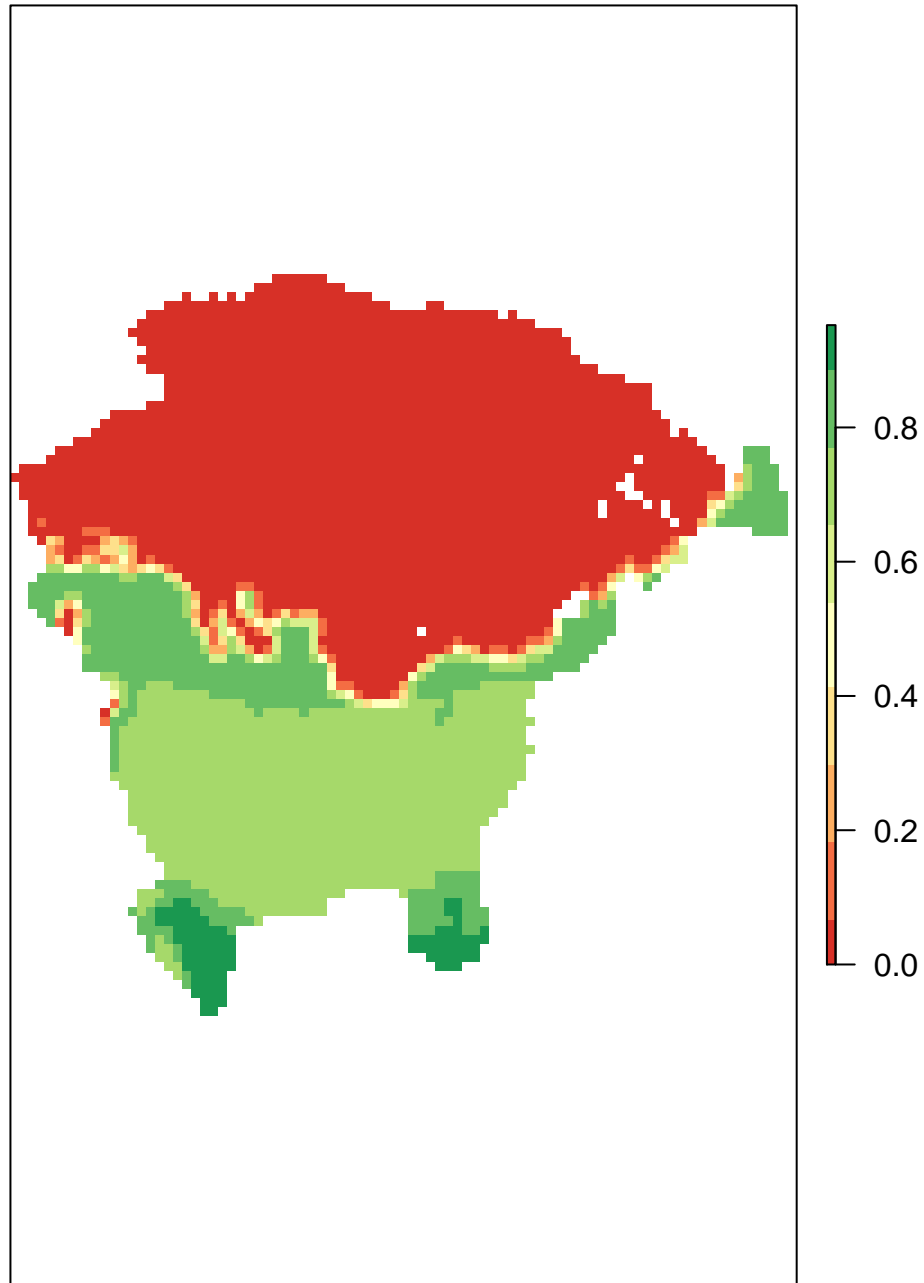


MAX without sp removed, X13000.ybp

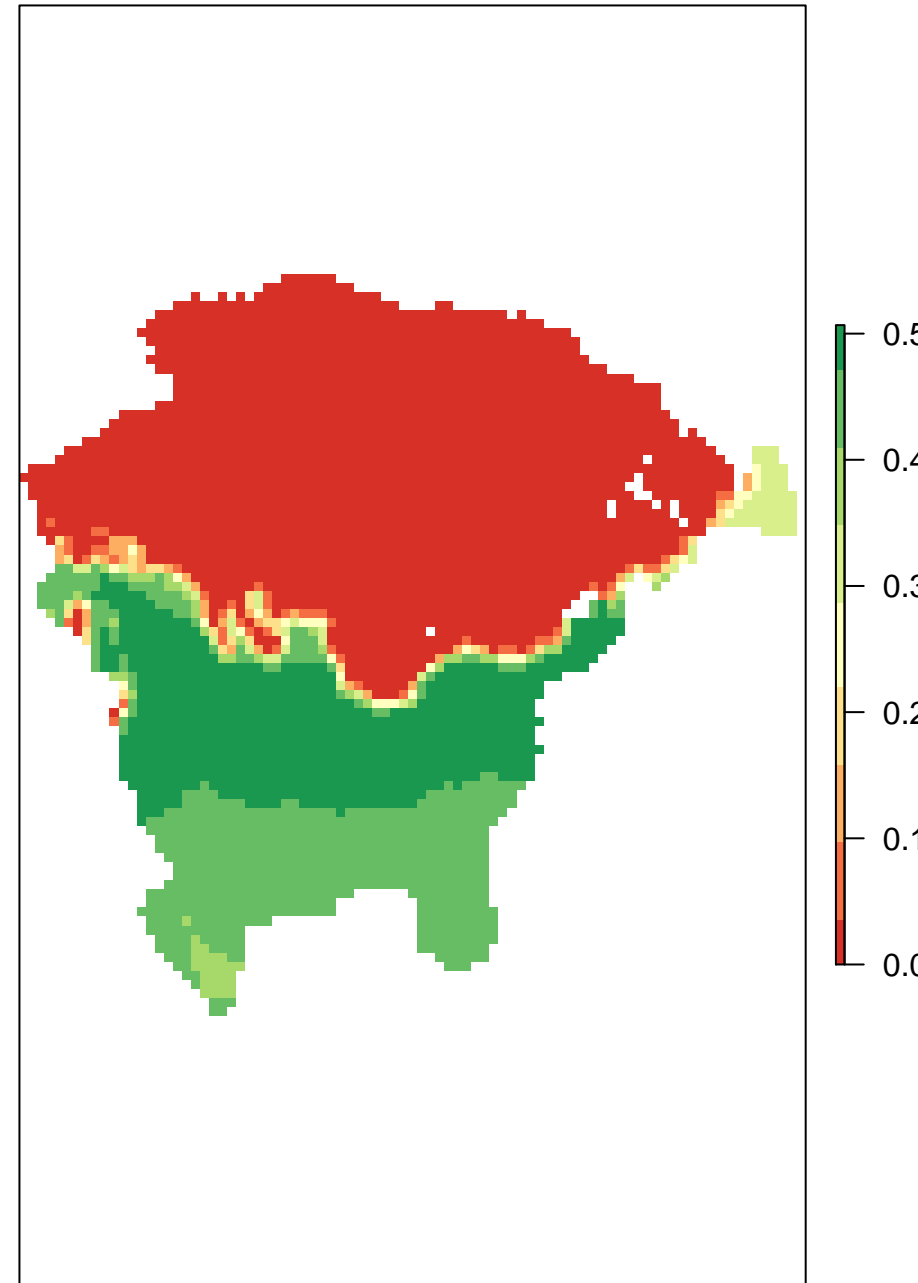


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X12000.ybp

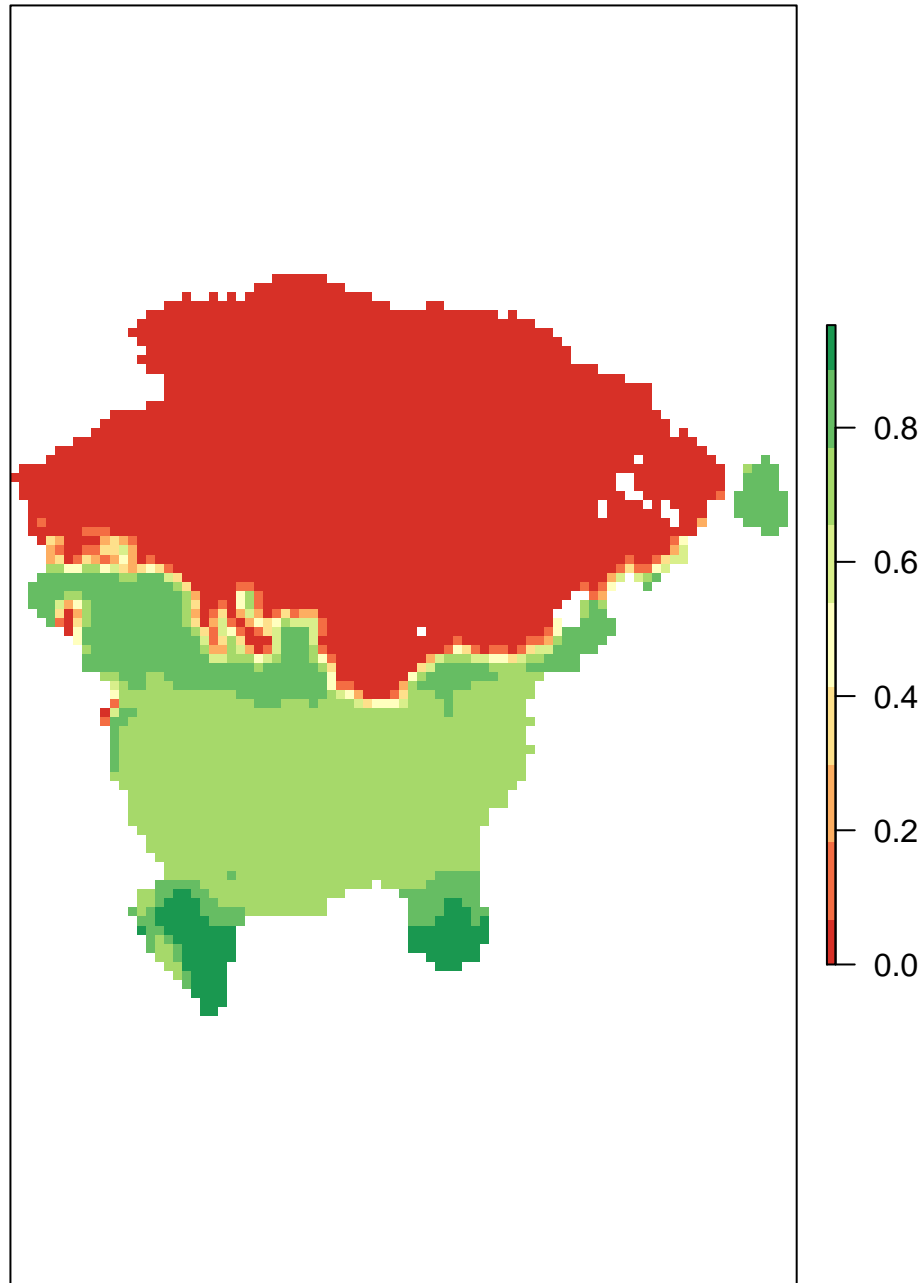


MAX without sp removed, X12000.ybp

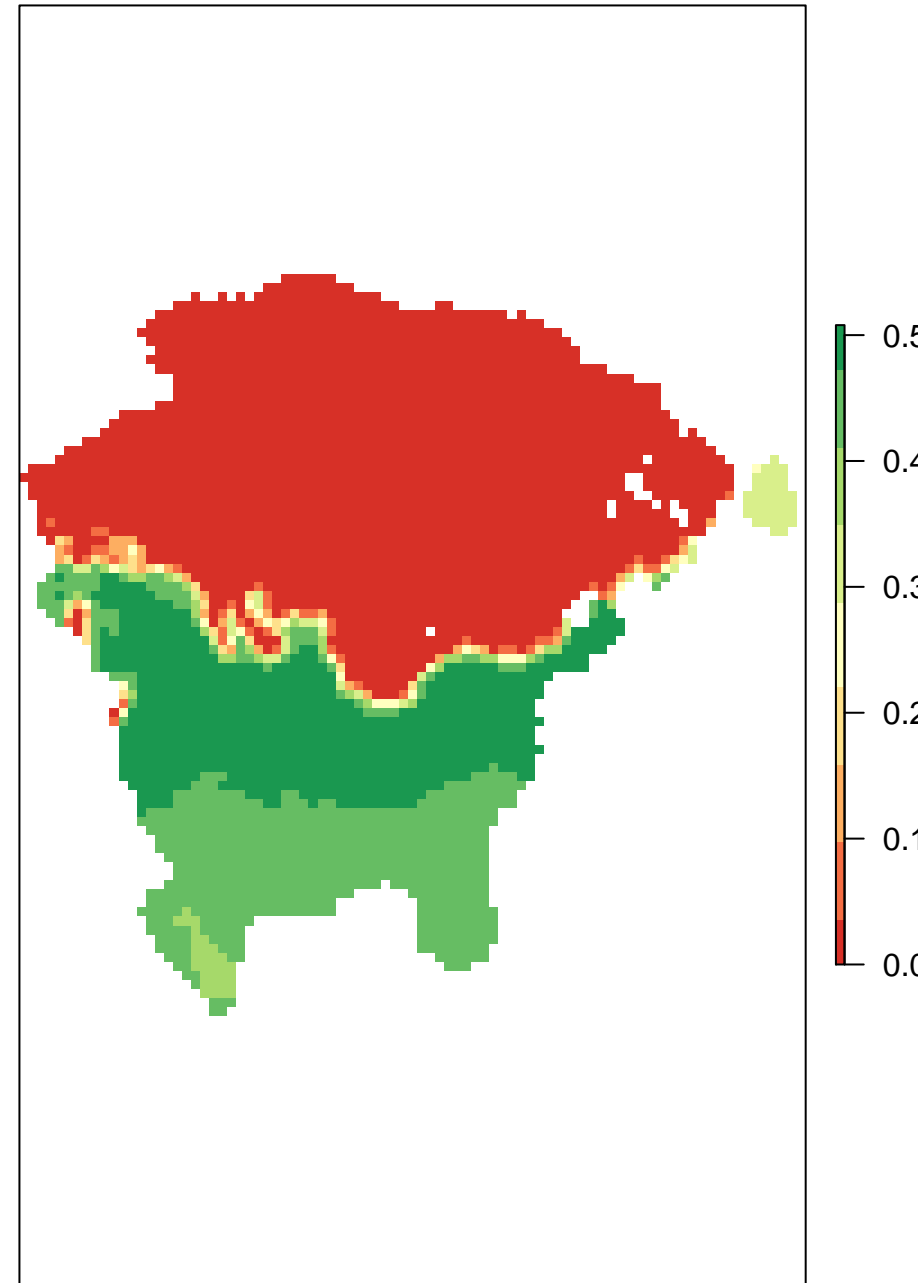


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X11000.ybp

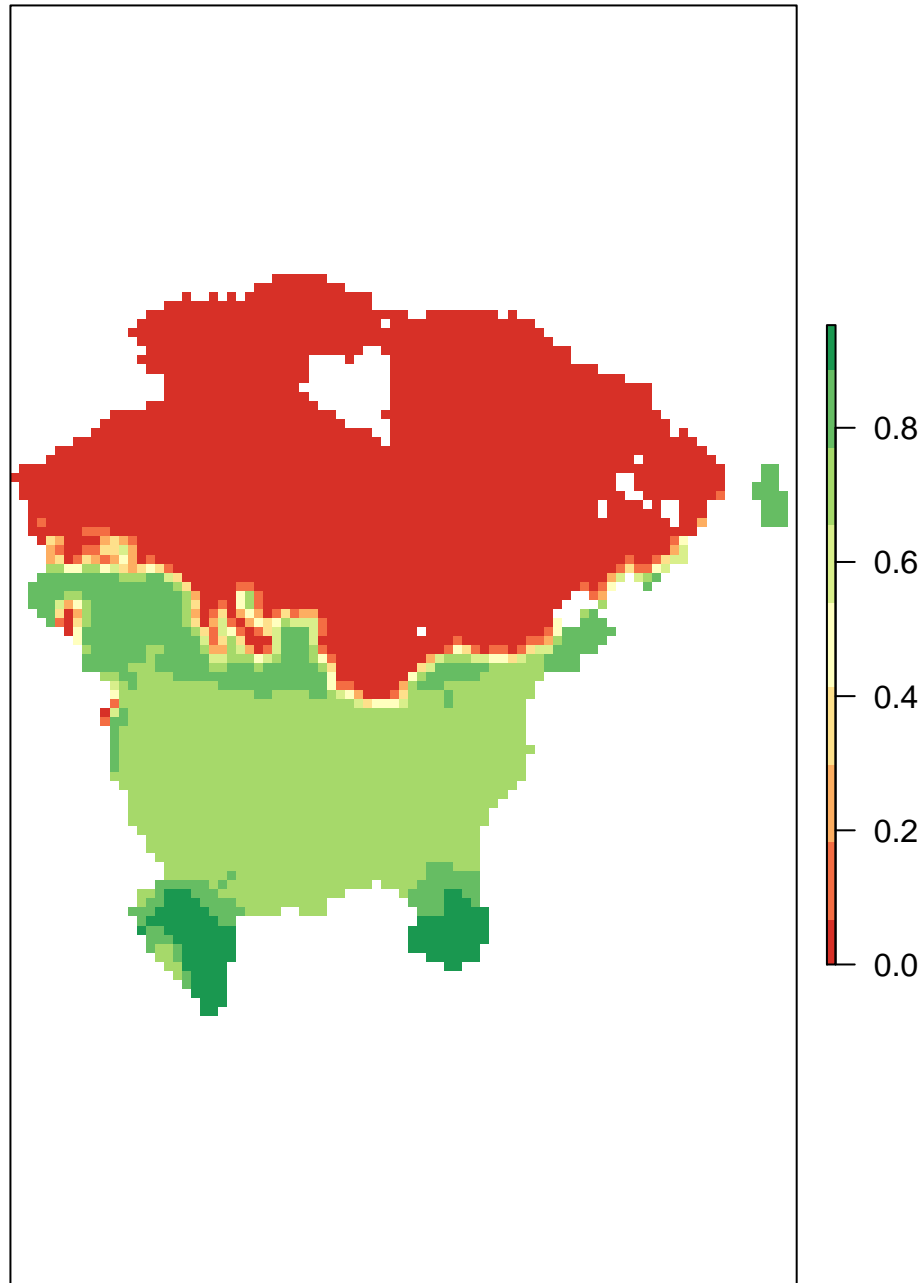


MAX without sp removed, X11000.ybp

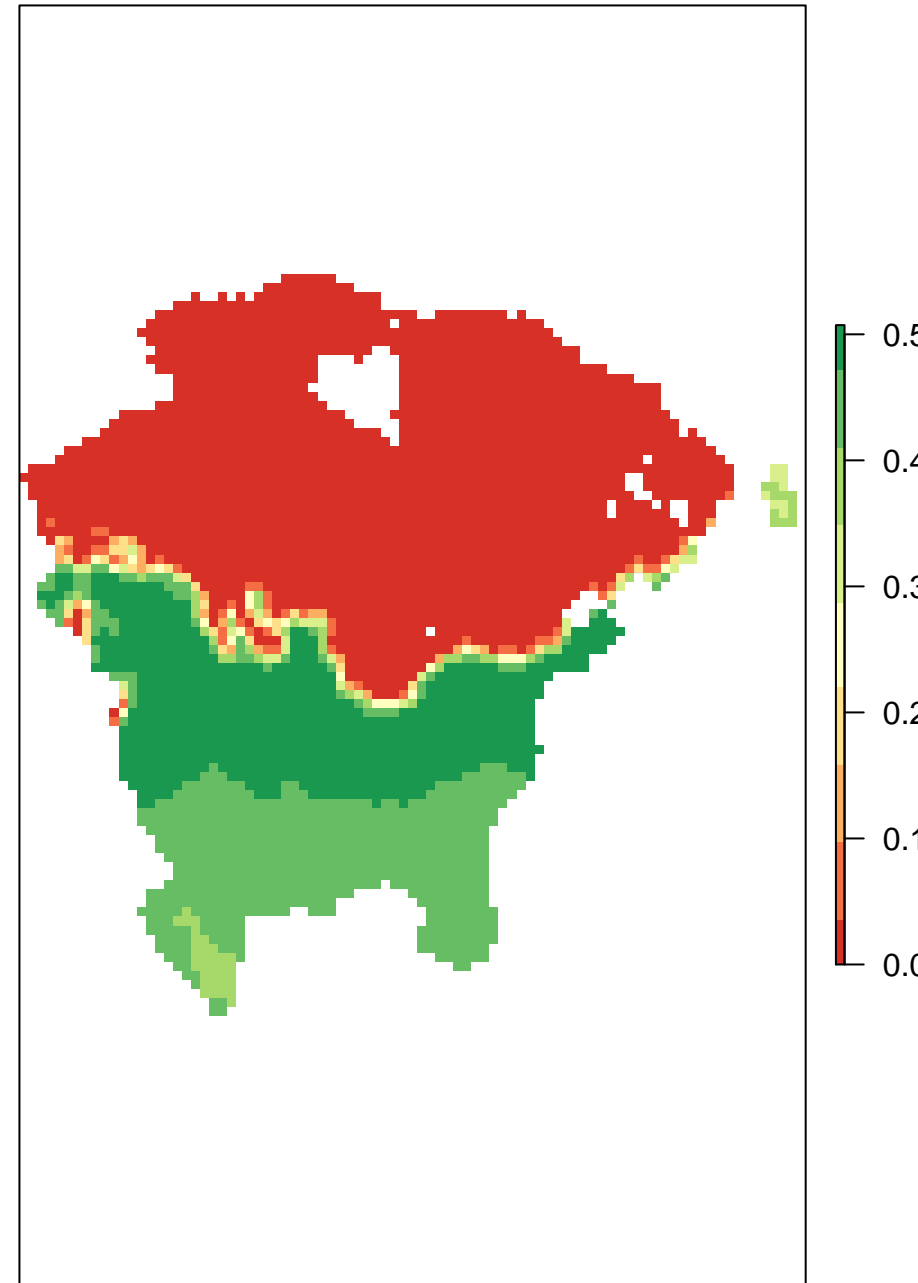


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X10000.ybp

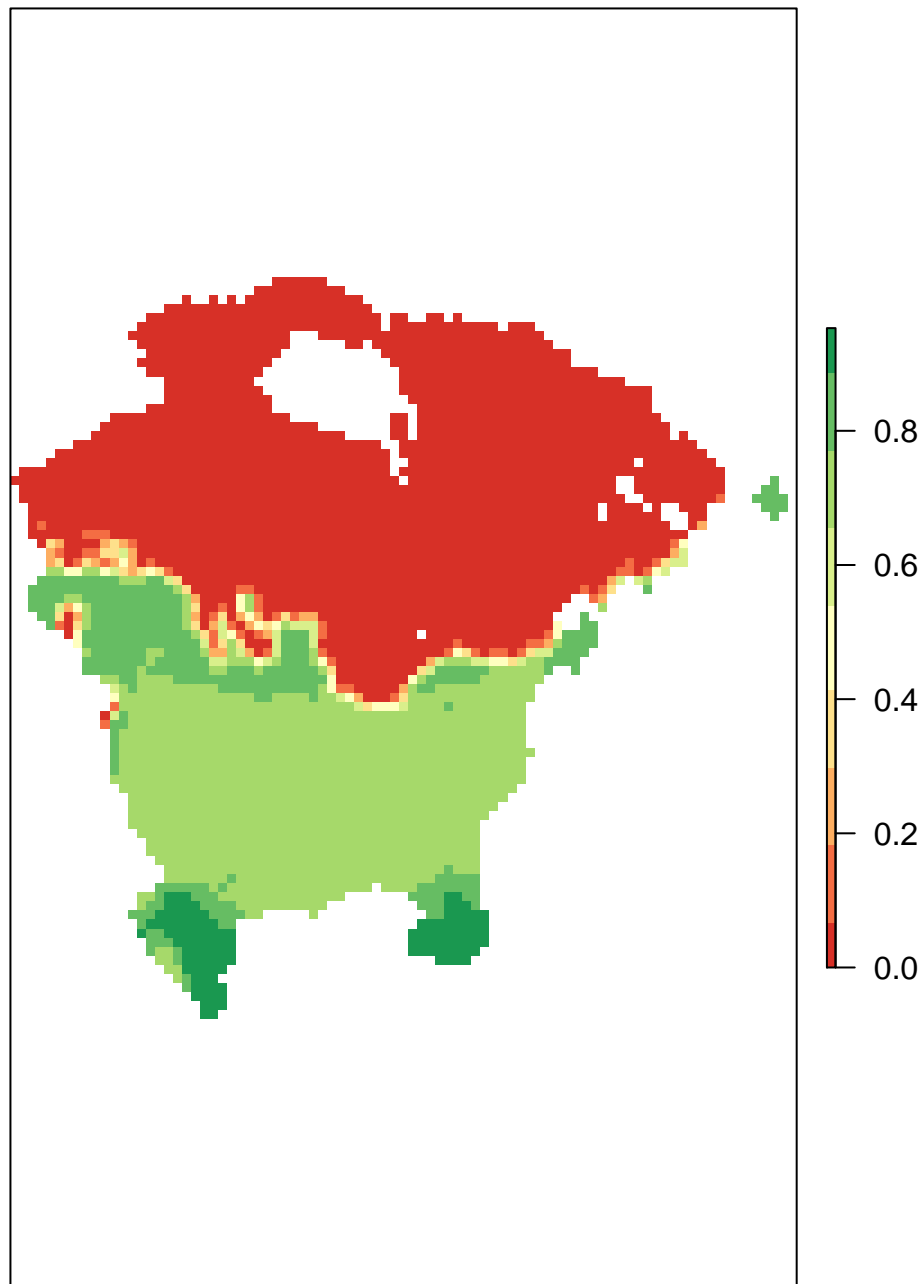


MAX without sp removed, X10000.ybp

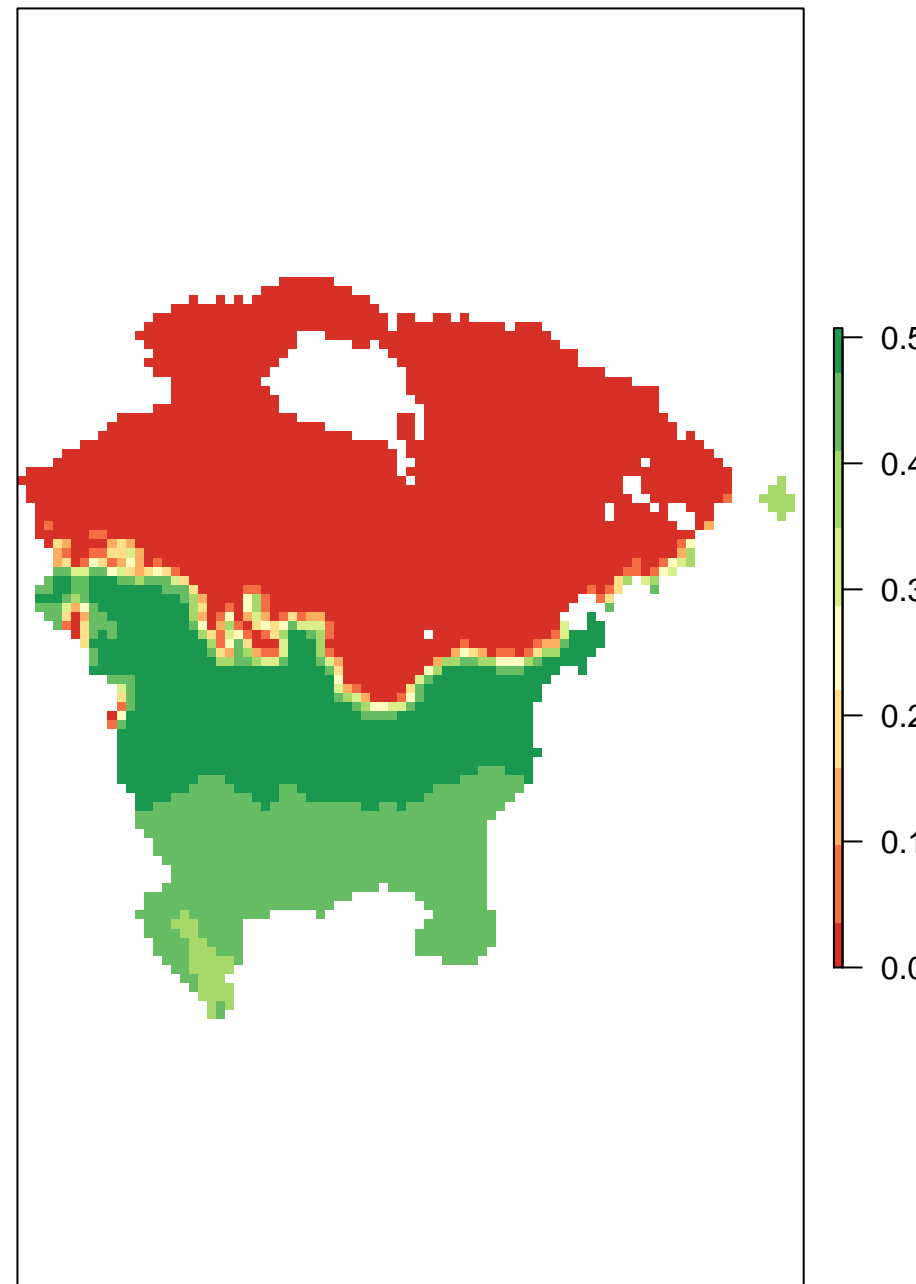


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X9000.ybp

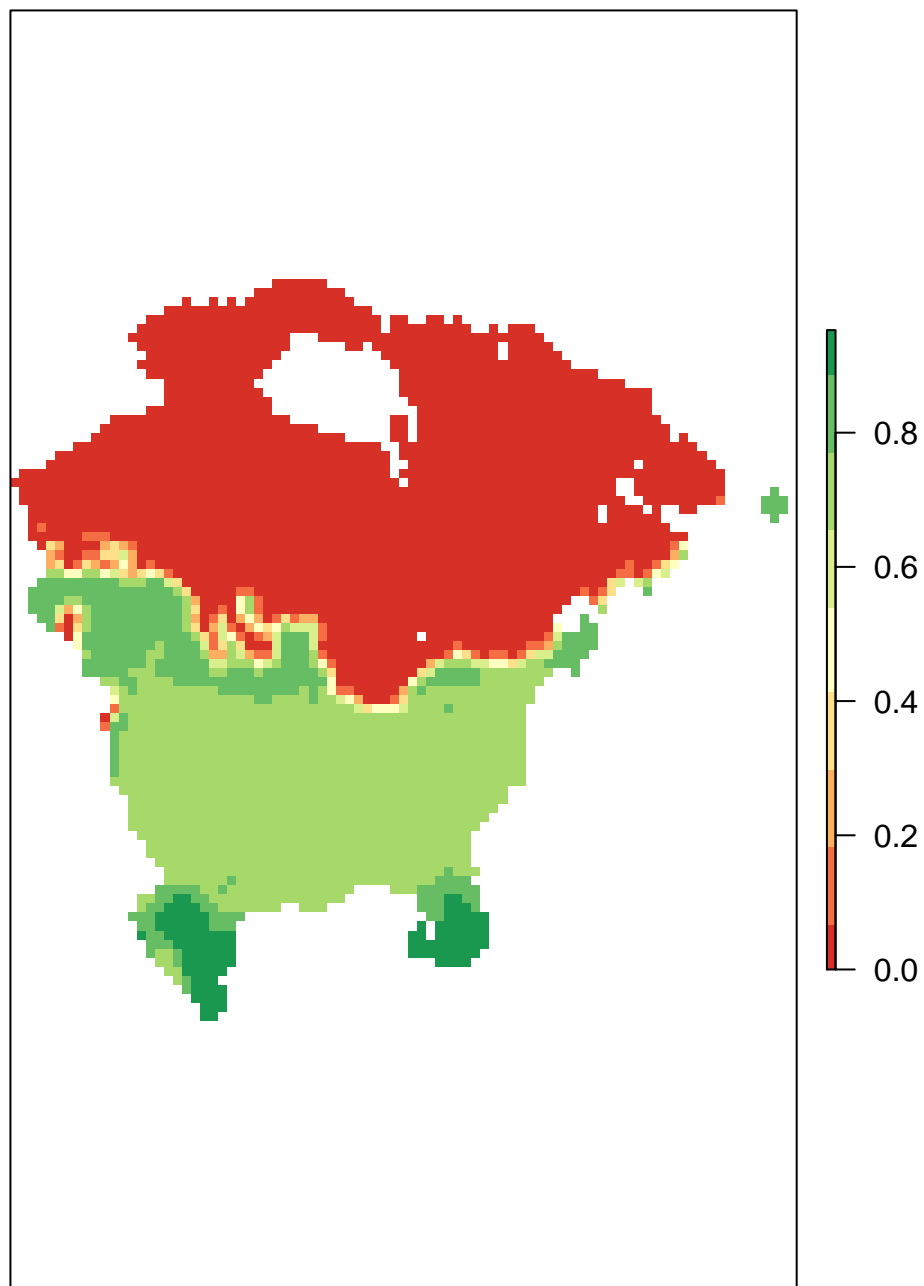


MAX without sp removed, X9000.ybp

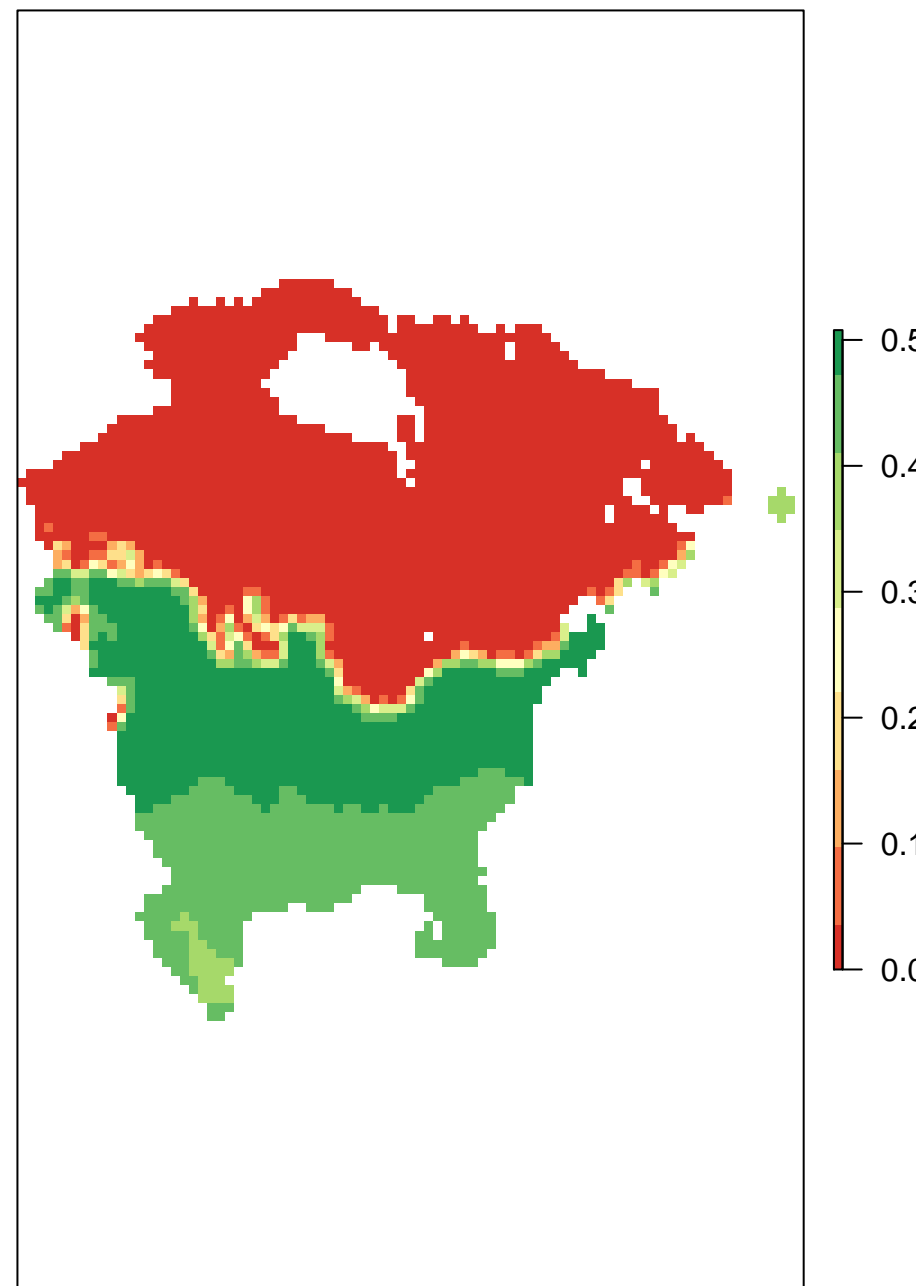


Species skipped = *Fraxinus americana*, GCM = *ecbilt*

MAX, X8000.ybp

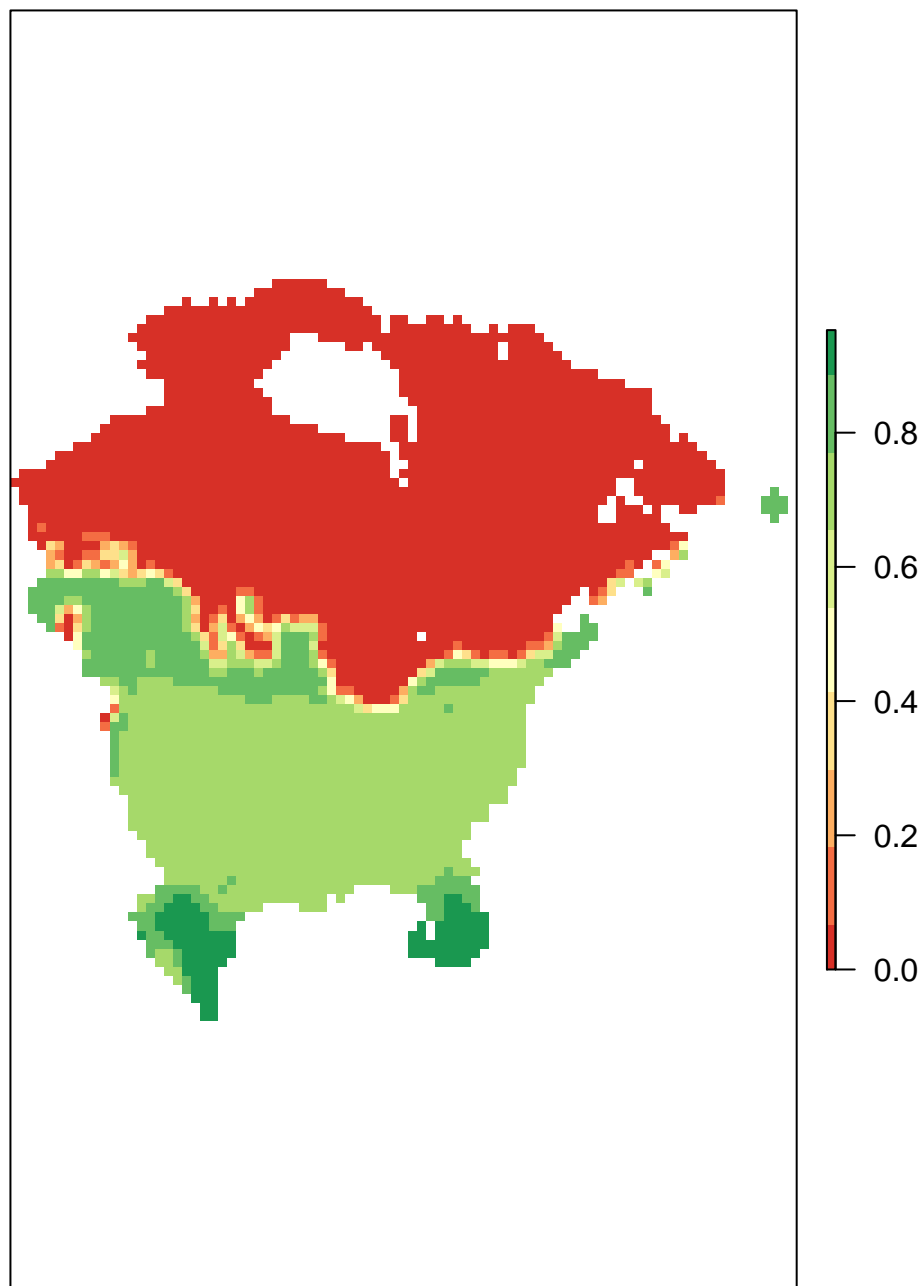


MAX without sp removed, X8000.ybp

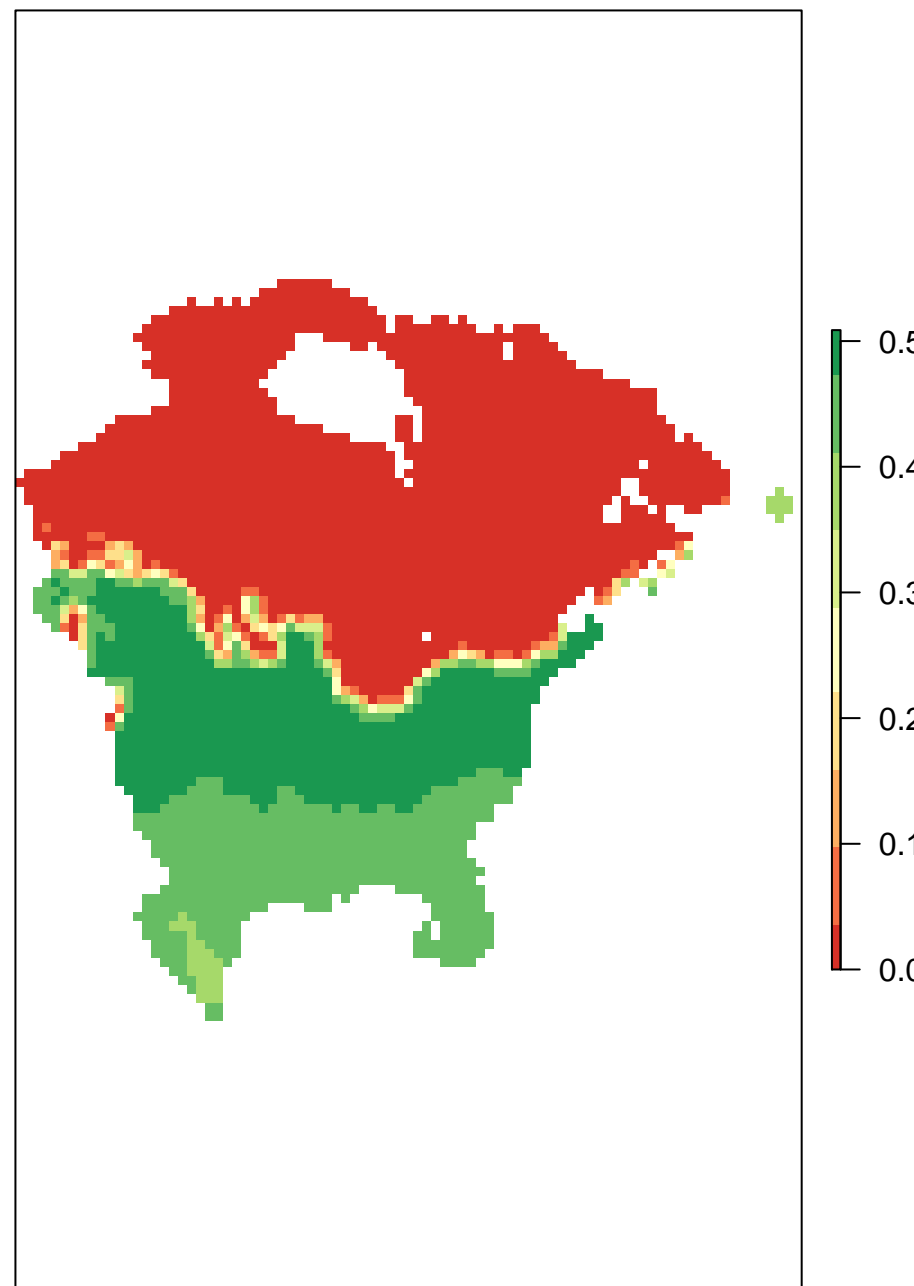


Species skipped = Fraxinus americana, GCM = ecbilt

MAX, X7000.ybp

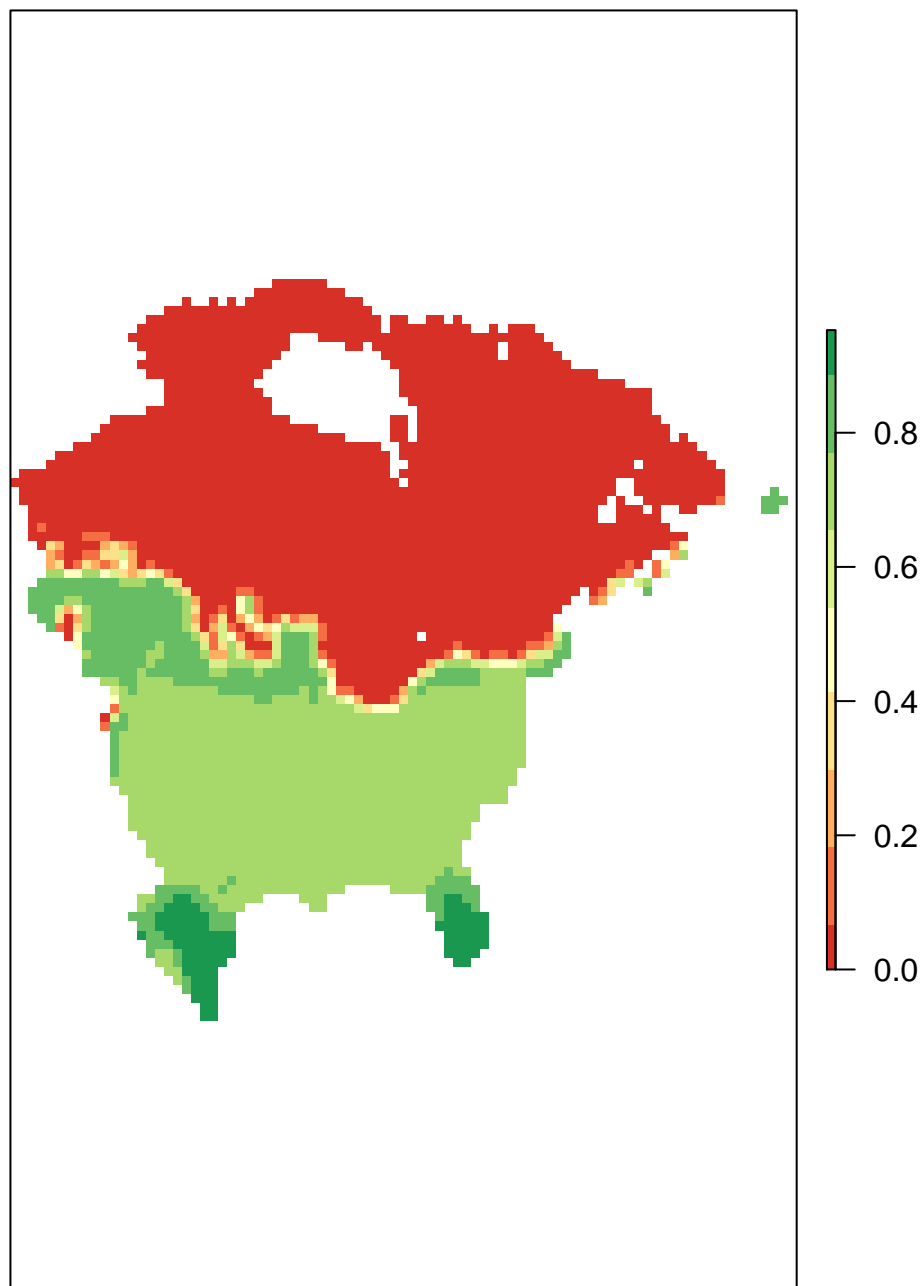


MAX without sp removed, X7000.ybp

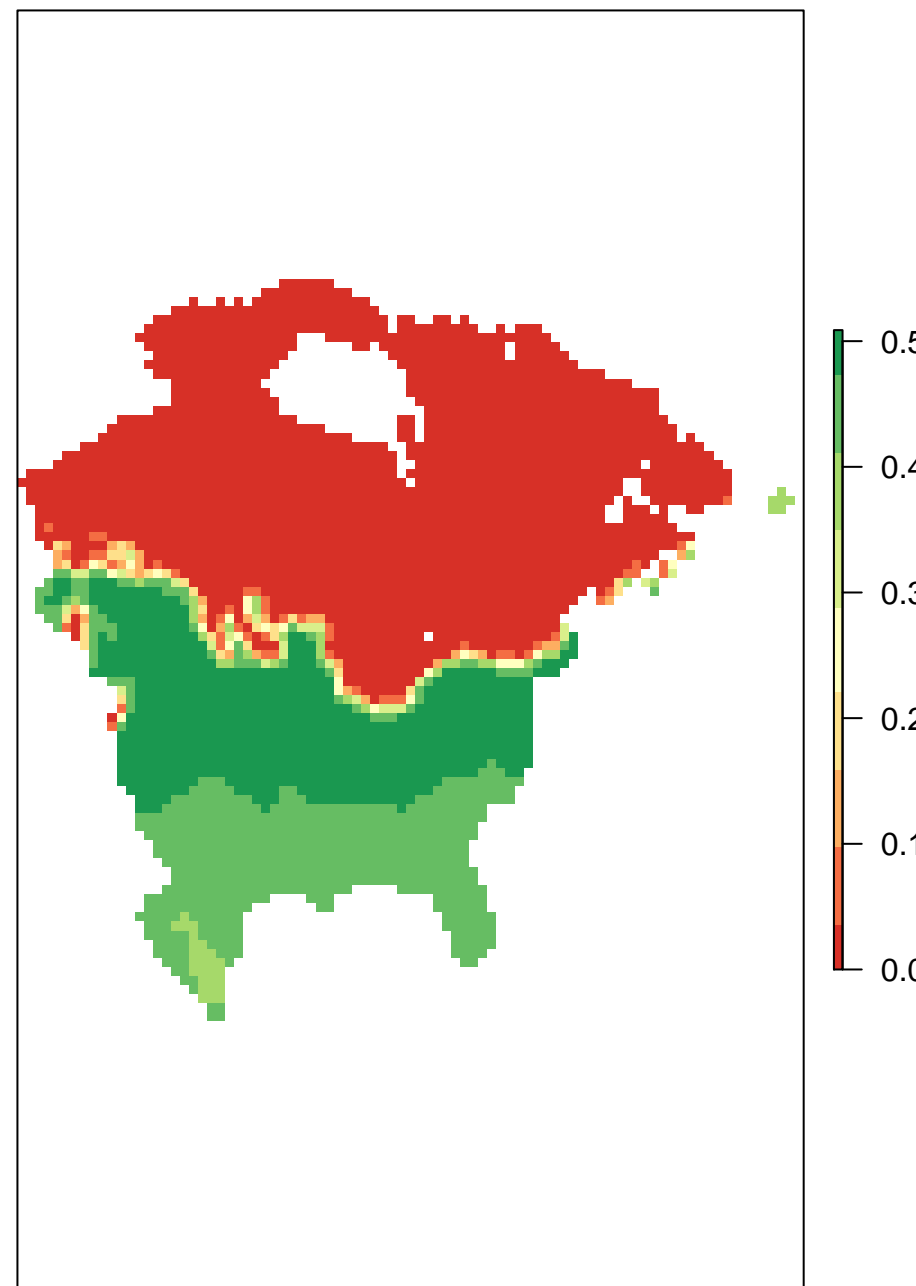


Species skipped = Fraxinus americana, GCM = ecbilt

MAX, X6000.ybp

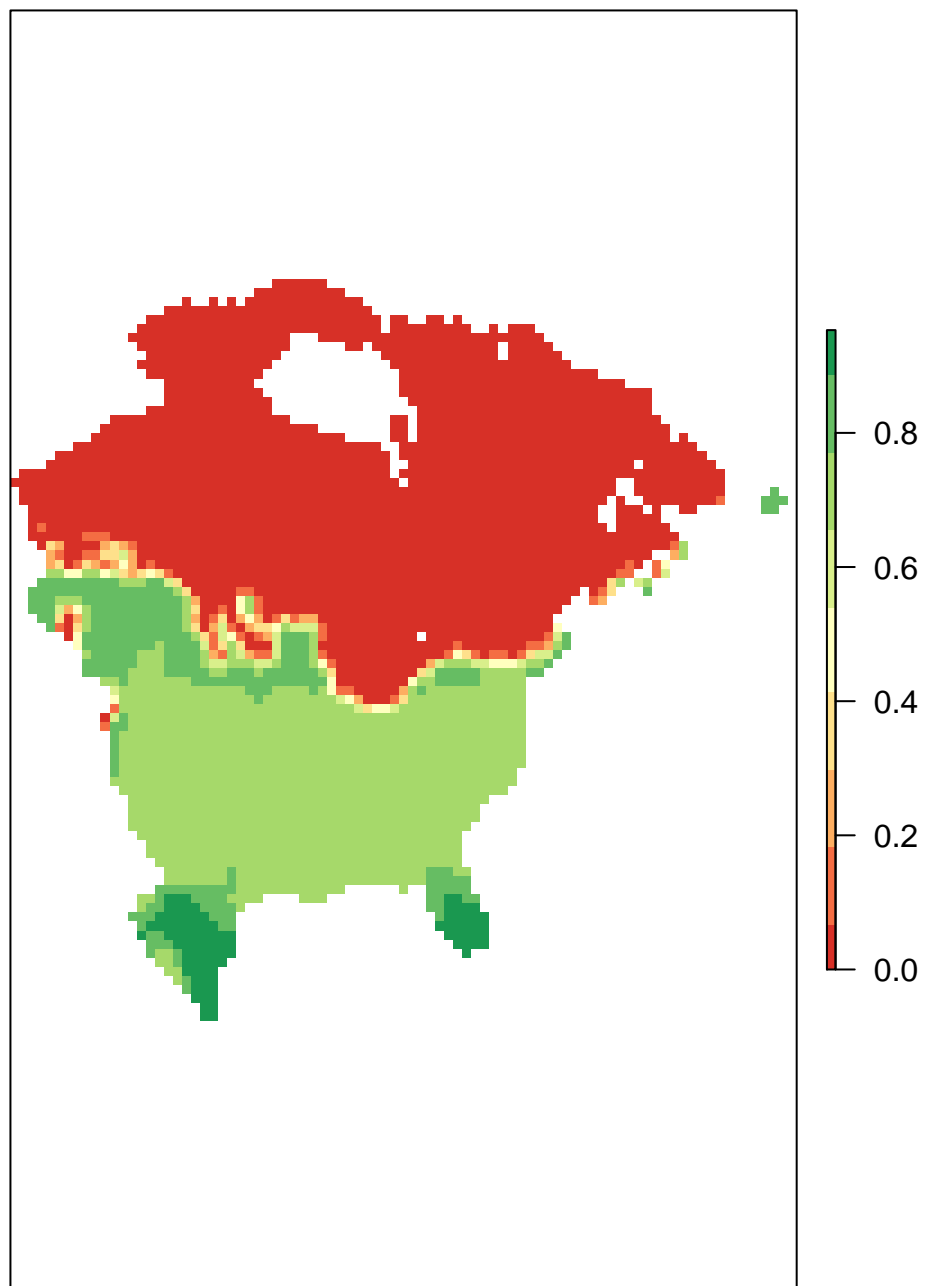


MAX without sp removed, X6000.ybp

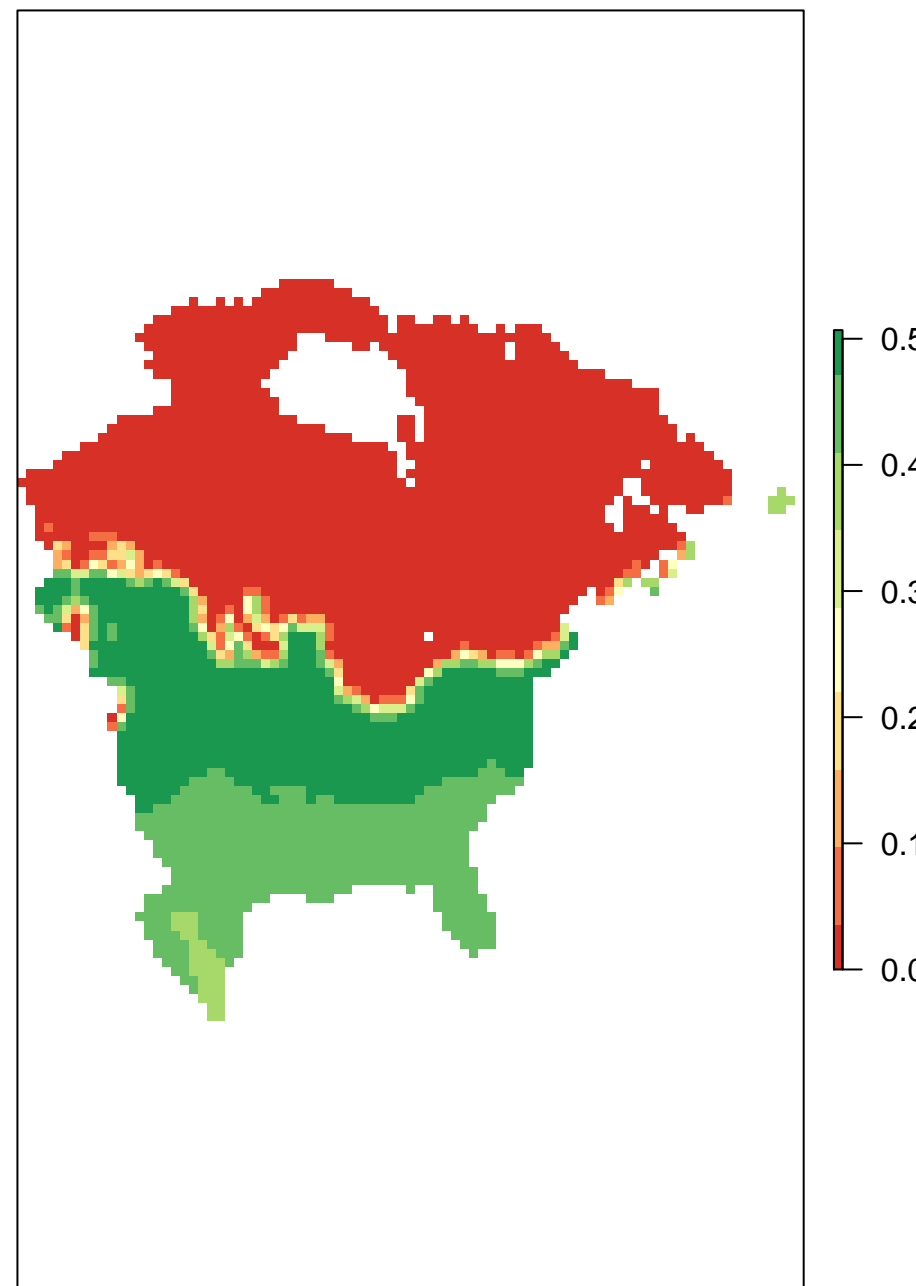


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X5000.ybp

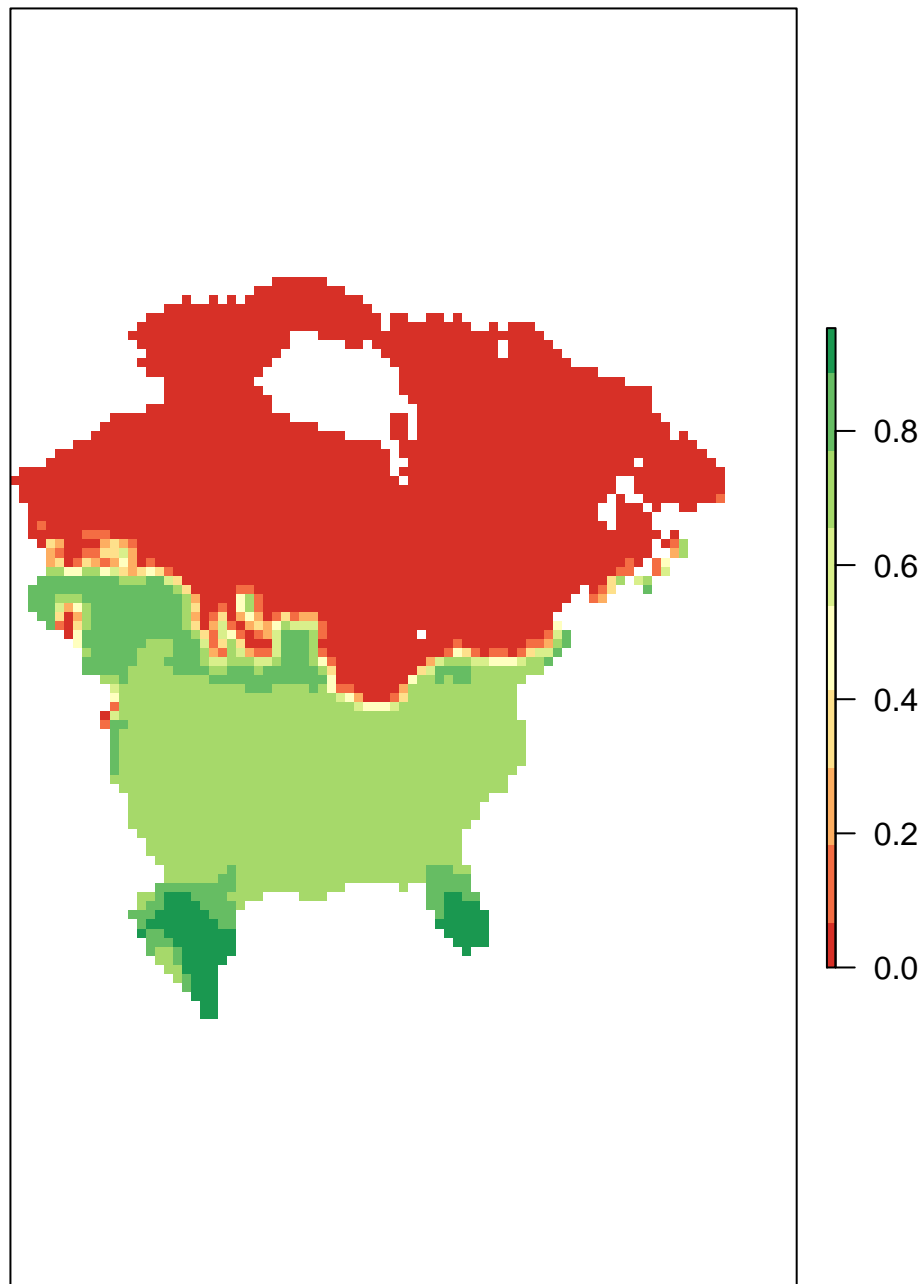


MAX without sp removed, X5000.ybp

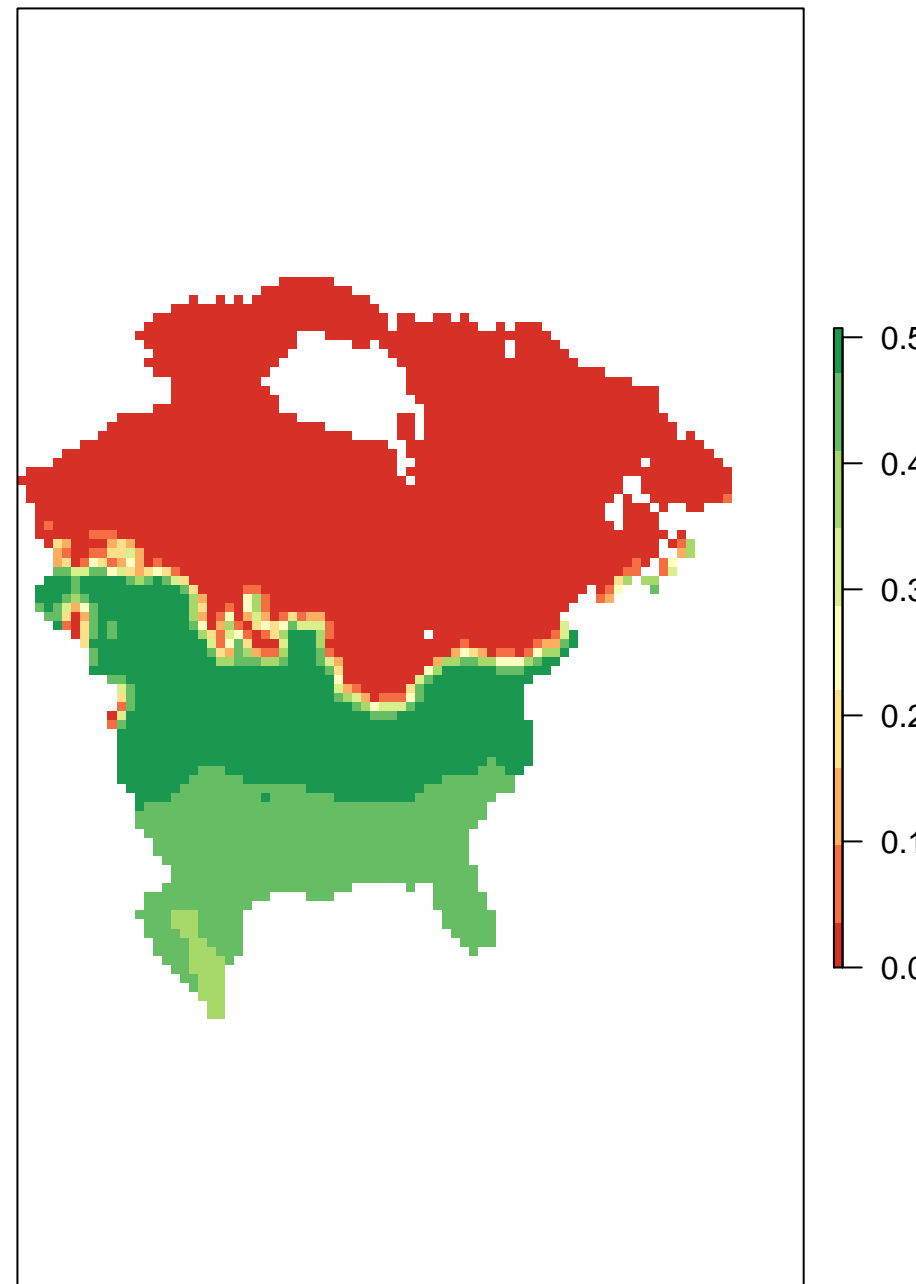


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X4000.ybp

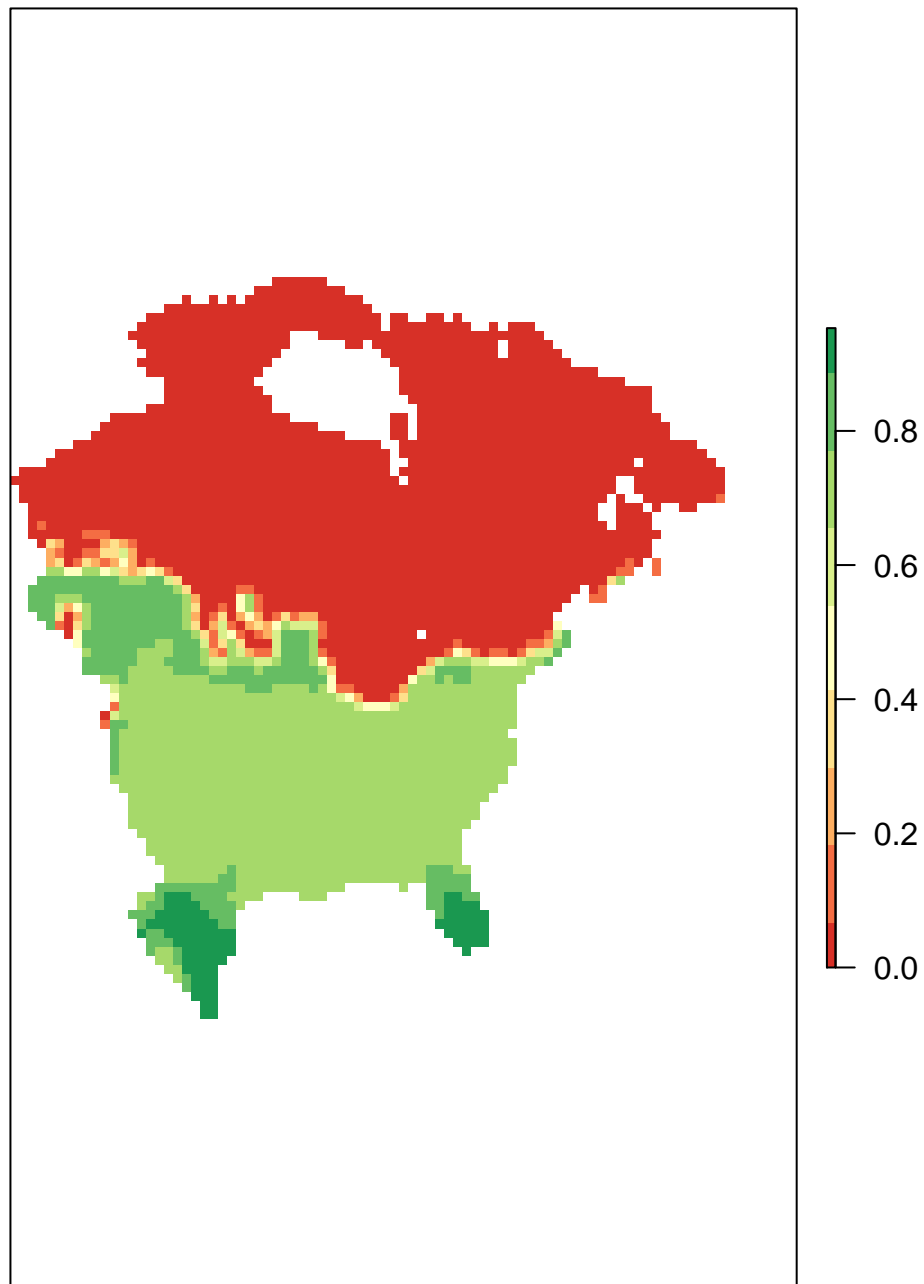


MAX without sp removed, X4000.ybp

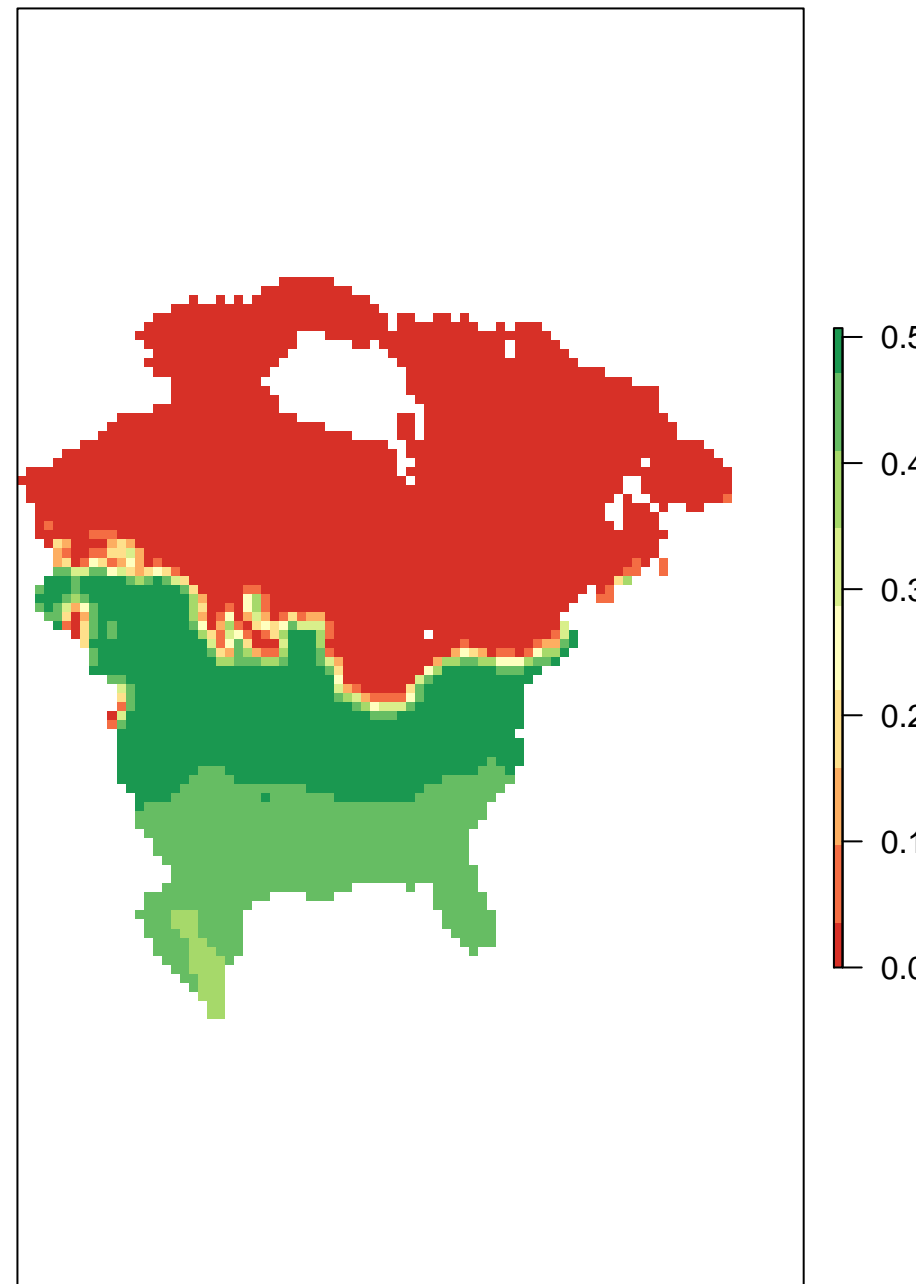


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X3000.ybp

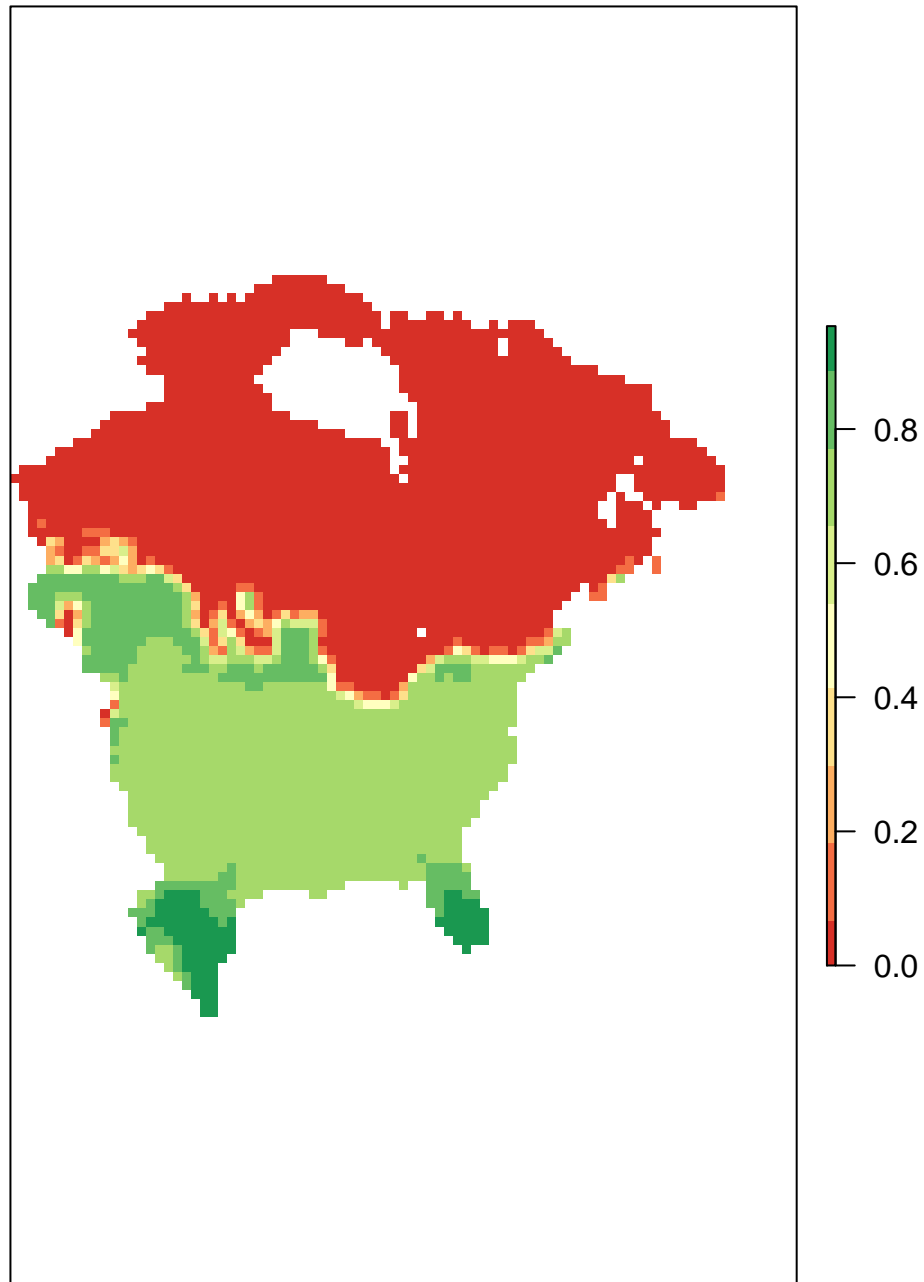


MAX without sp removed, X3000.ybp

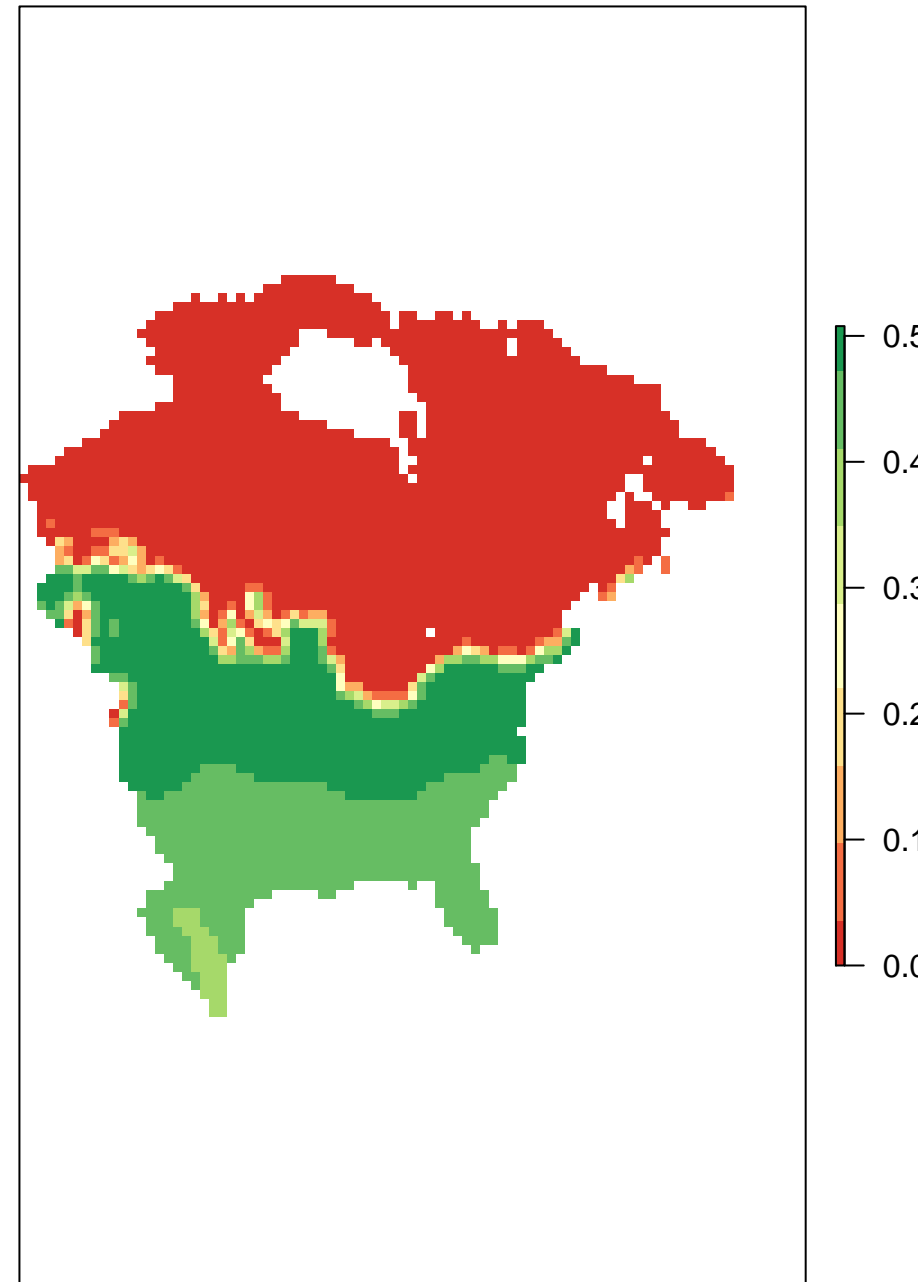


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X2000.ybp

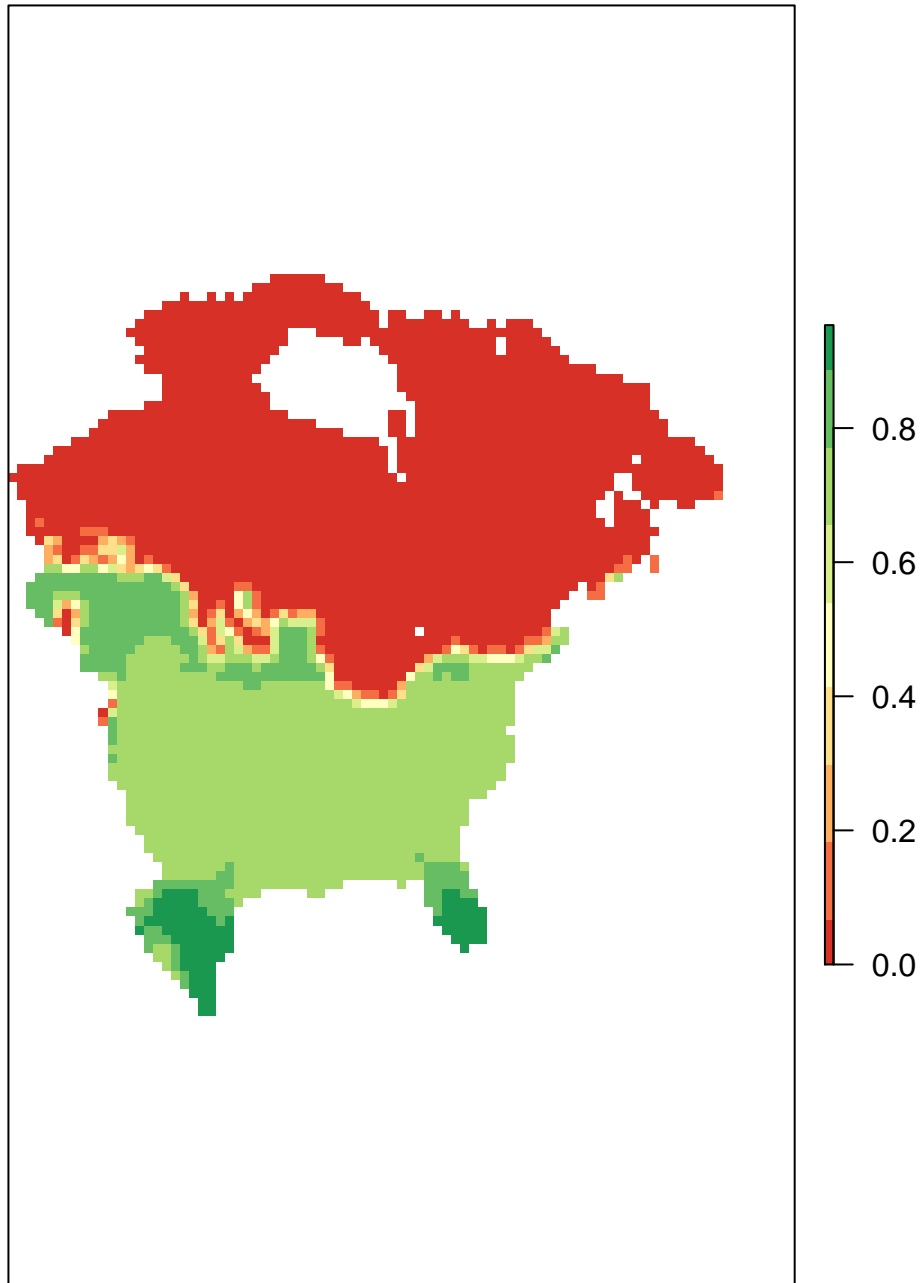


MAX without sp removed, X2000.ybp

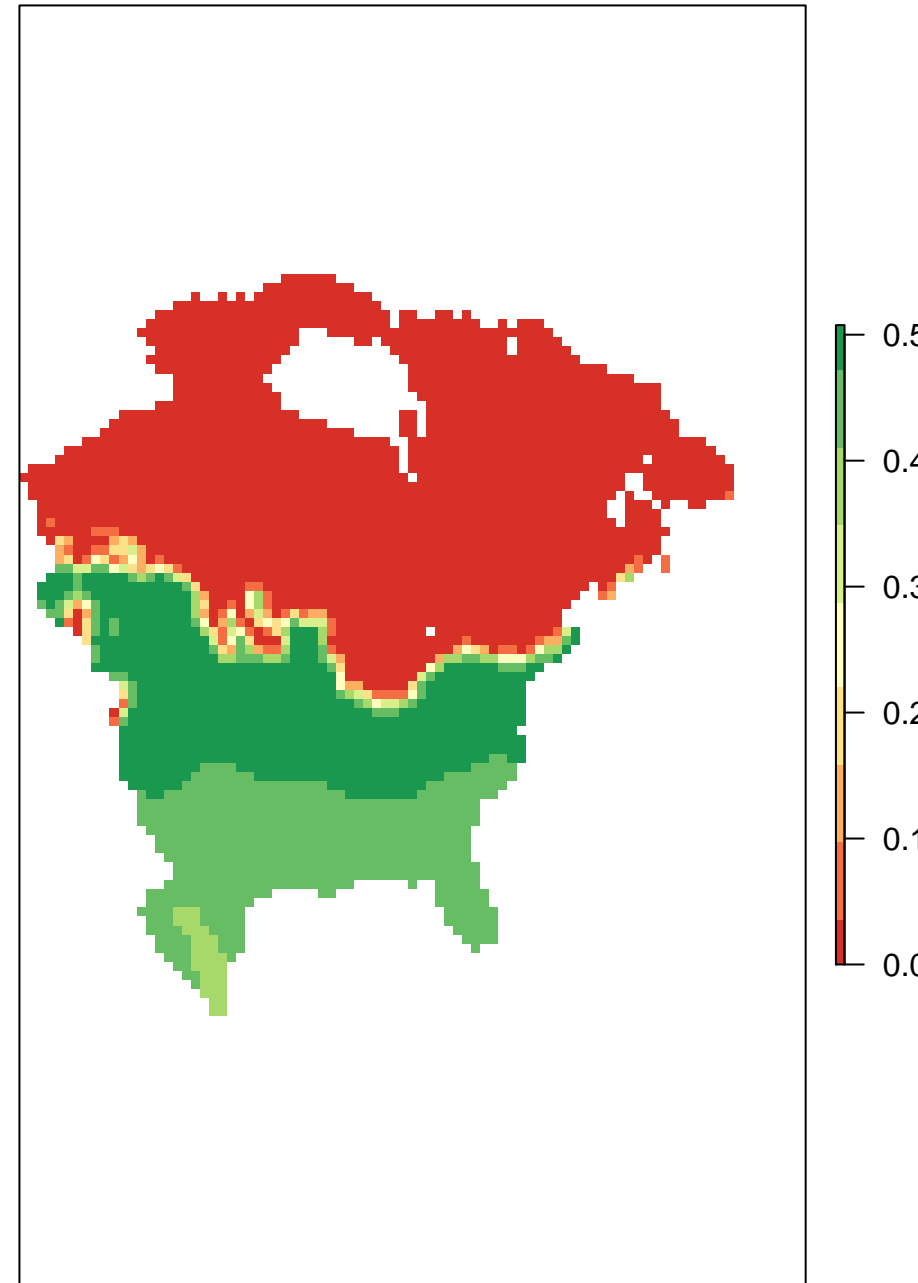


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X1000.ybp

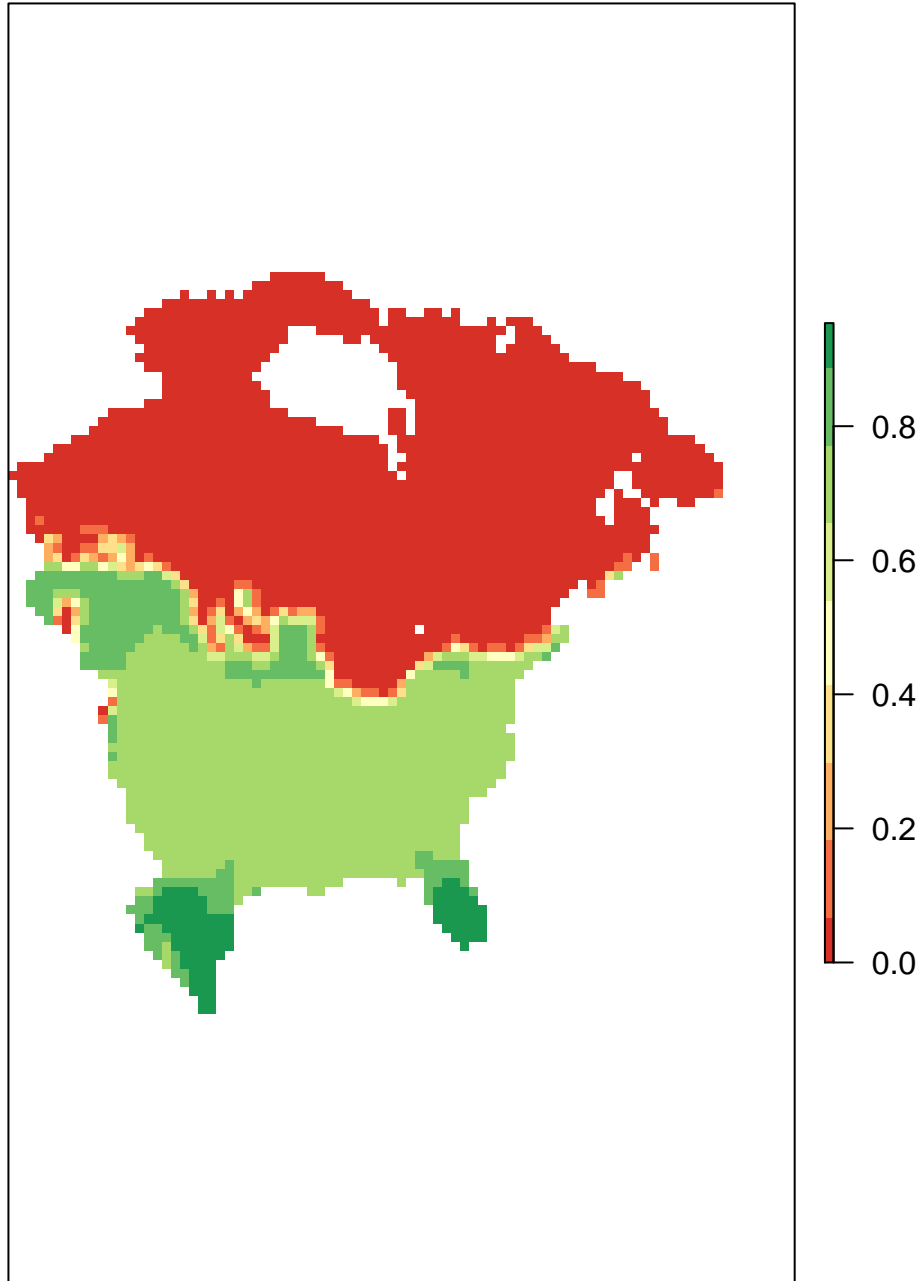


MAX without sp removed, X1000.ybp

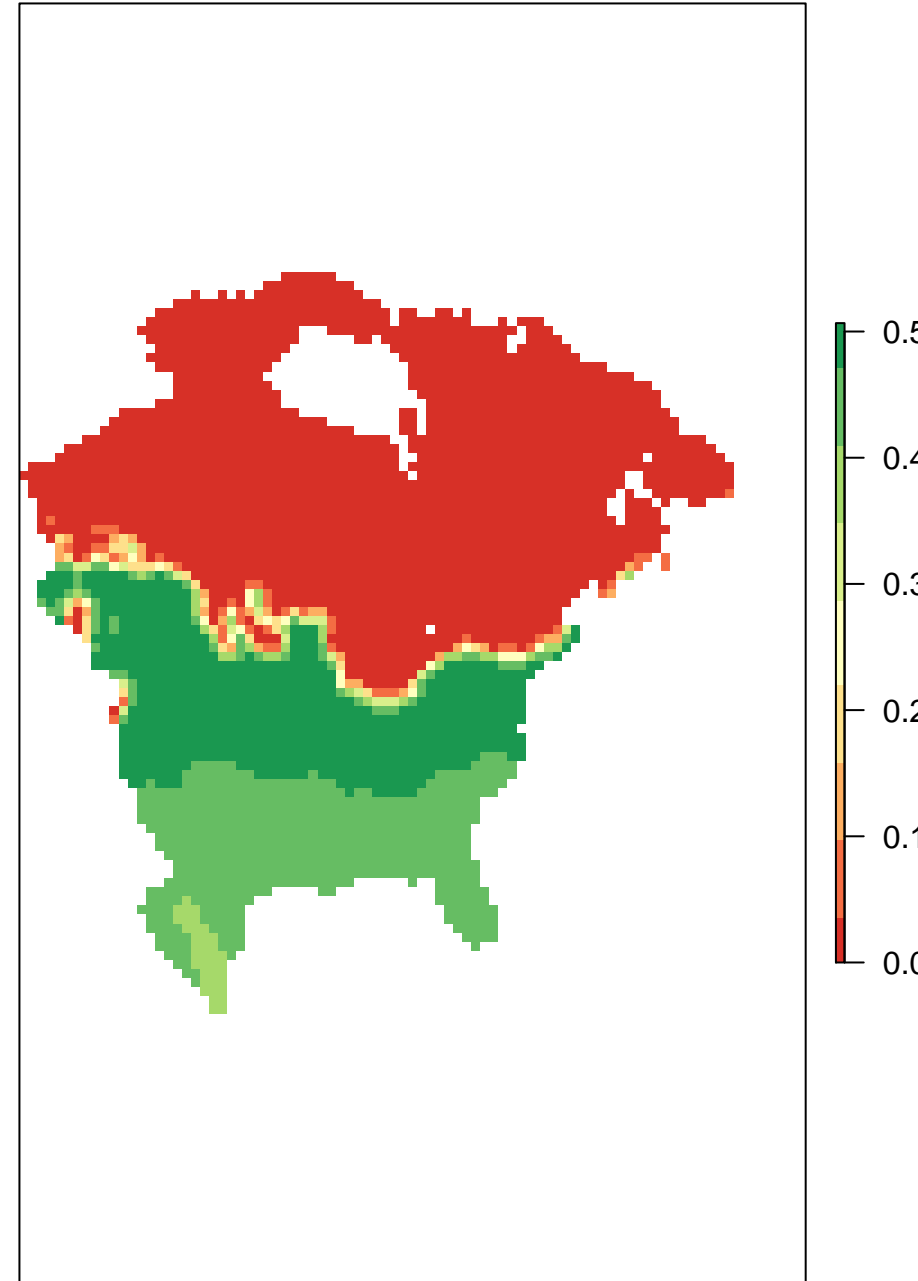


Species skipped = *Fraxinus americana*, GCM = ecbilt

MAX, X0.ybp



MAX without sp removed, X0.ybp



Species skipped = *Fraxinus americana*, GCM = ecbilt