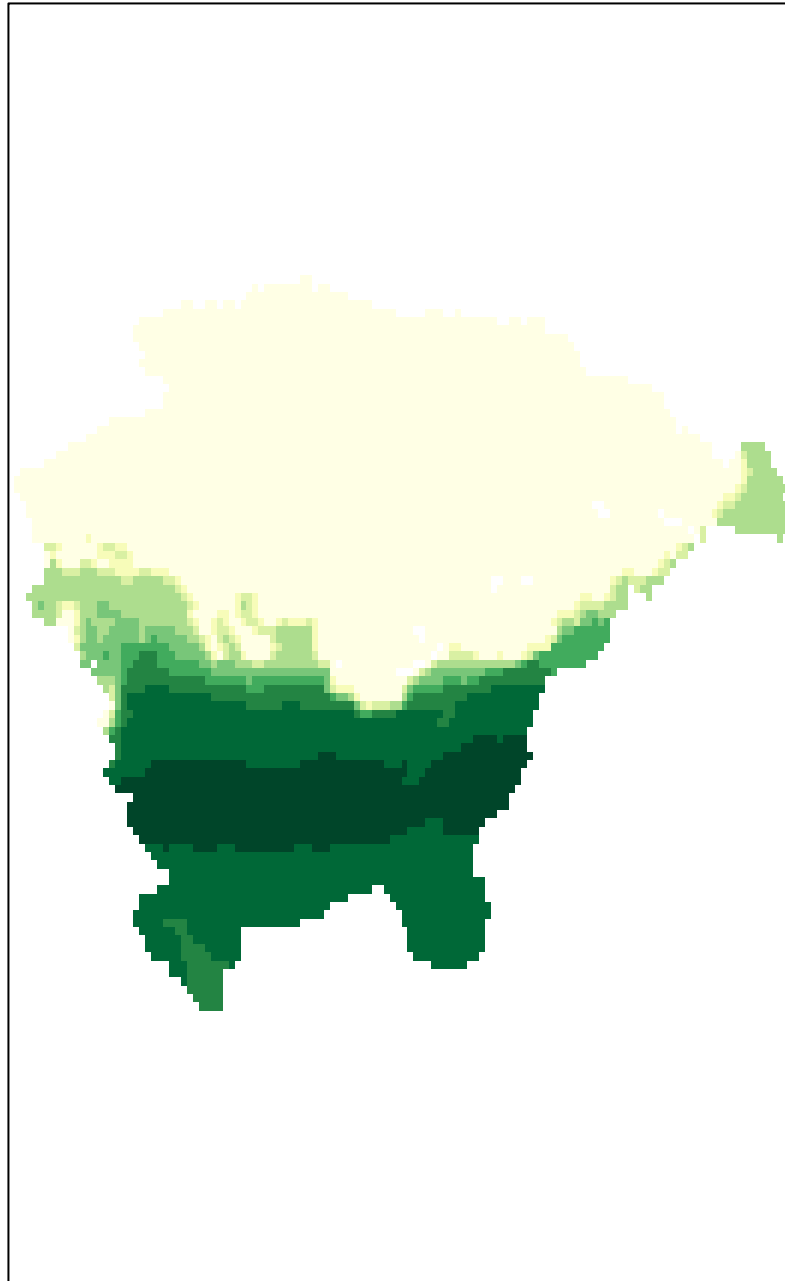
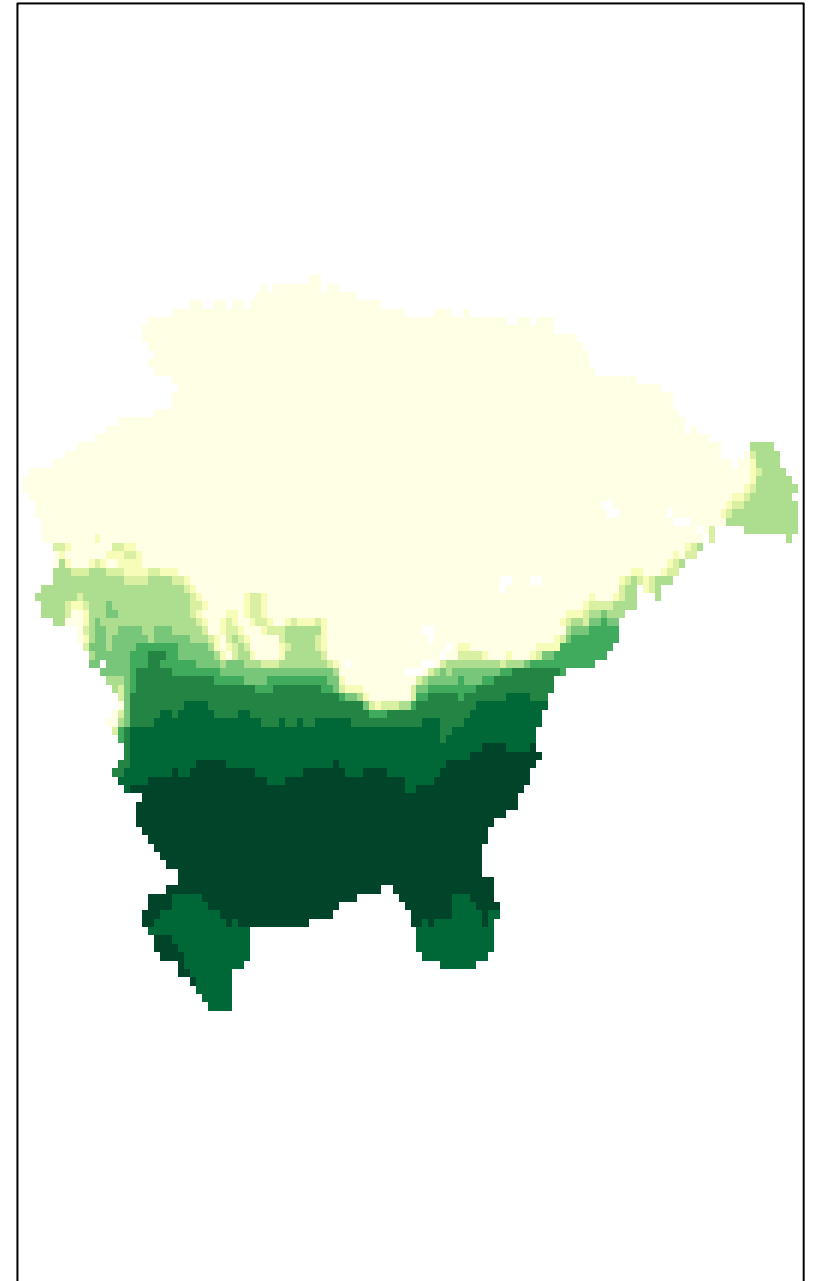


SUM, X21000.ybp

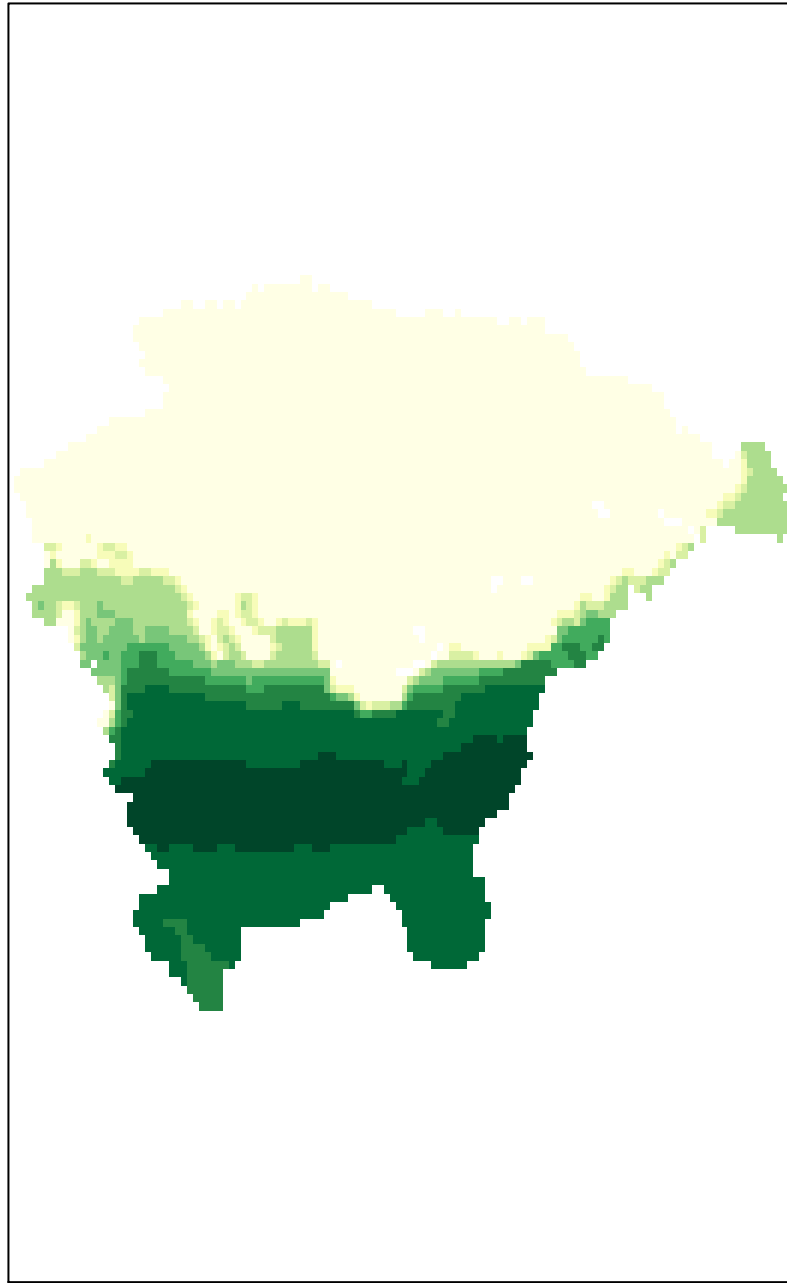


SUM, X21000.ybp

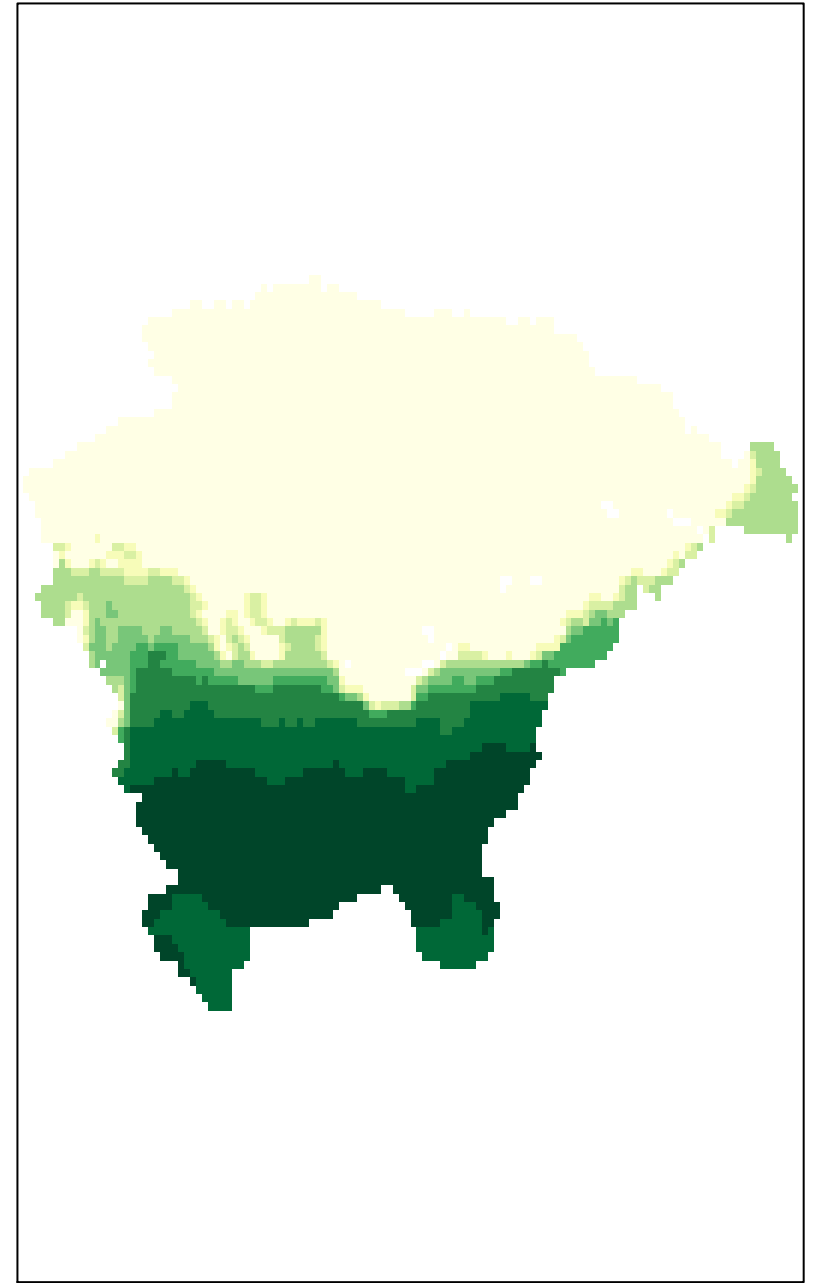


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X20000.ybp

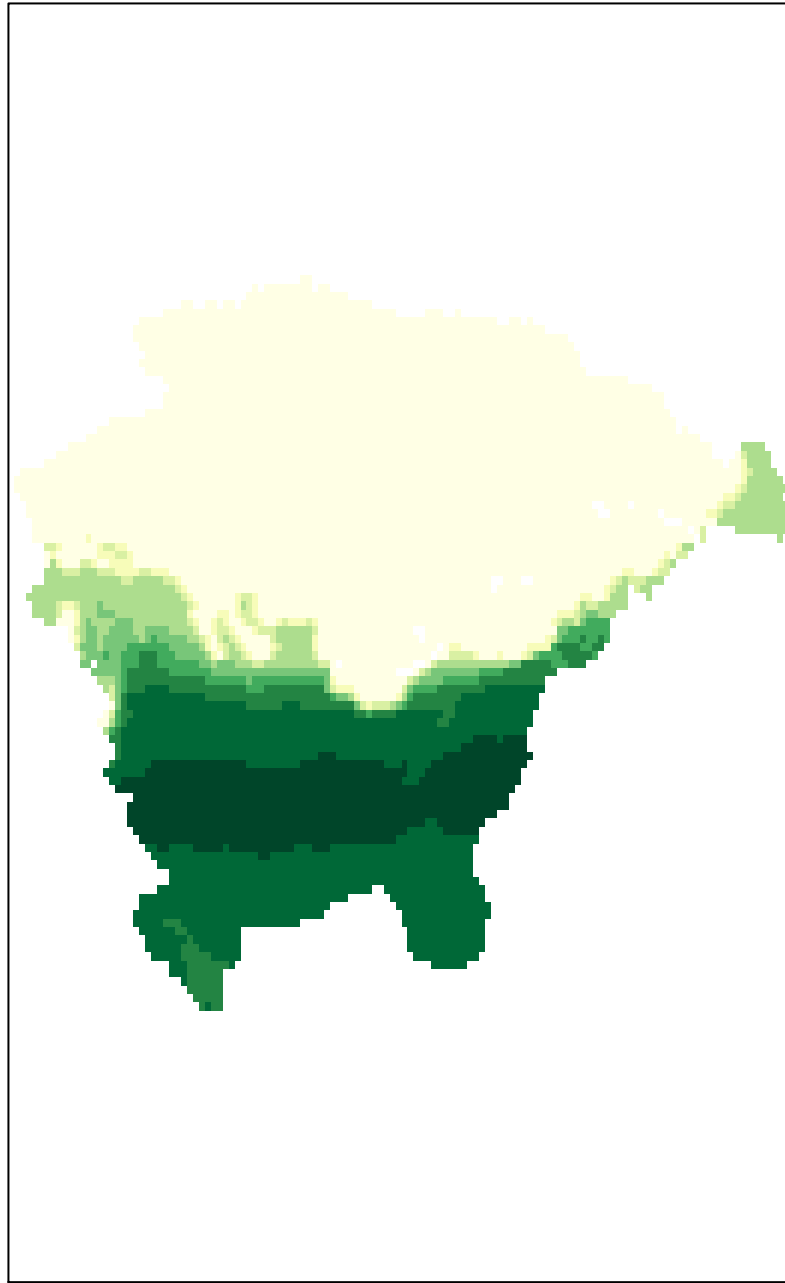


SUM, X20000.ybp

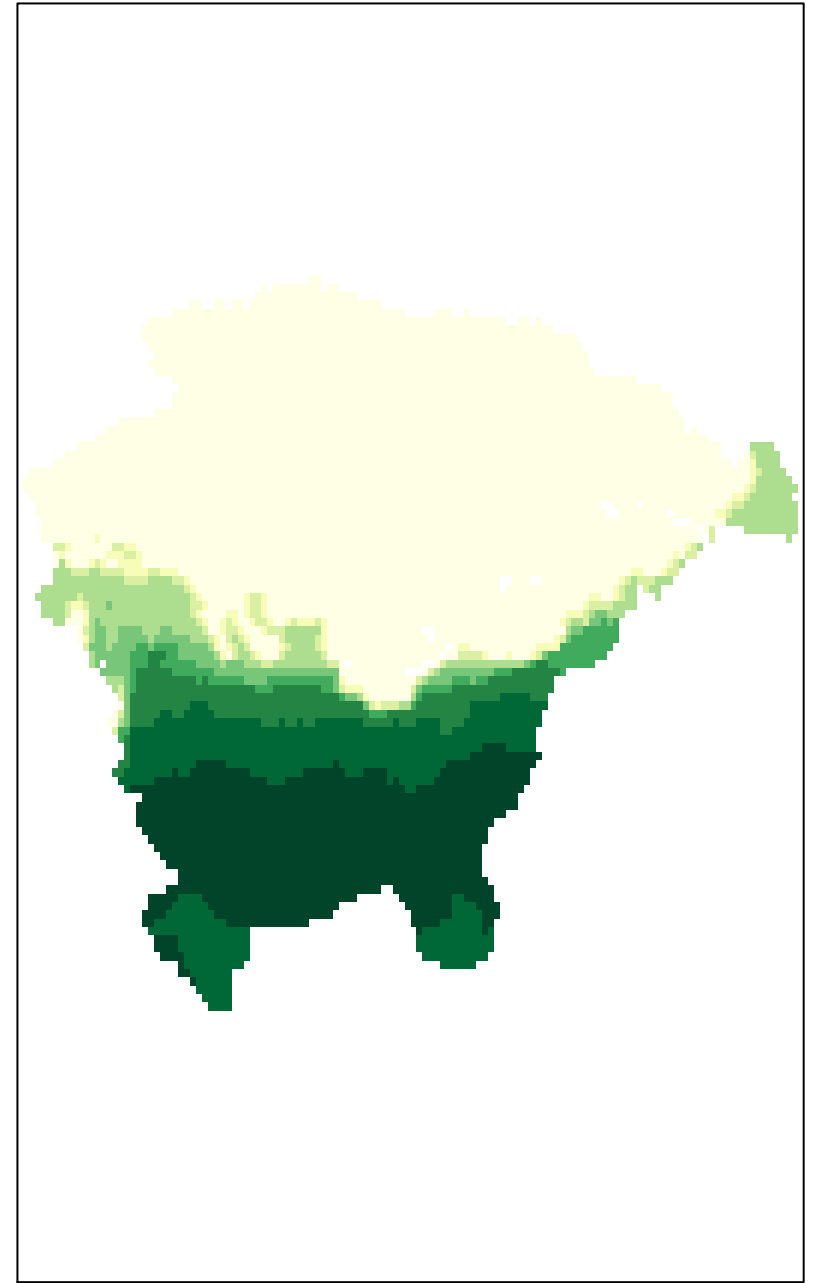


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SUM, X19000.ybp

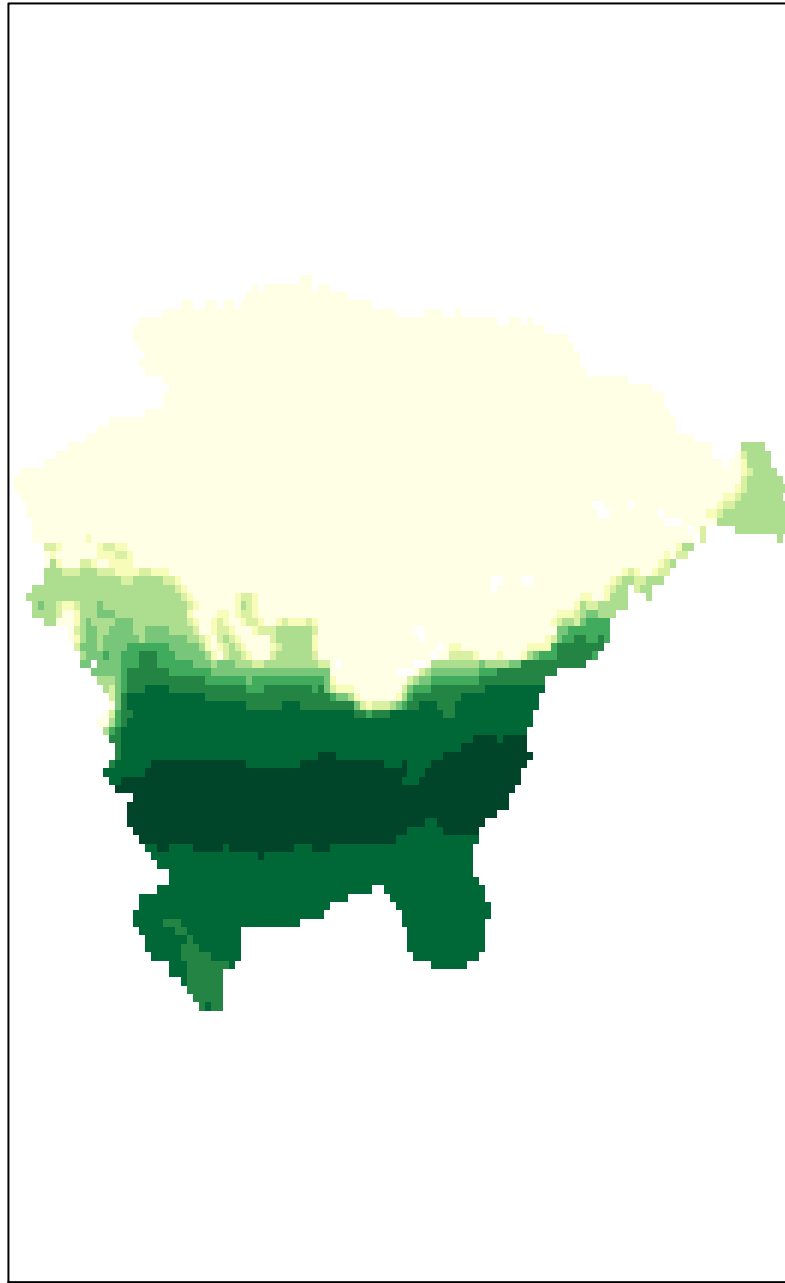


SUM, X19000.ybp

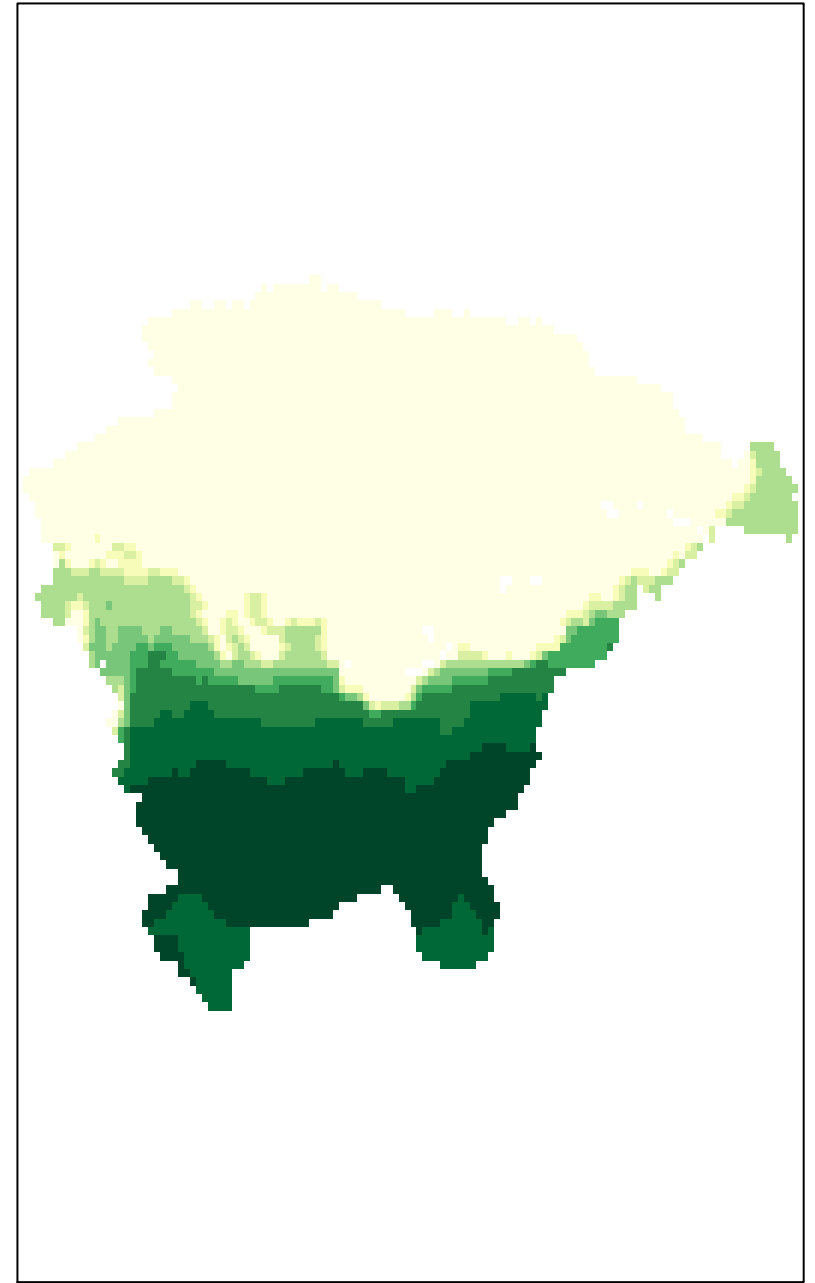


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X18000.ybp

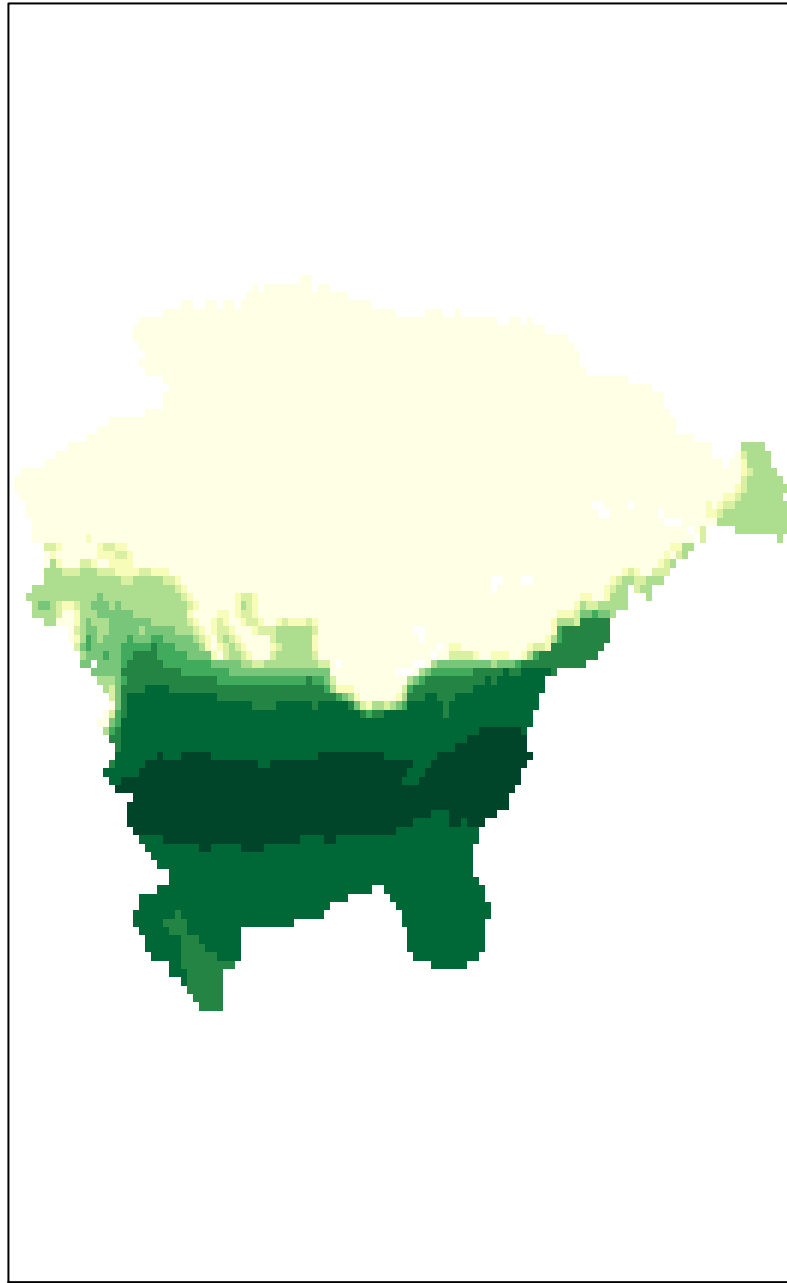


SUM, X18000.ybp

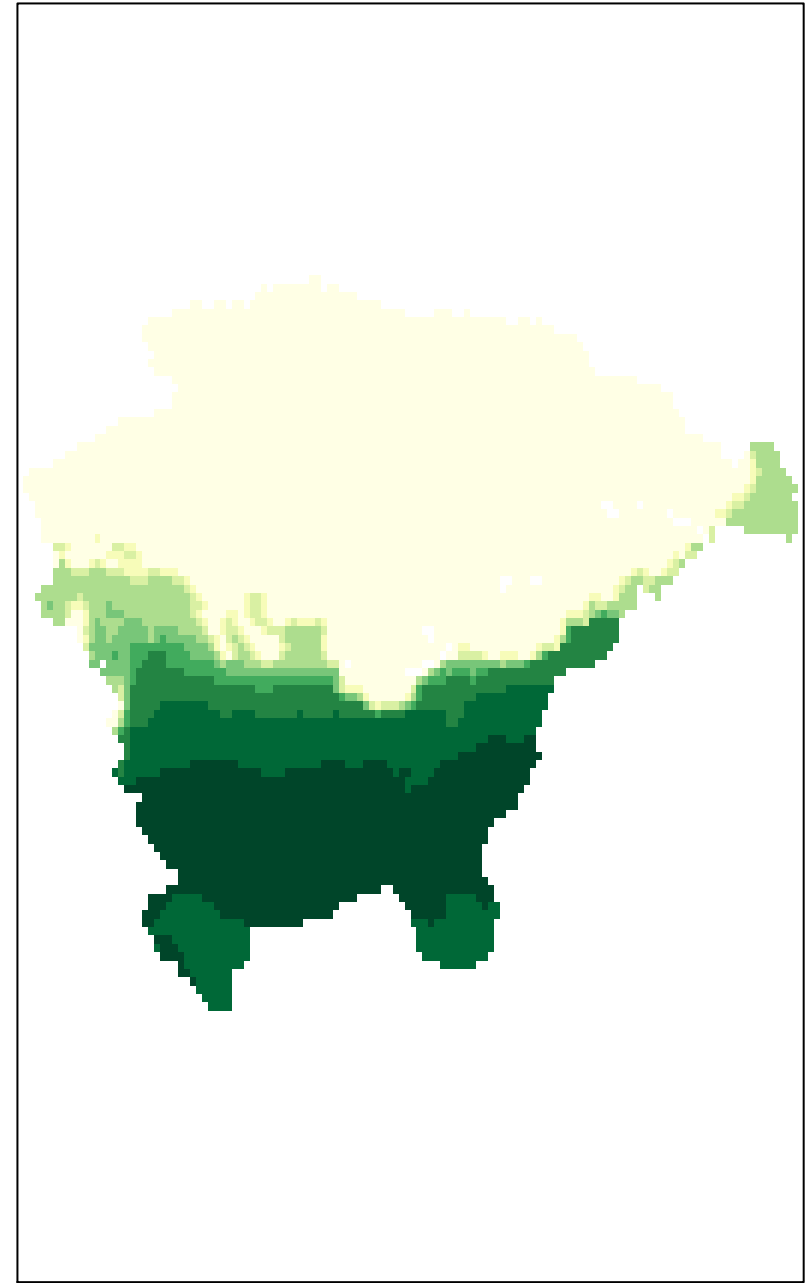


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X17000.ybp

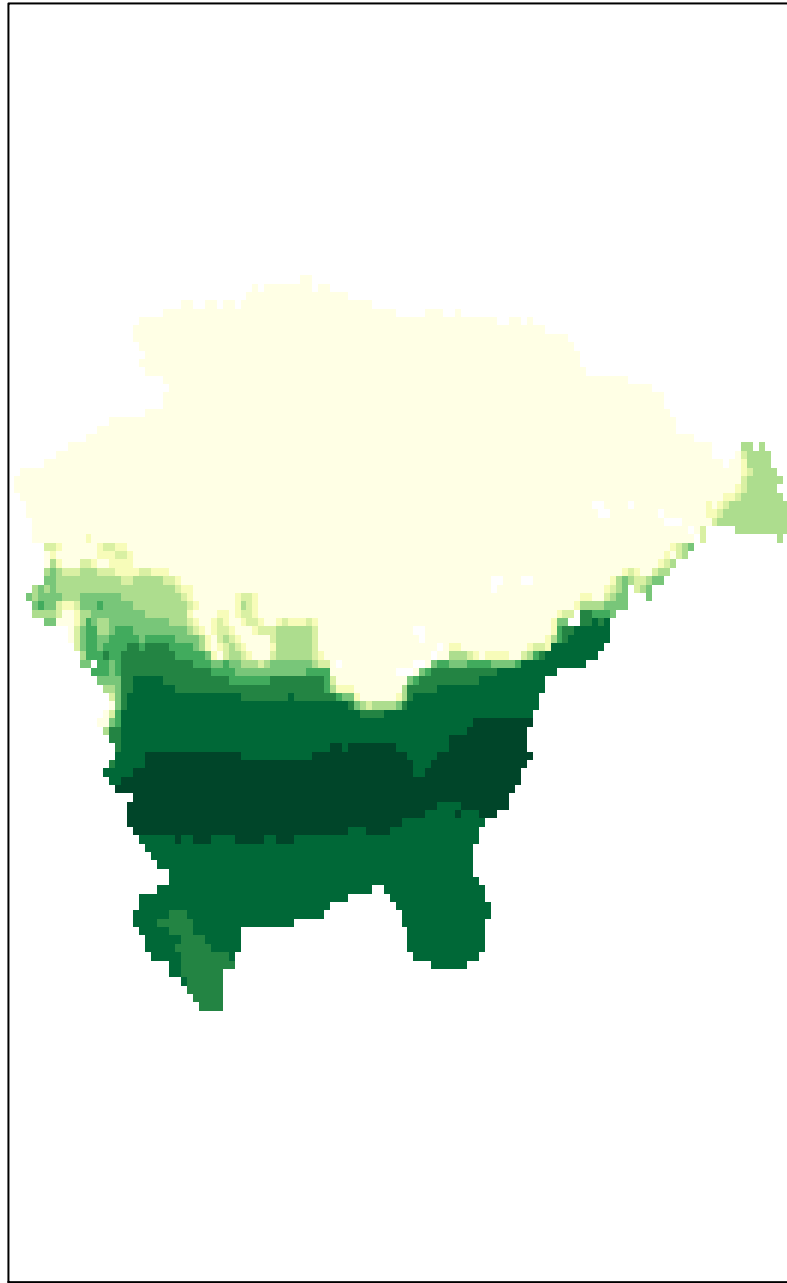


SUM, X17000.ybp

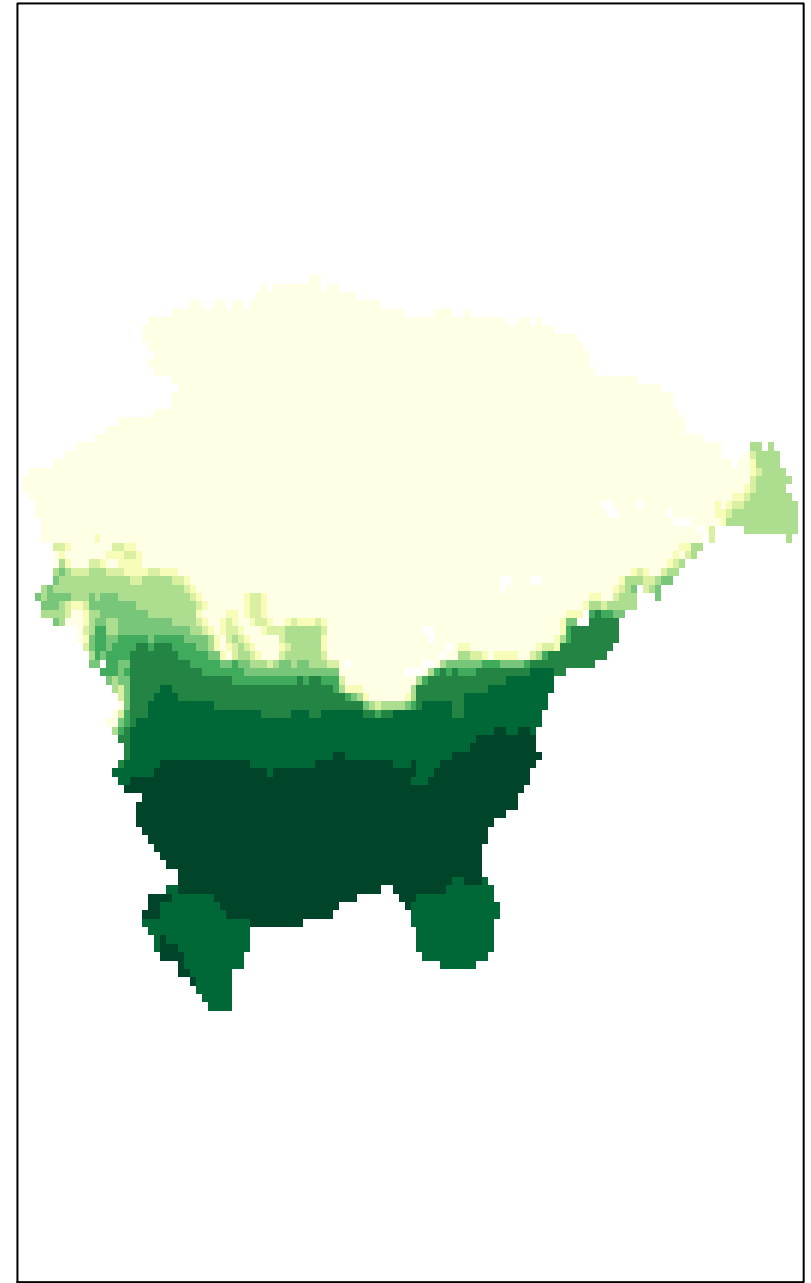


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X16000.ybp

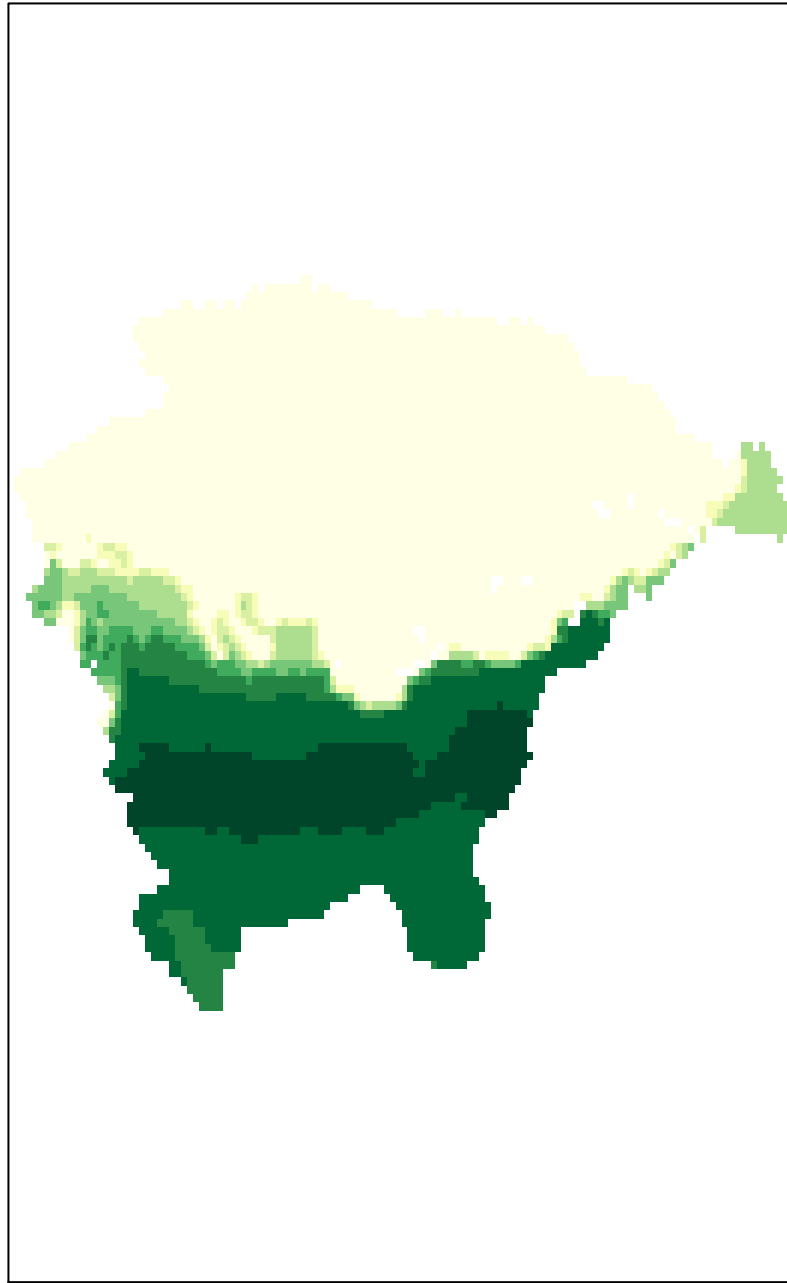


SUM, X16000.ybp

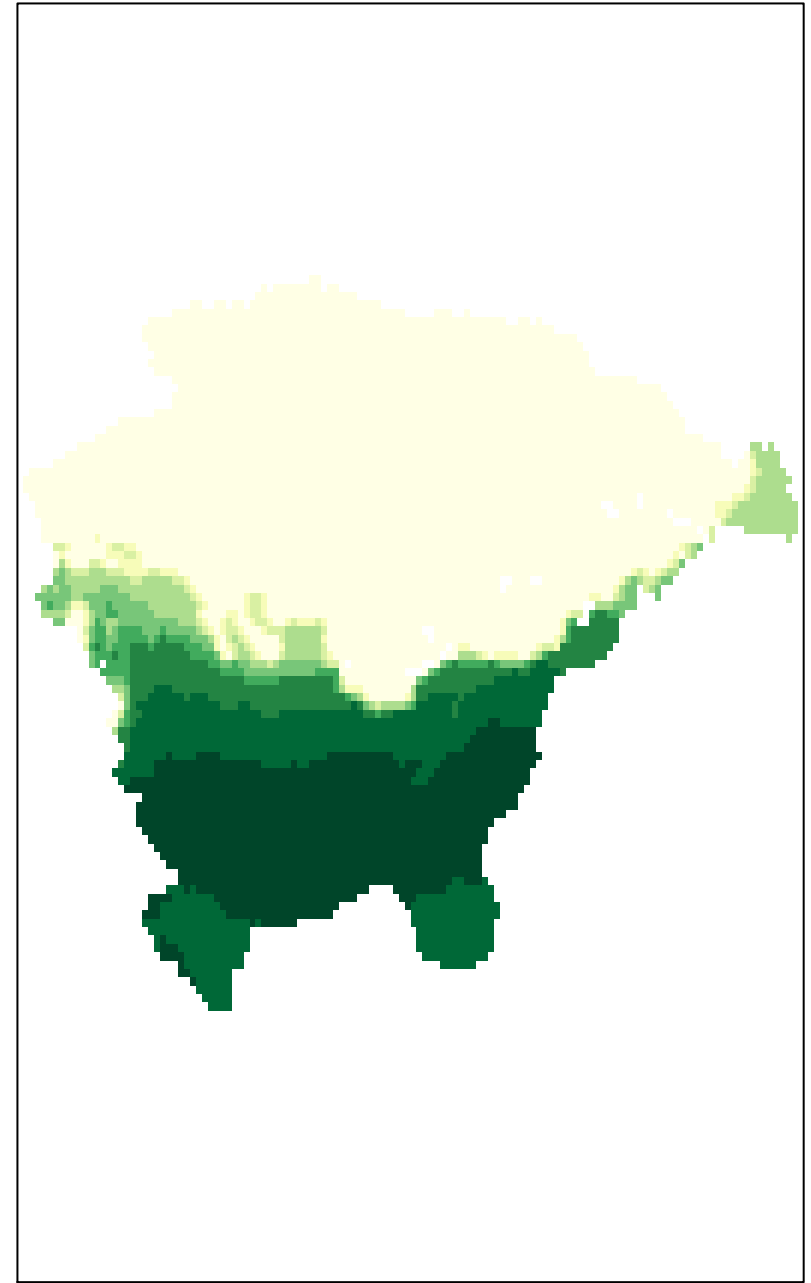


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X15000.ybp

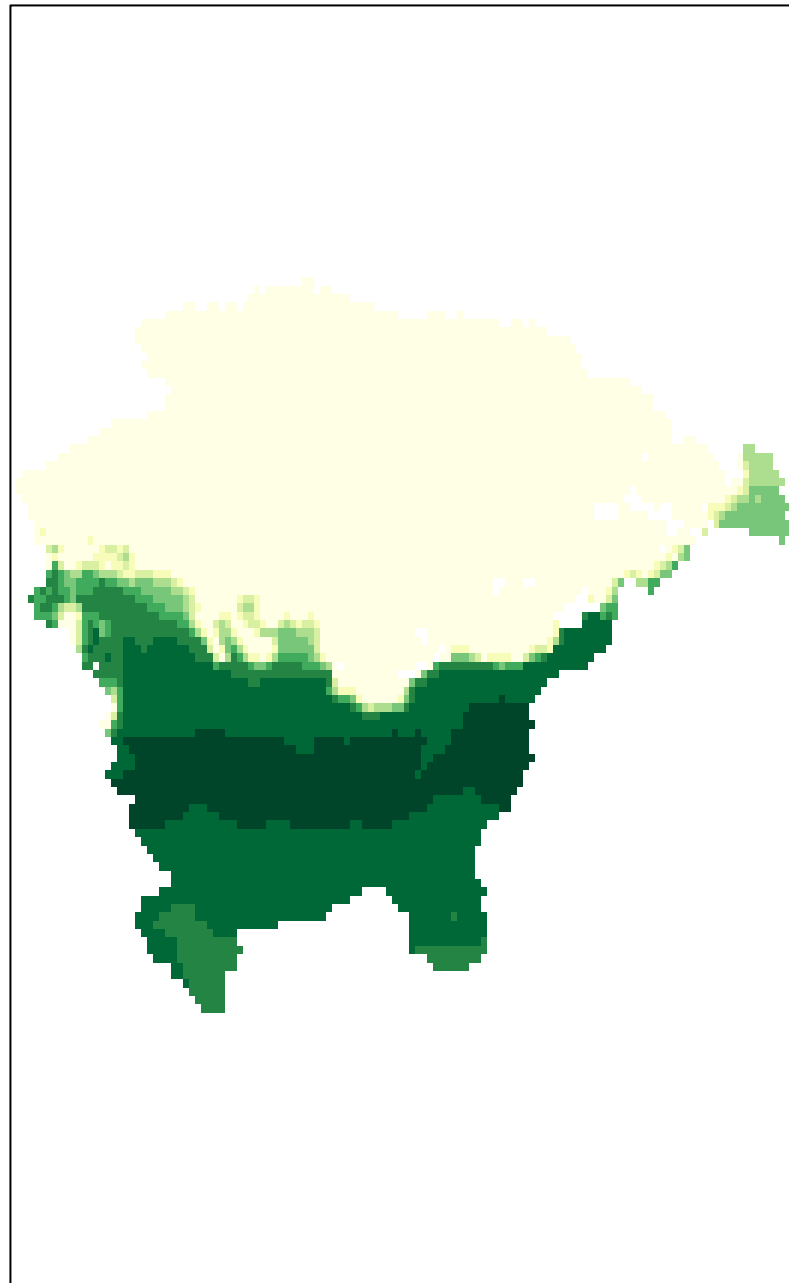


SUM, X15000.ybp

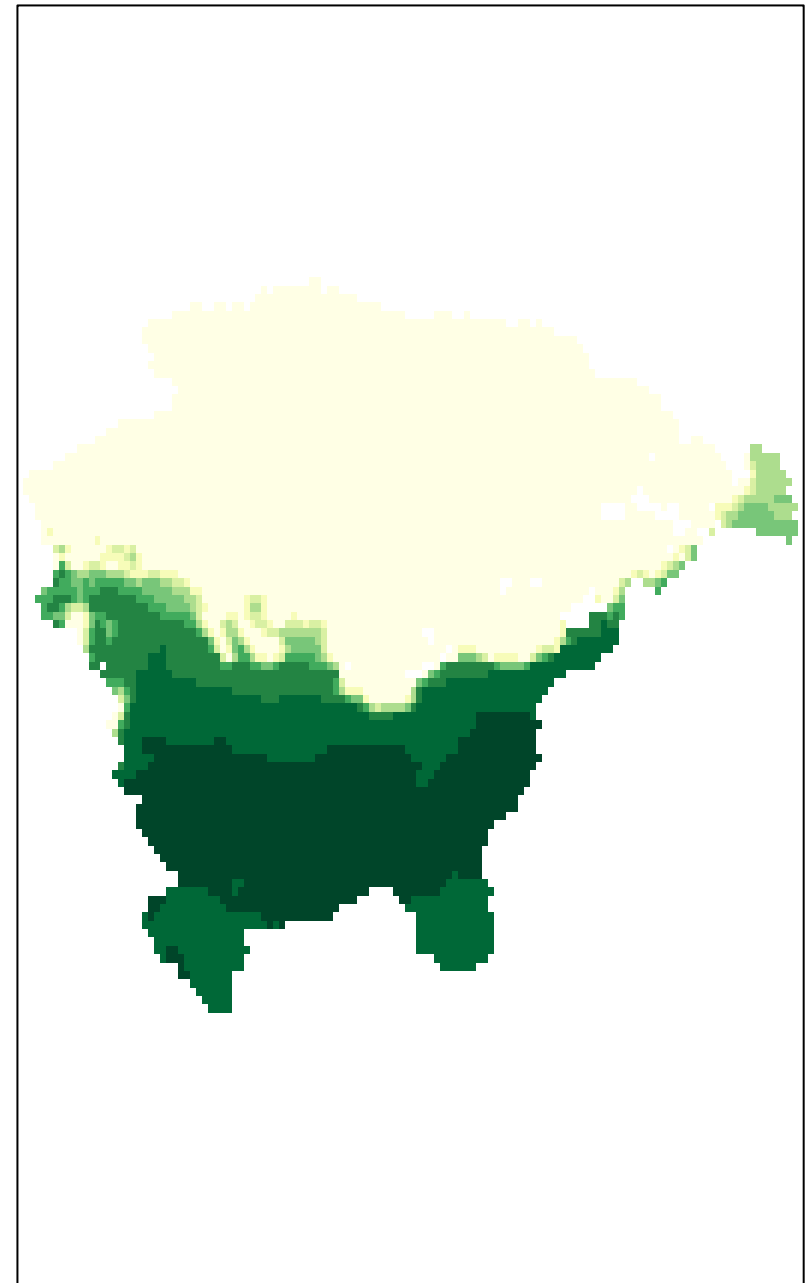


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X14000.ybp



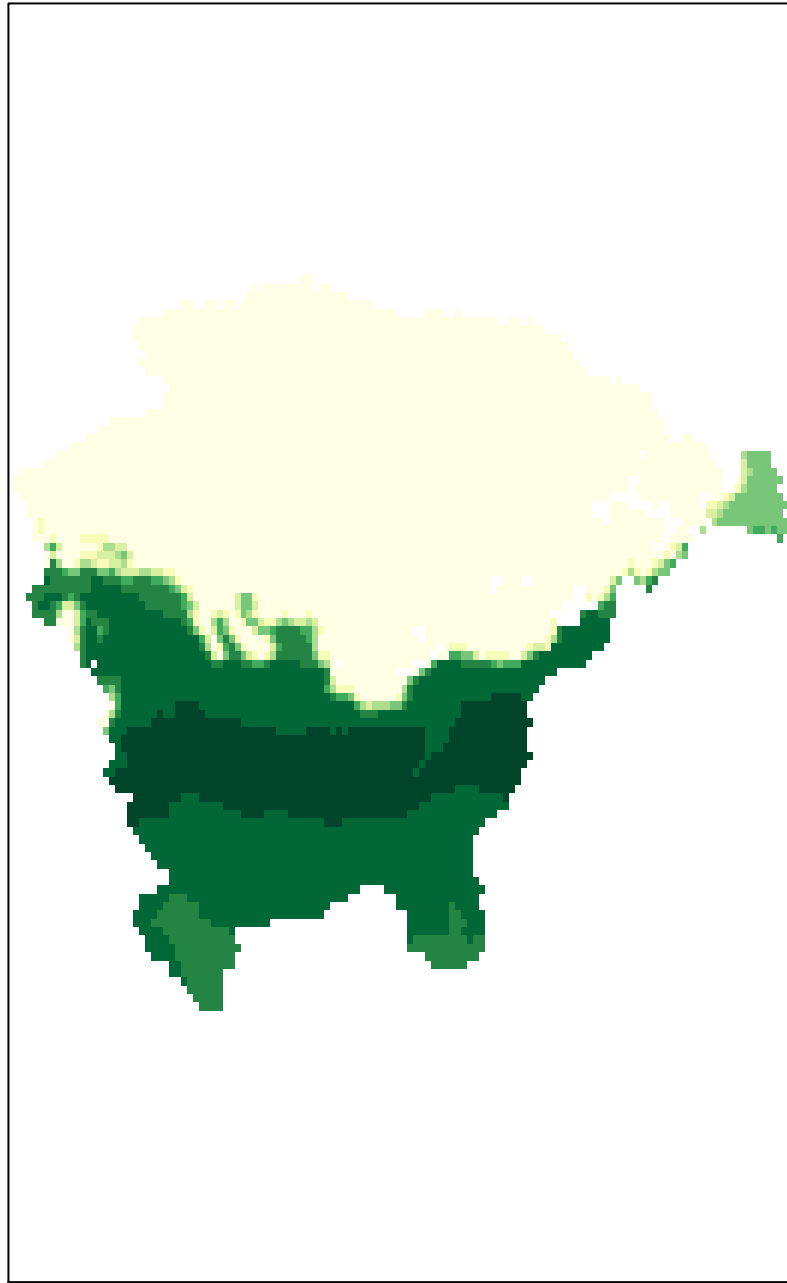
SUM, X14000.ybp



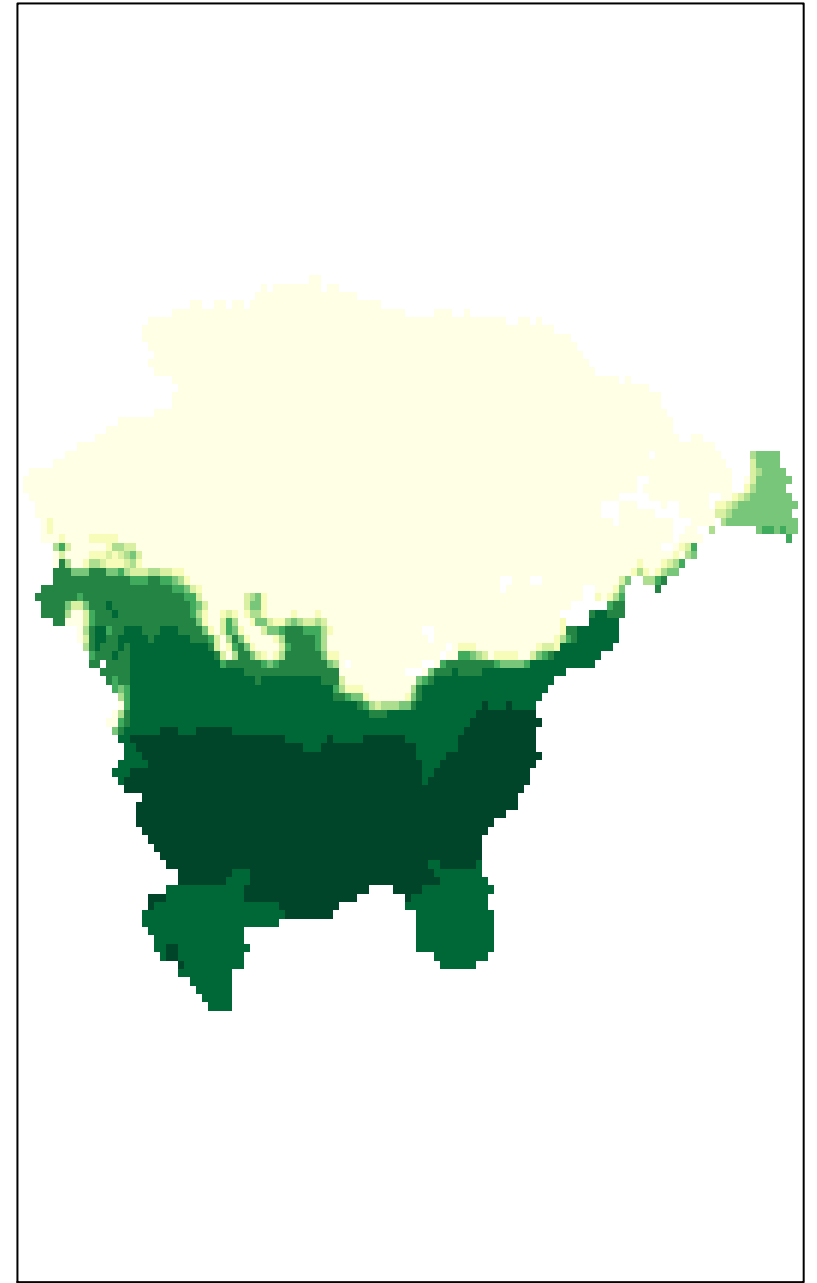
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SUM, X13000.ybp

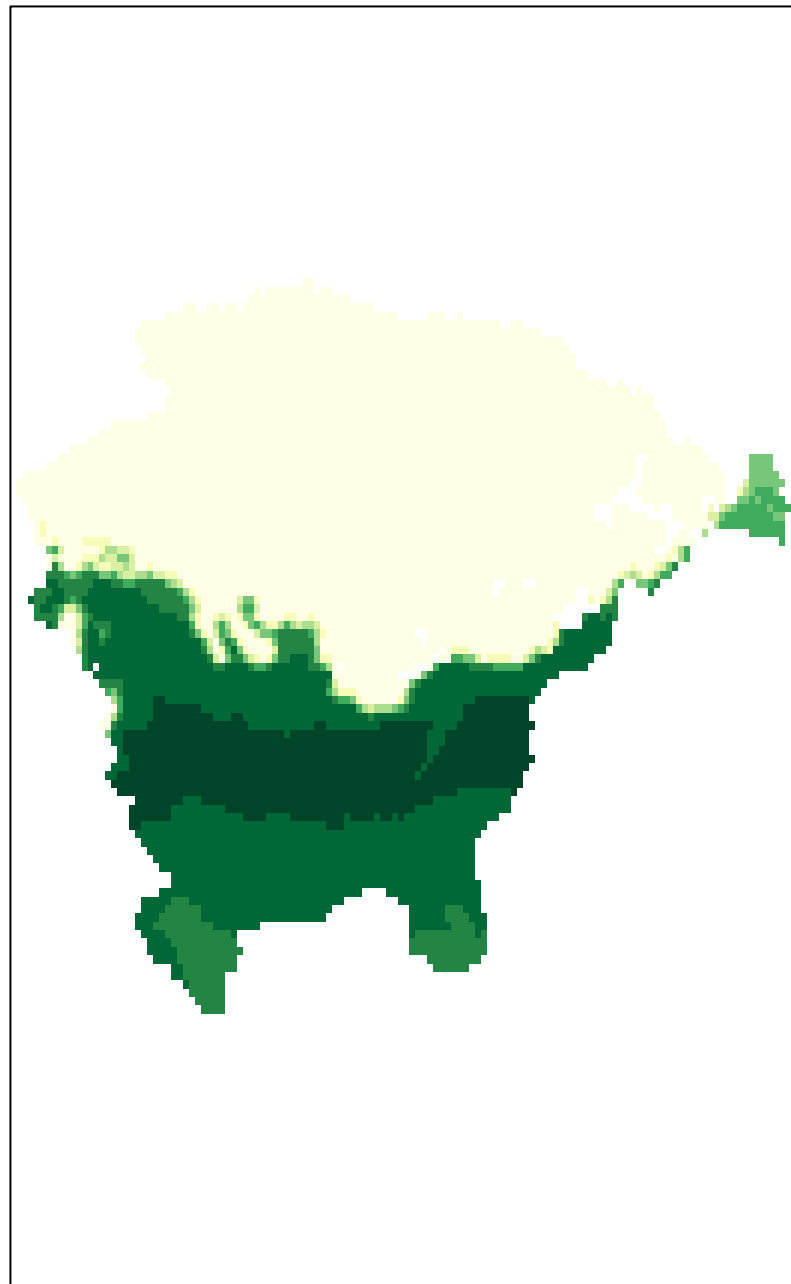


SUM, X13000.ybp

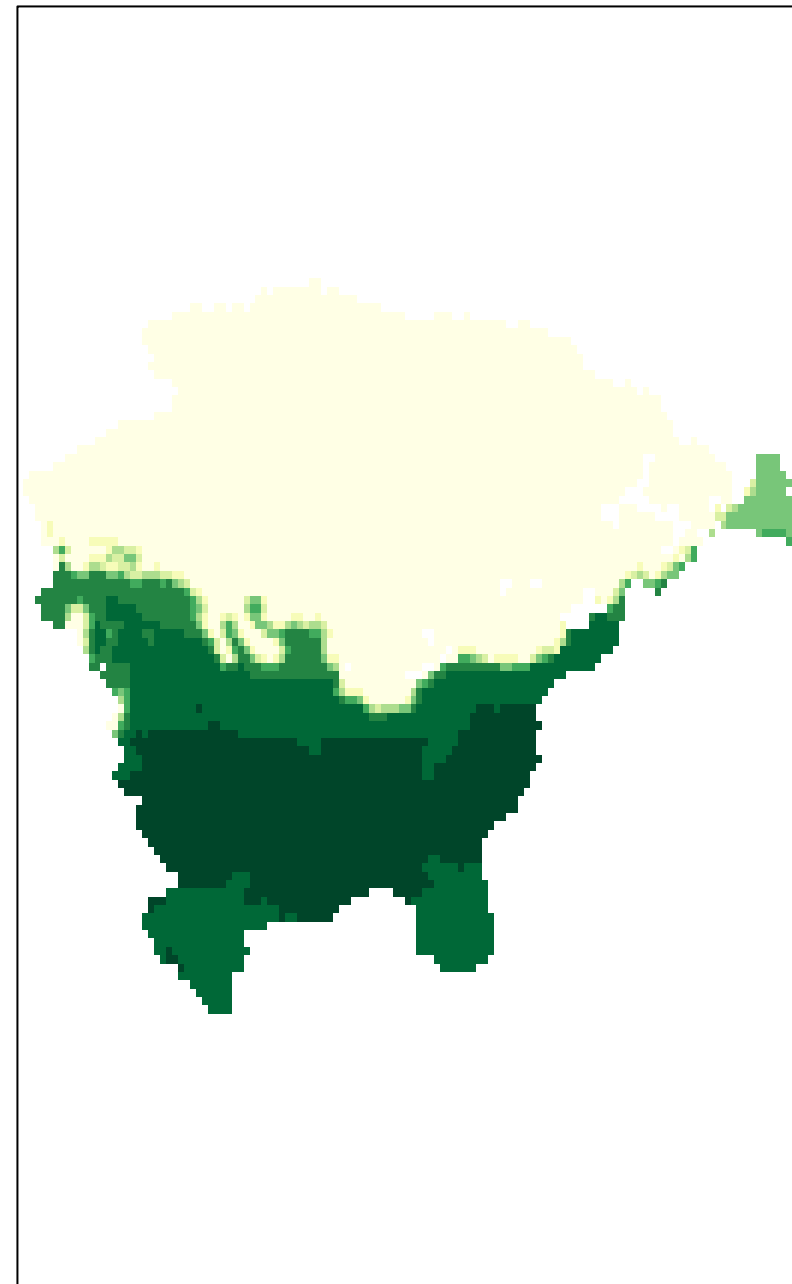


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X12000.ybp

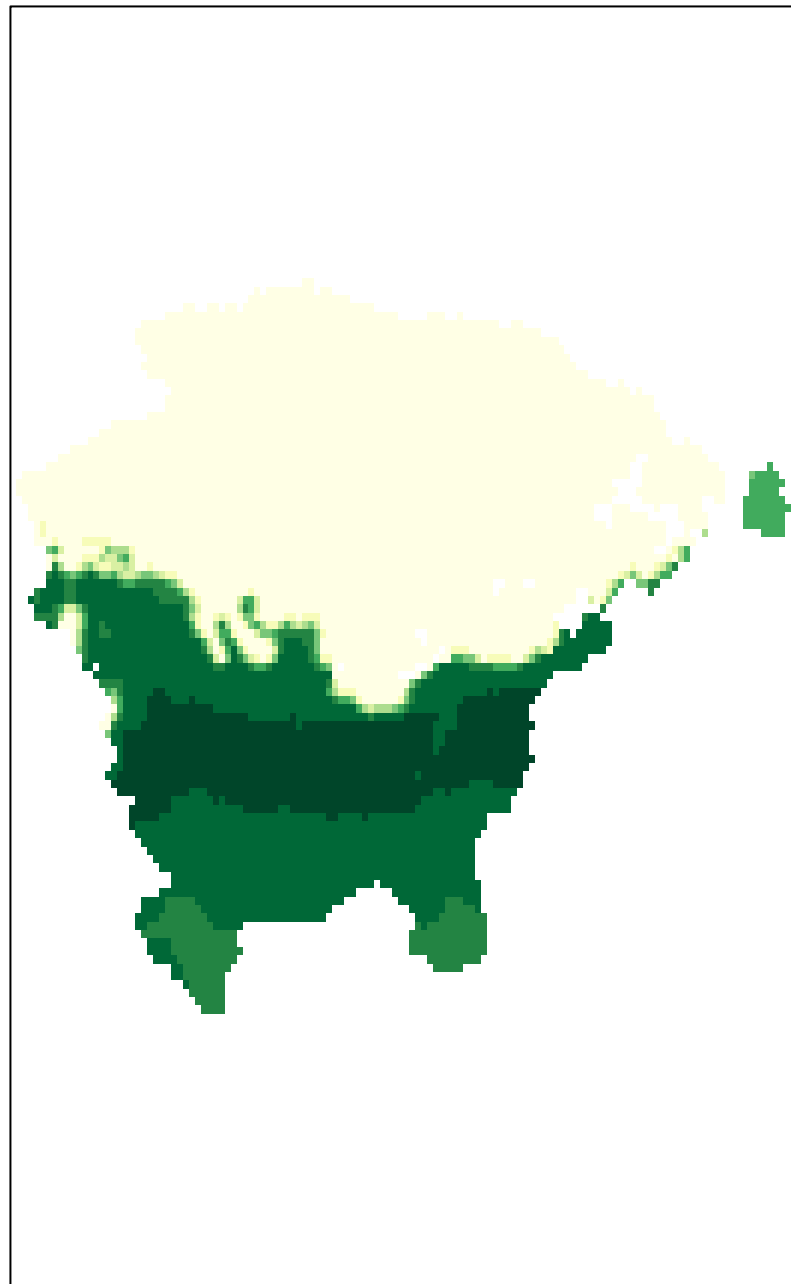


SUM, X12000.ybp

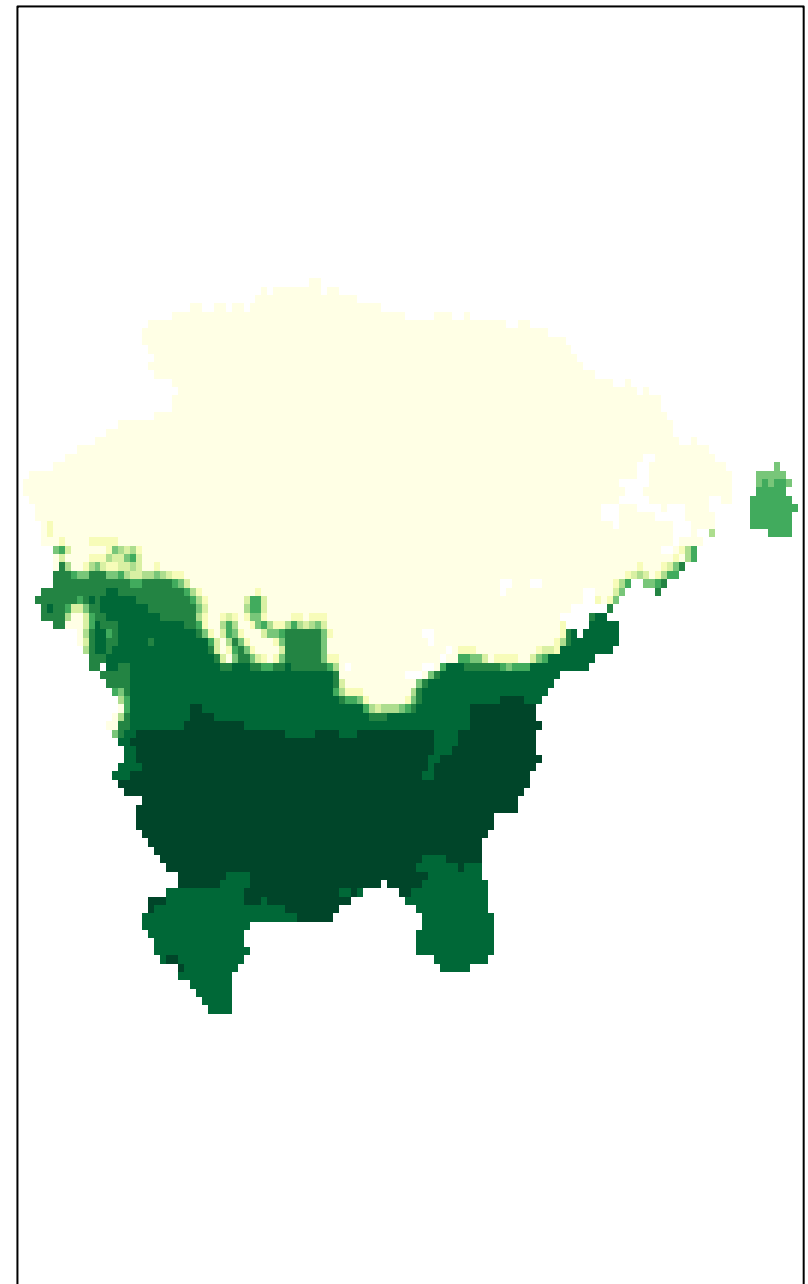


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X11000.ybp

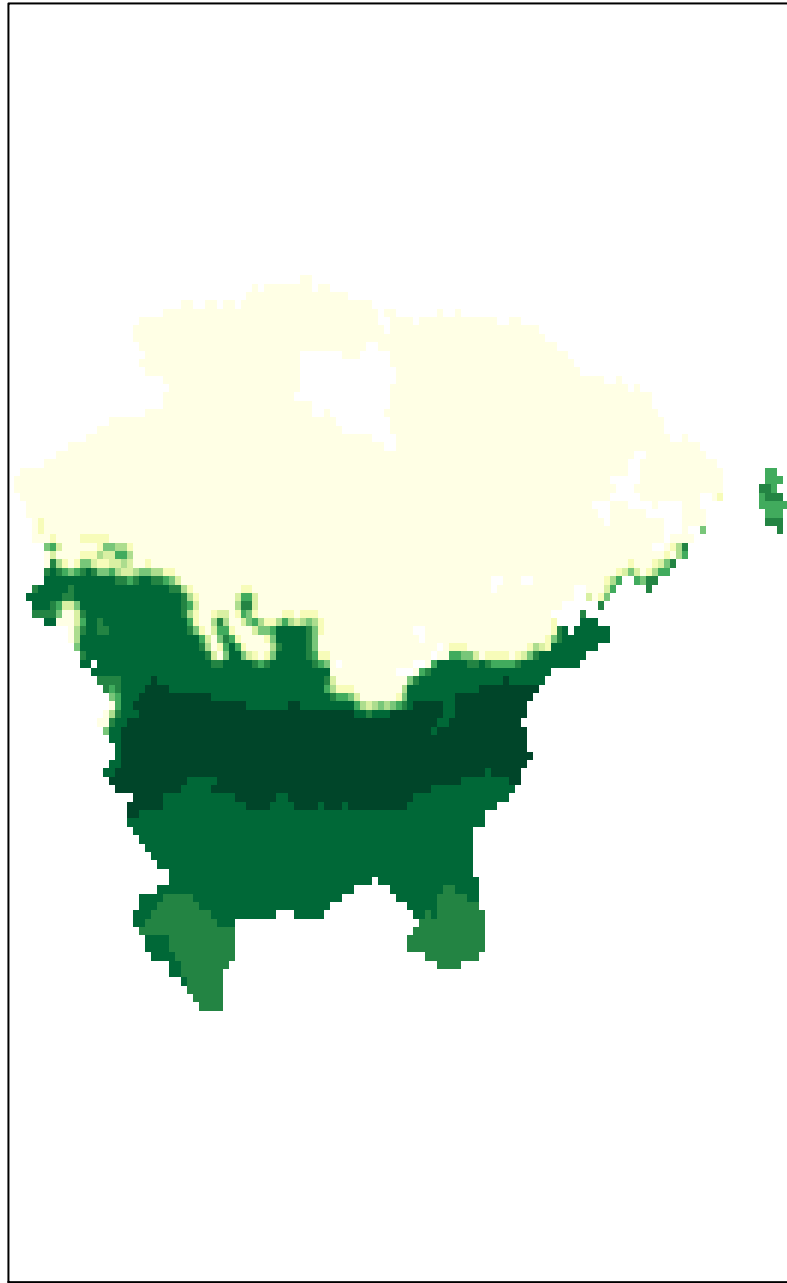


SUM, X11000.ybp

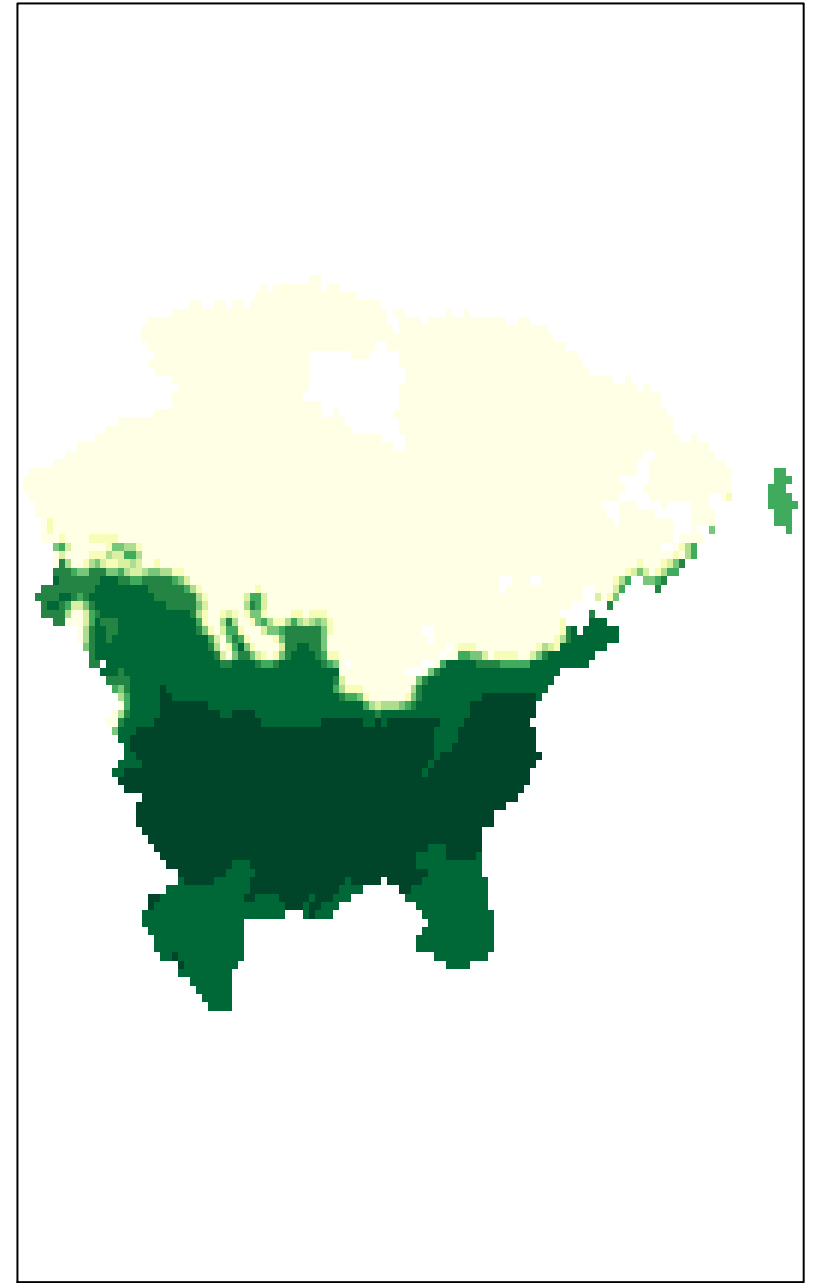


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SUM, X10000.ybp

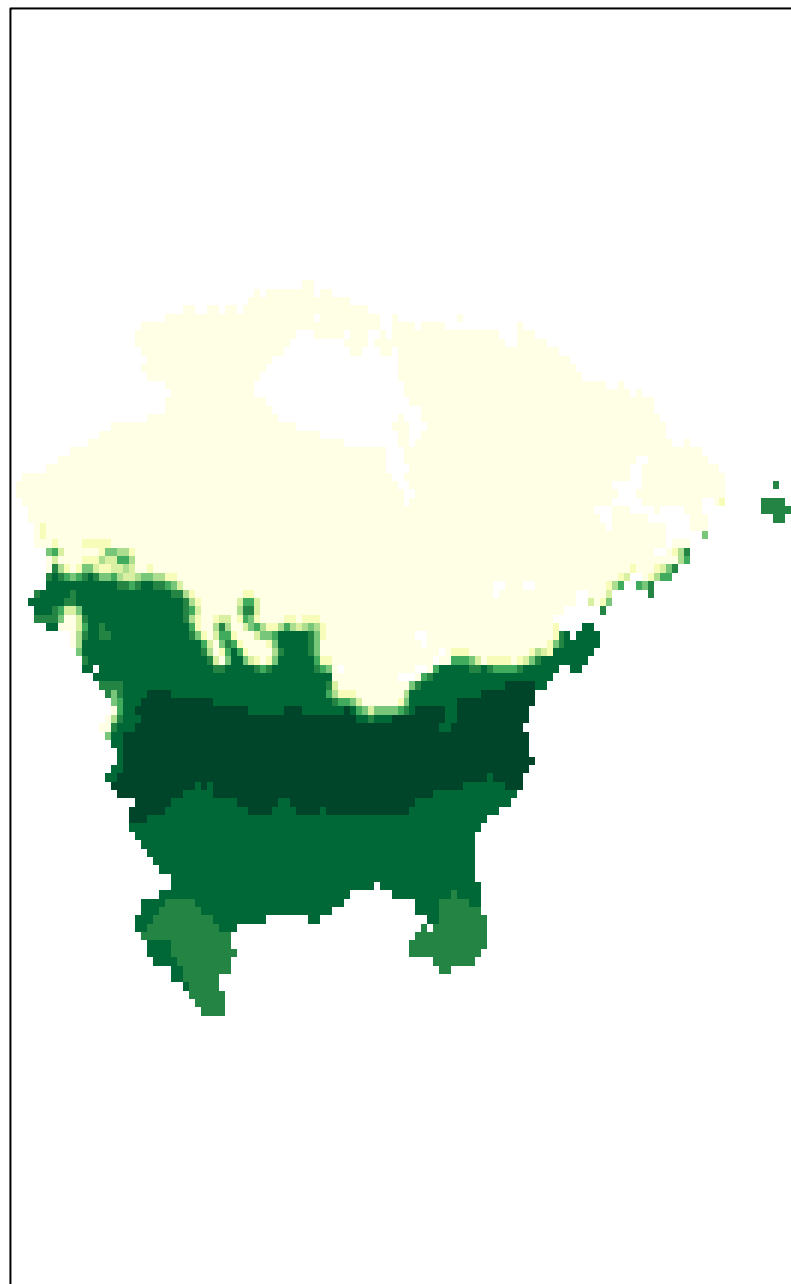


SUM, X10000.ybp

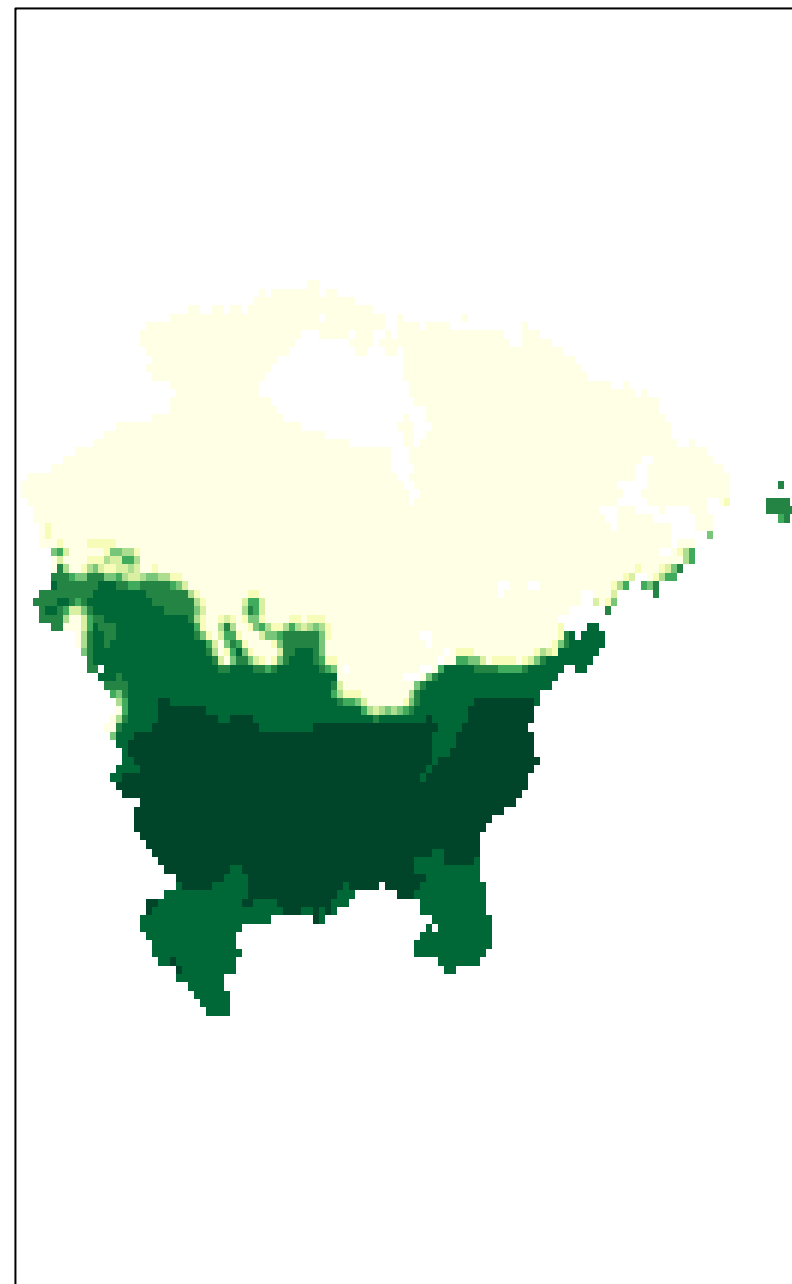


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X9000.ybp

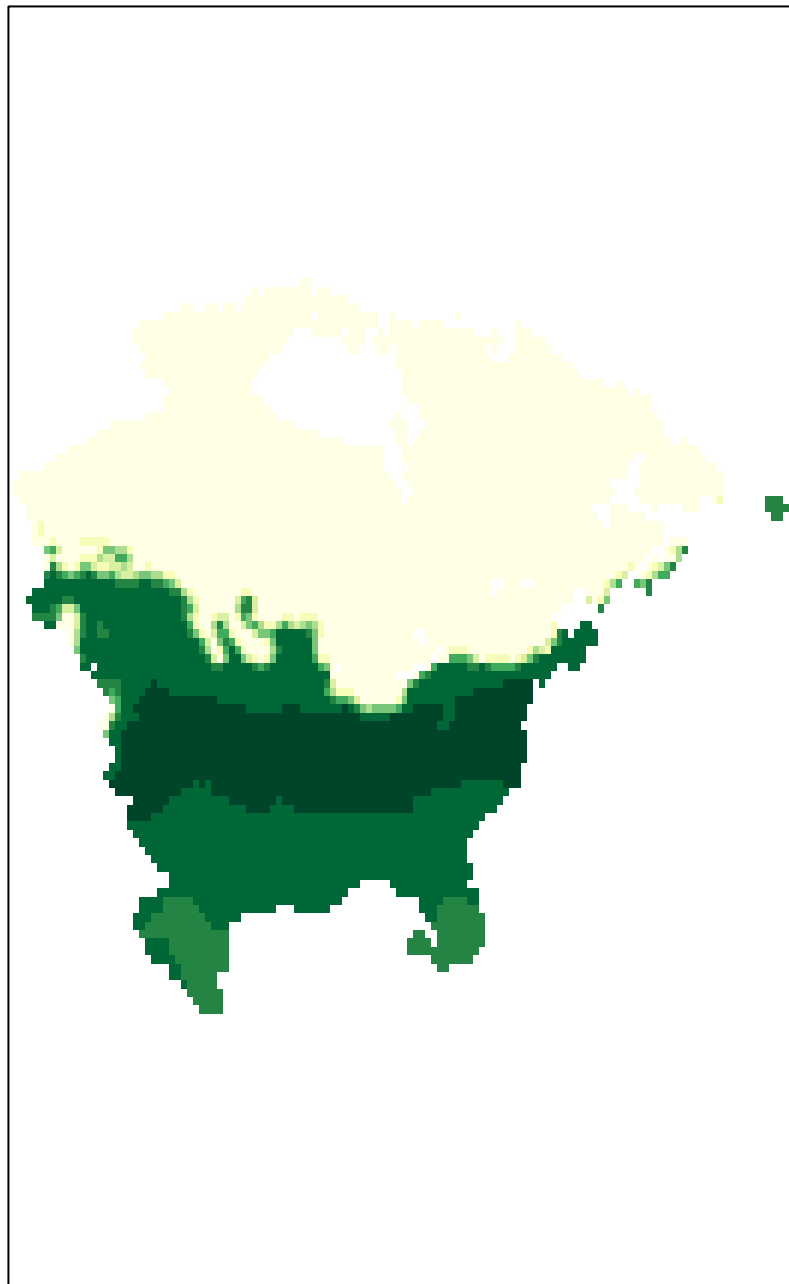


SUM, X9000.ybp

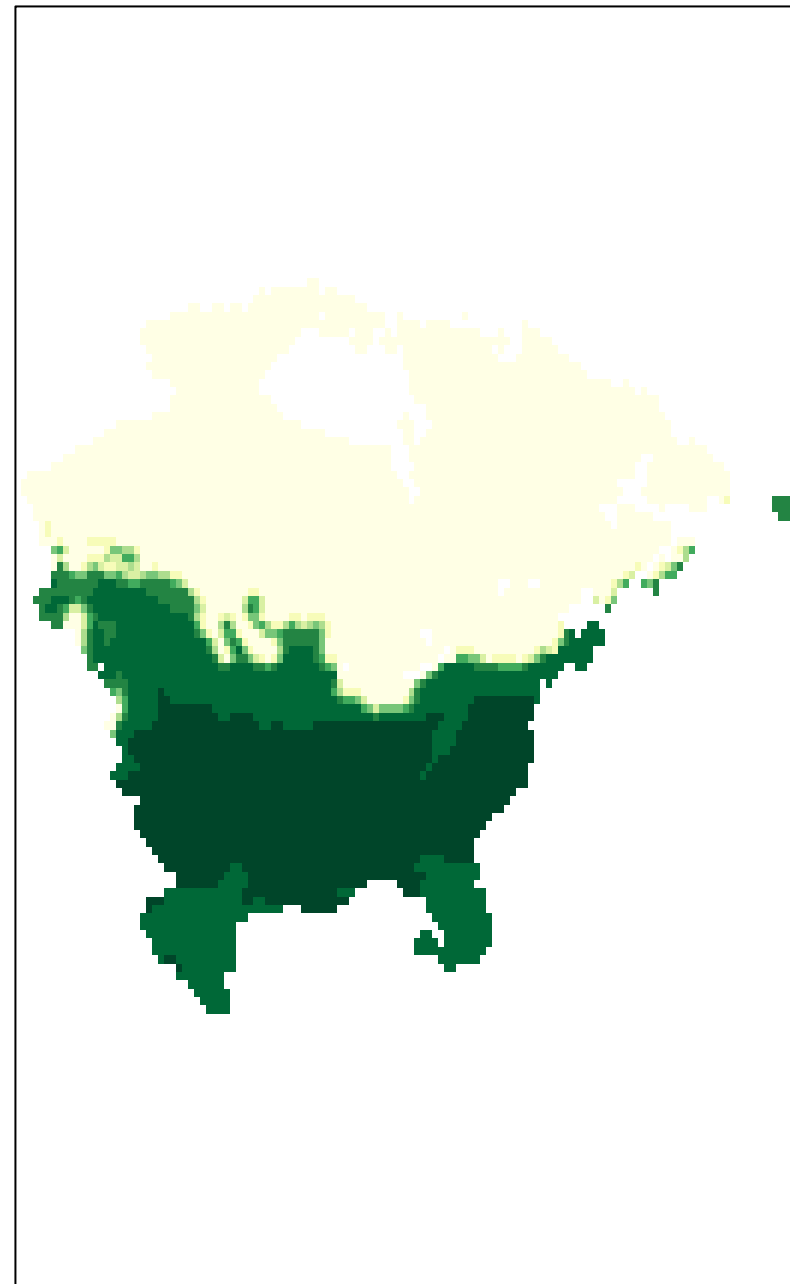


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SUM, X8000.ybp

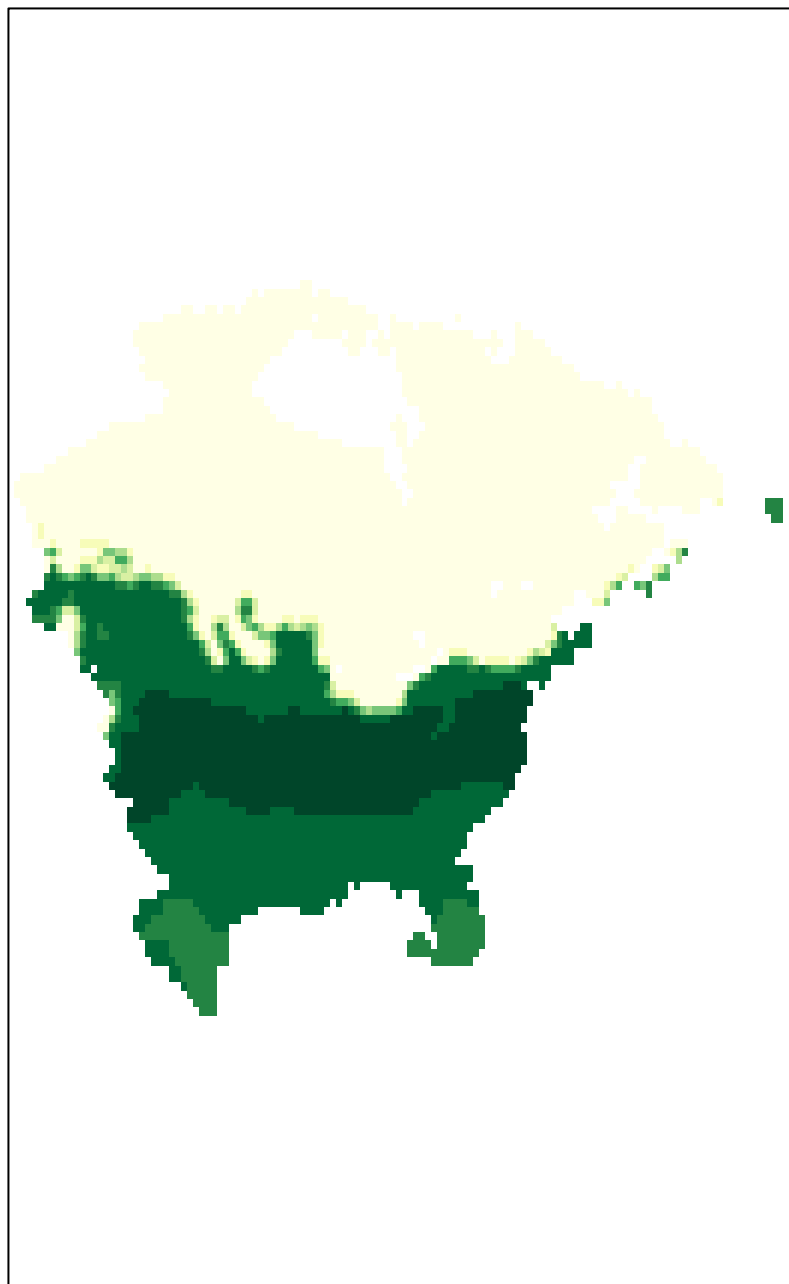


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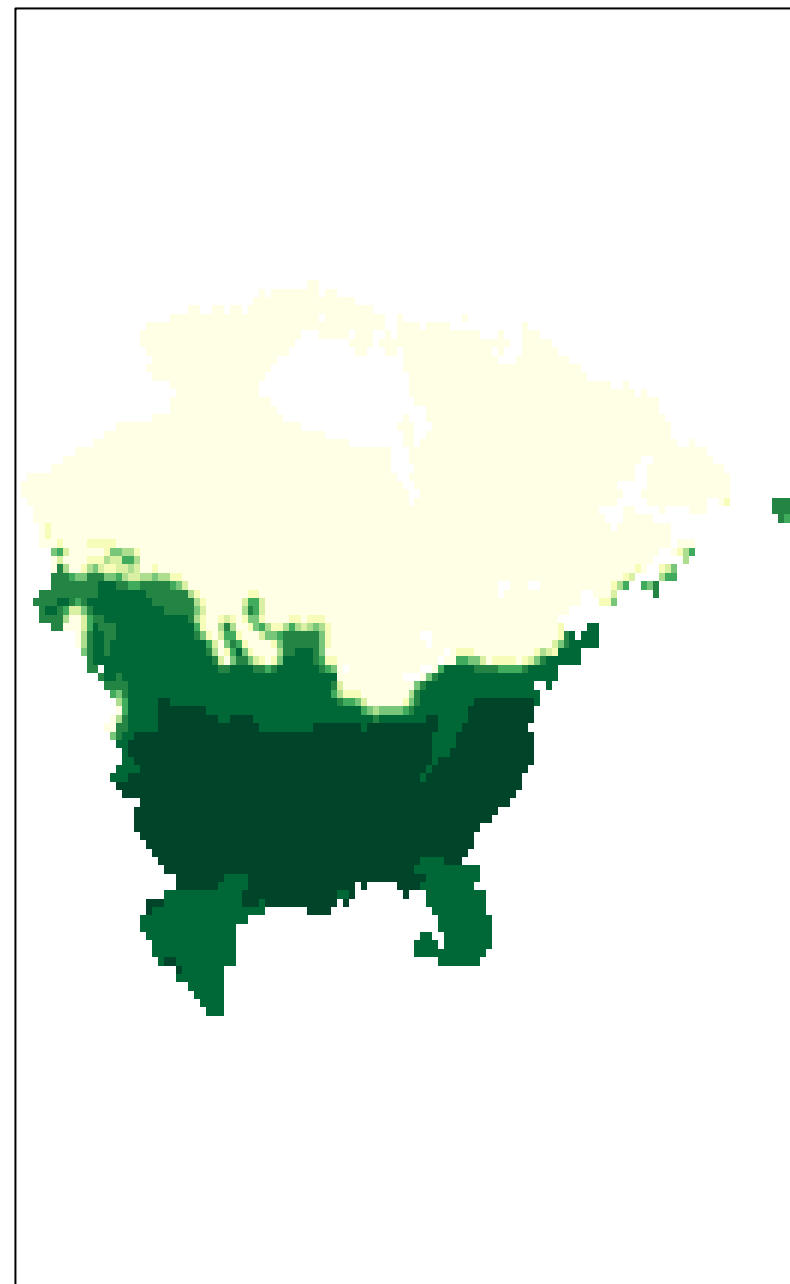


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SUM, X7000.ybp

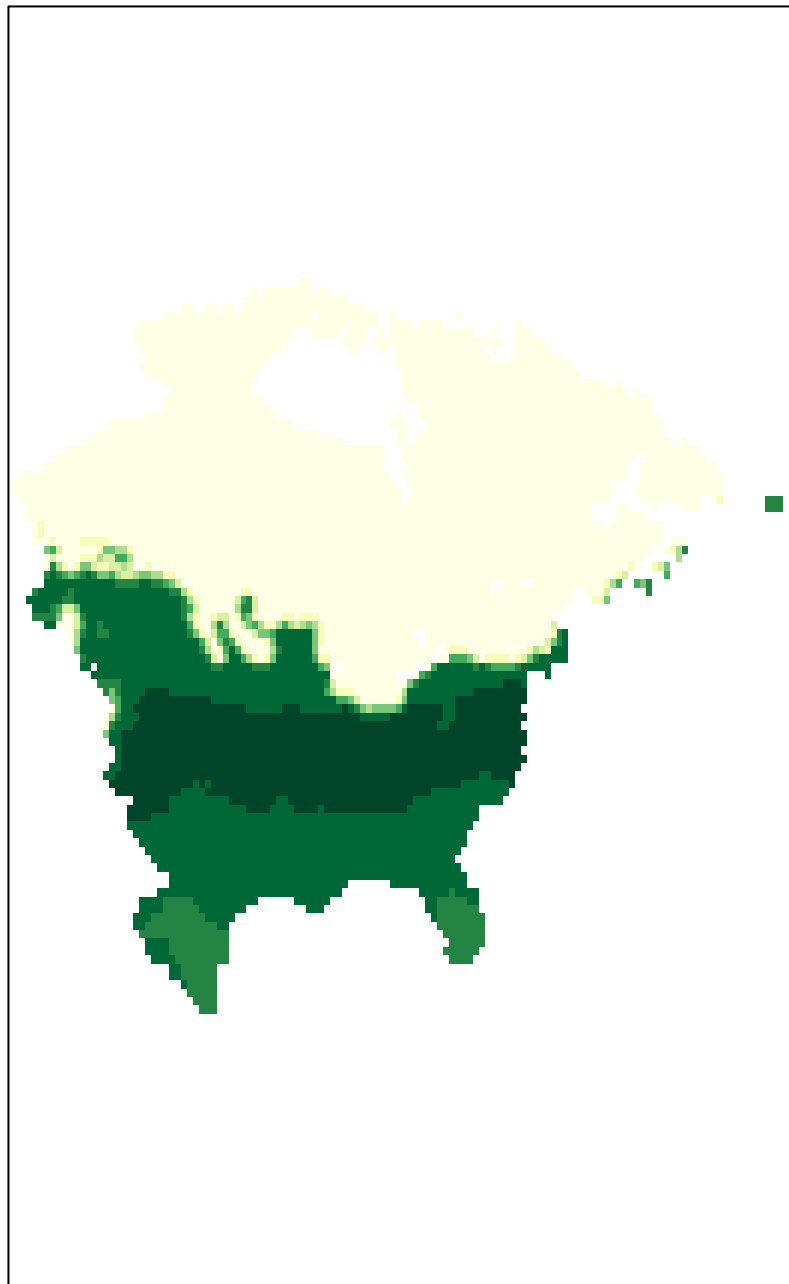


SUM, X7000.ybp

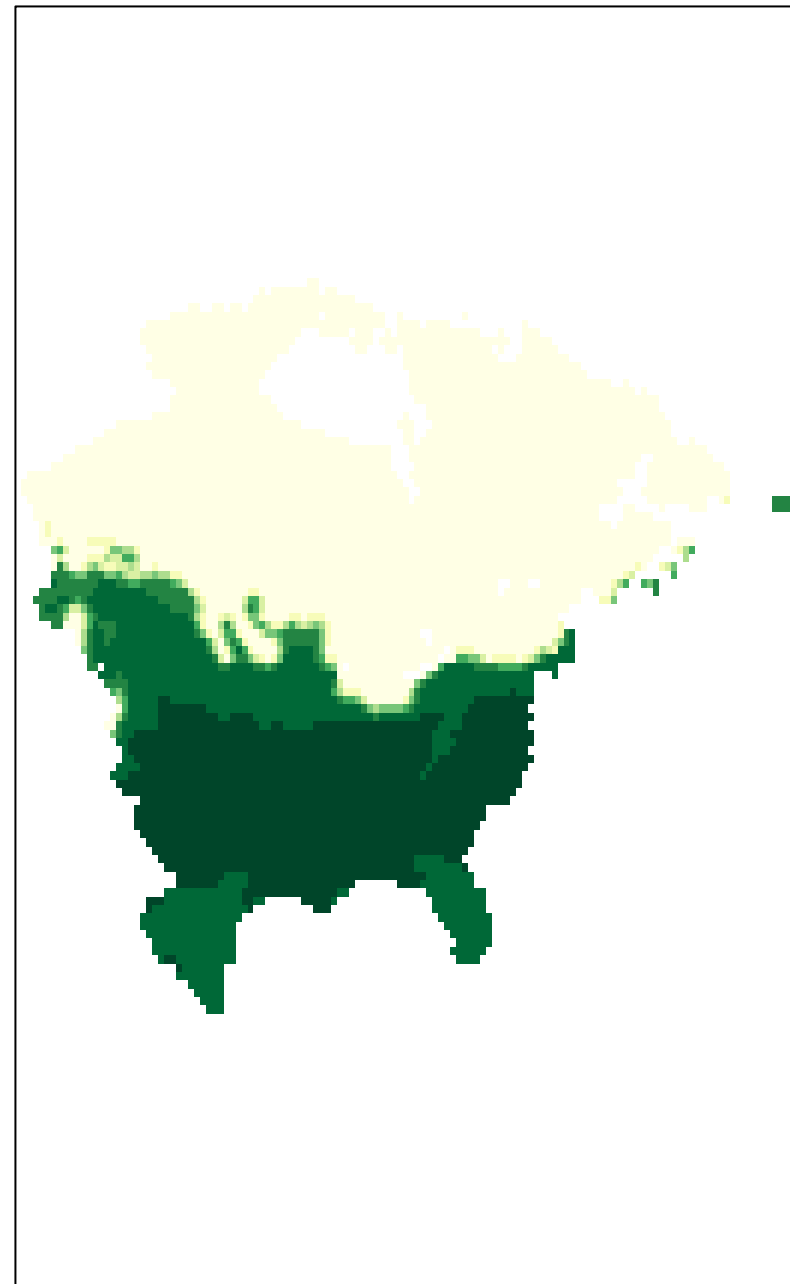


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X6000.ybp



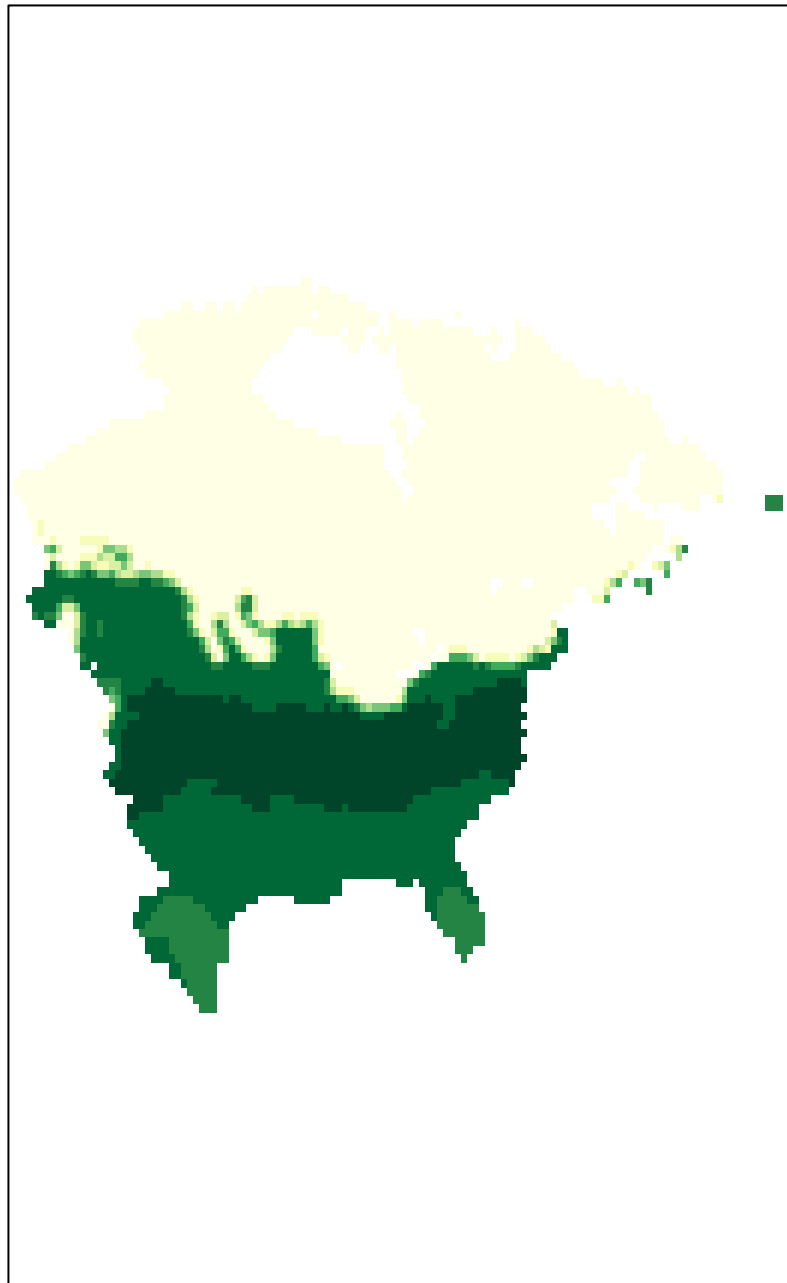
SUM, X6000.ybp



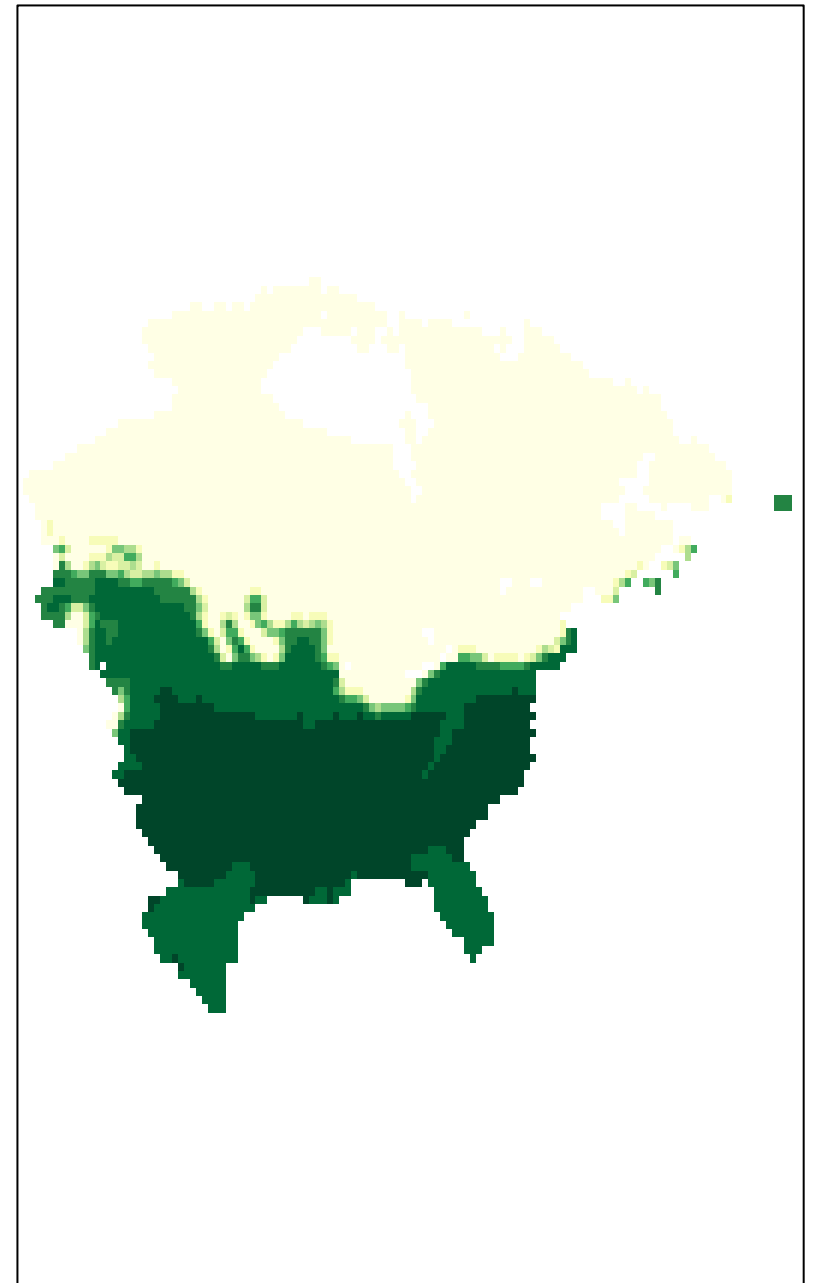
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SUM, X5000.ybp

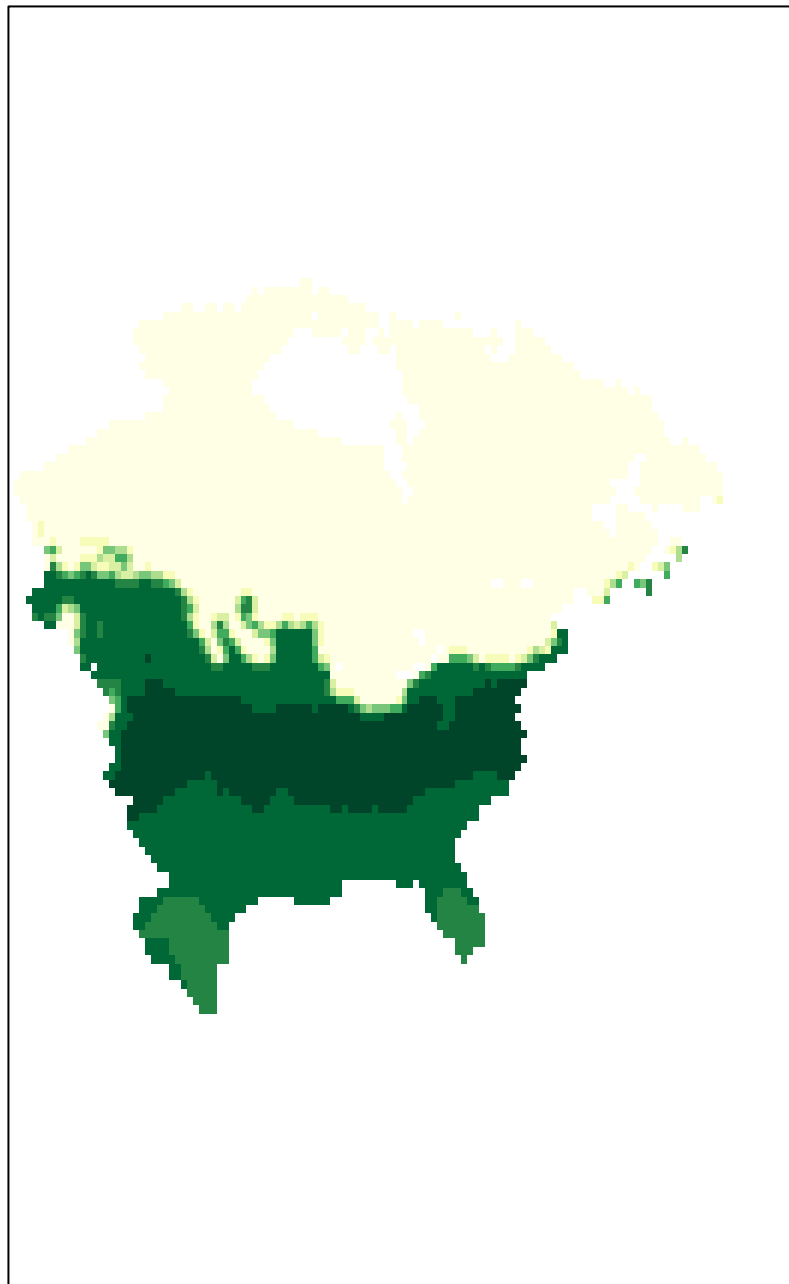


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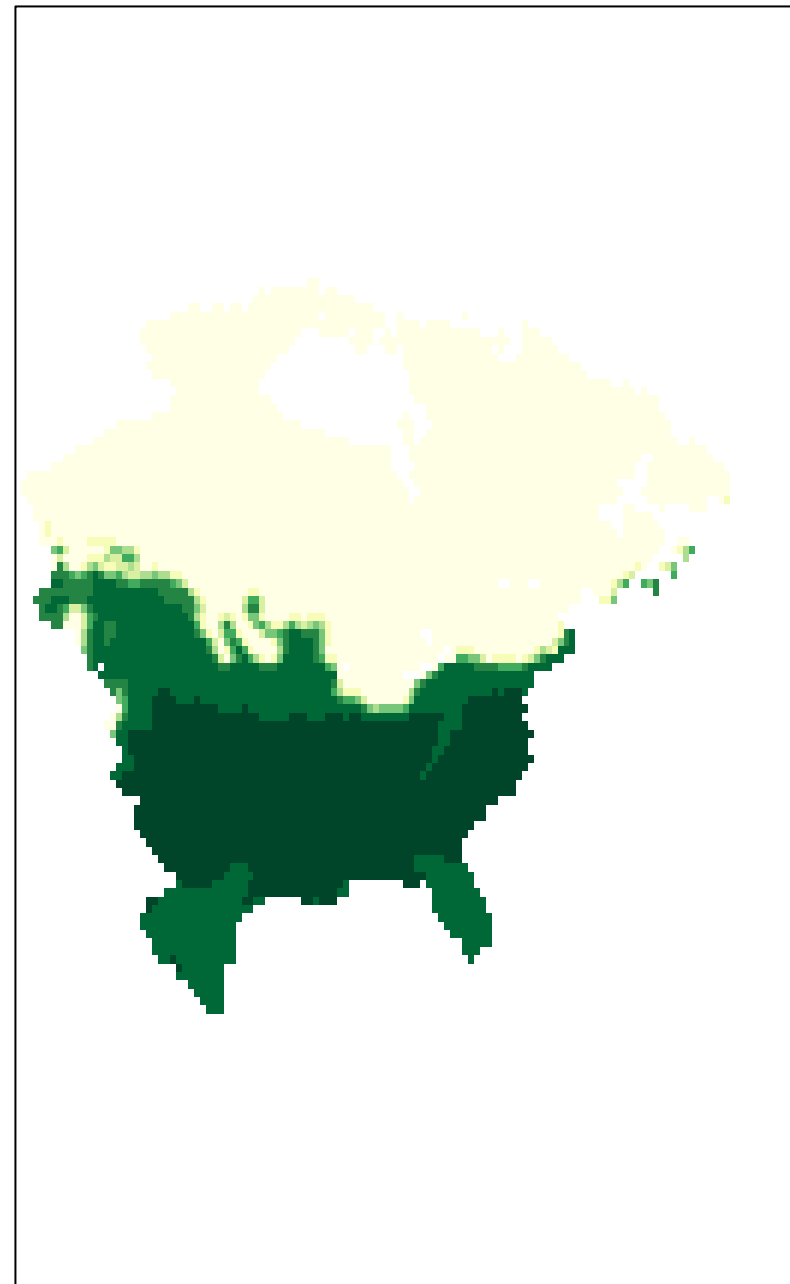


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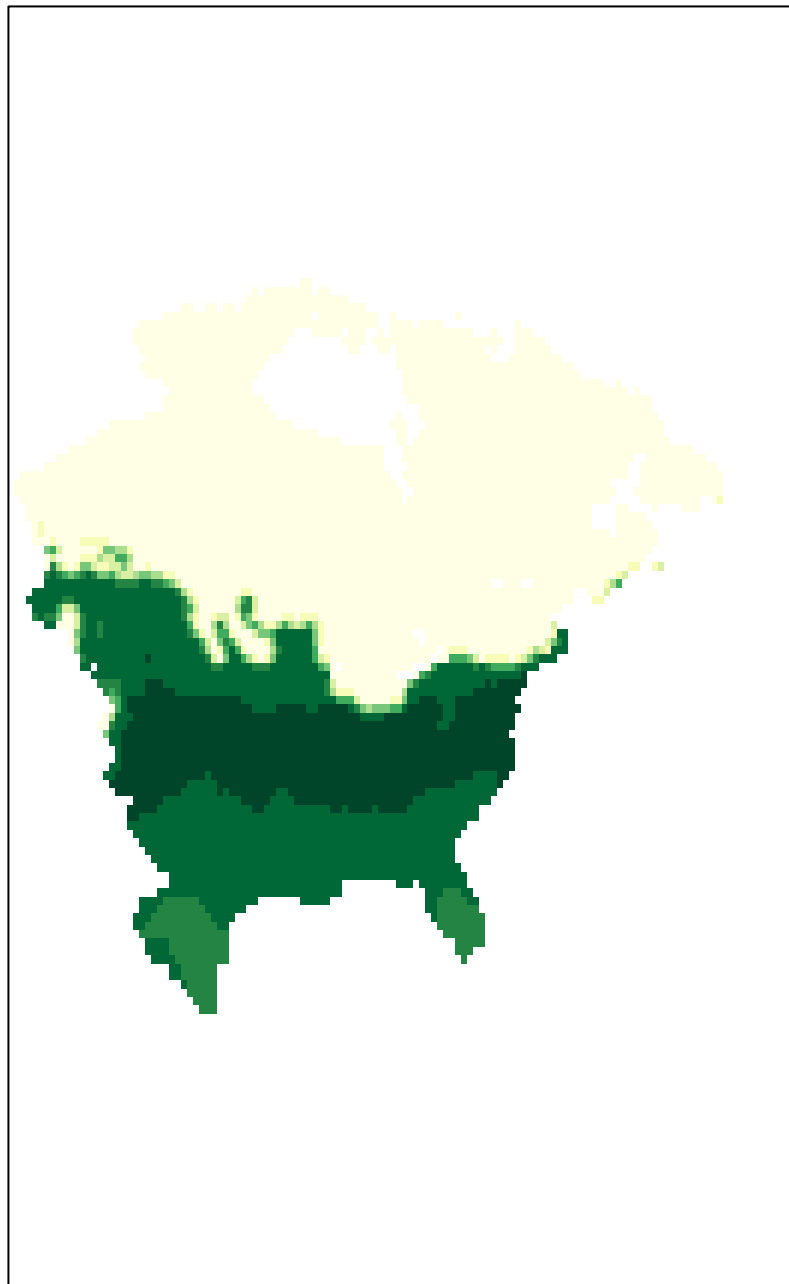


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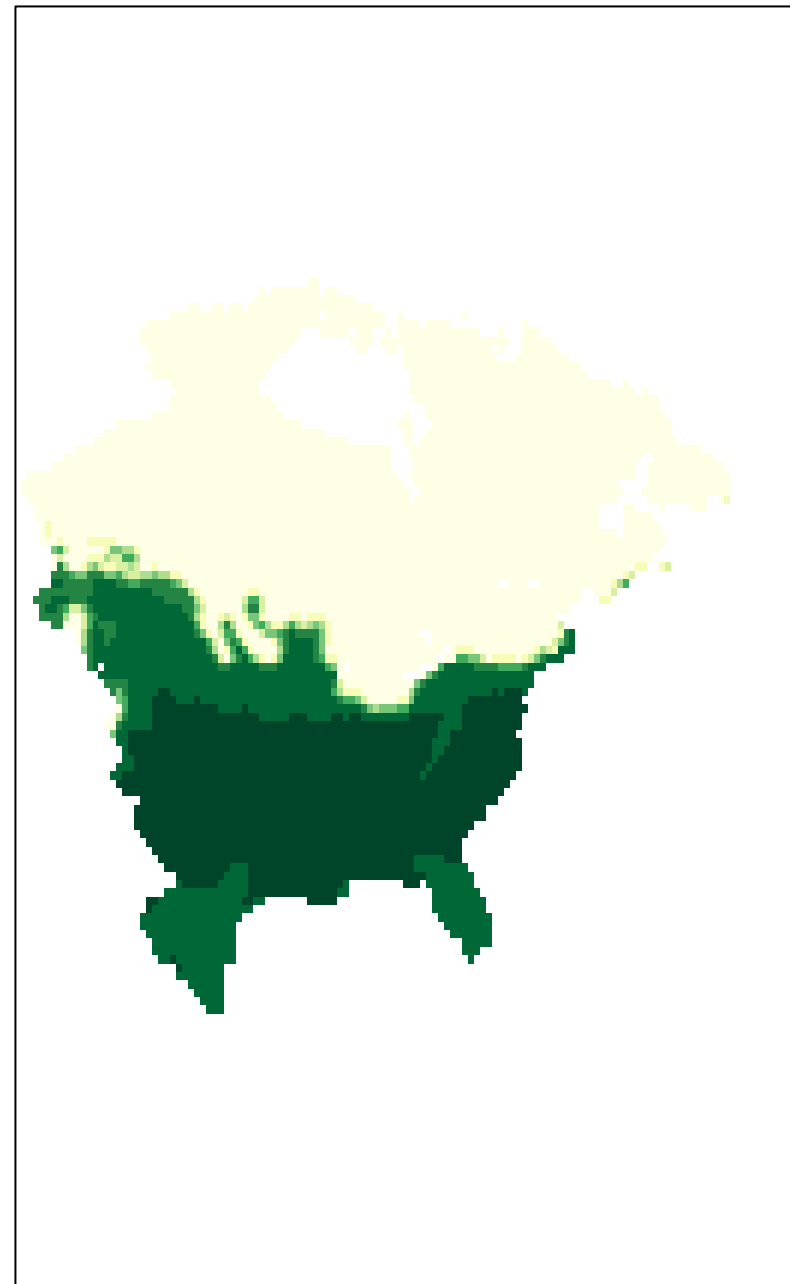


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SUM, X3000.ybp

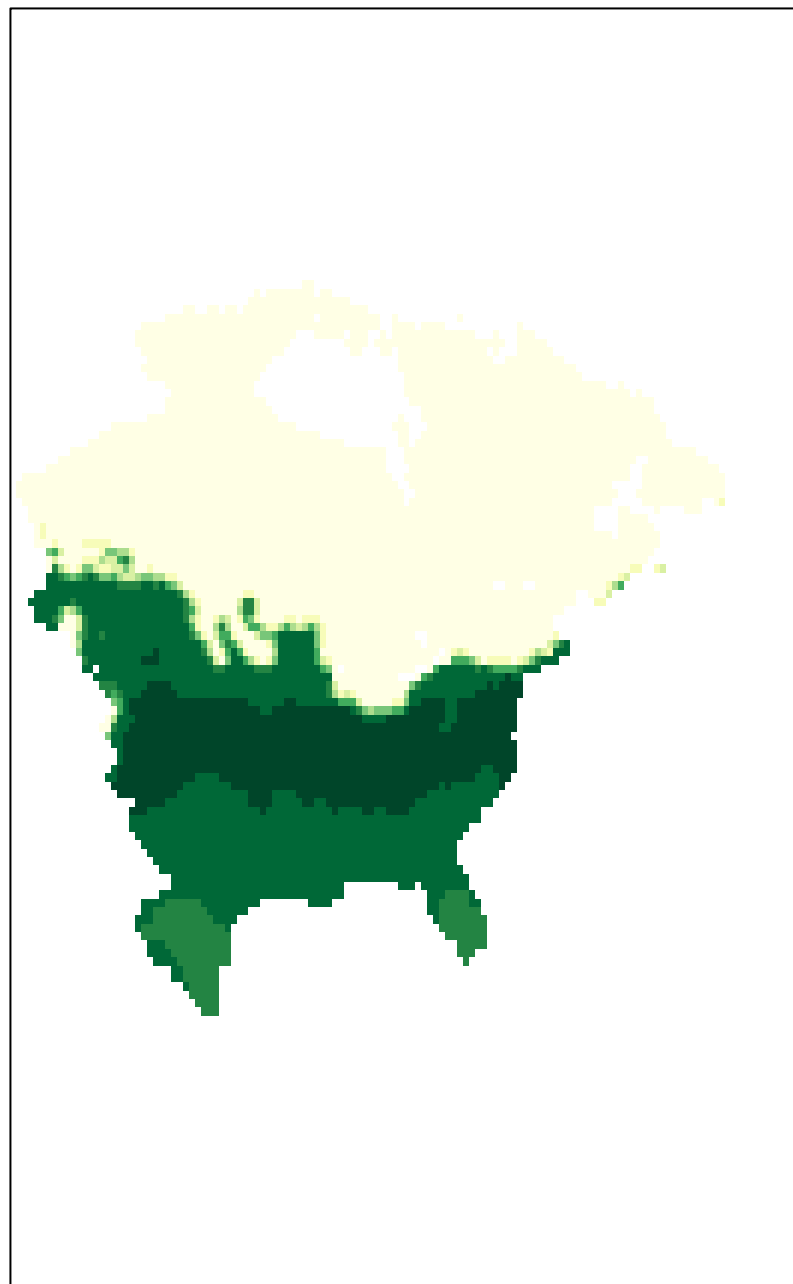


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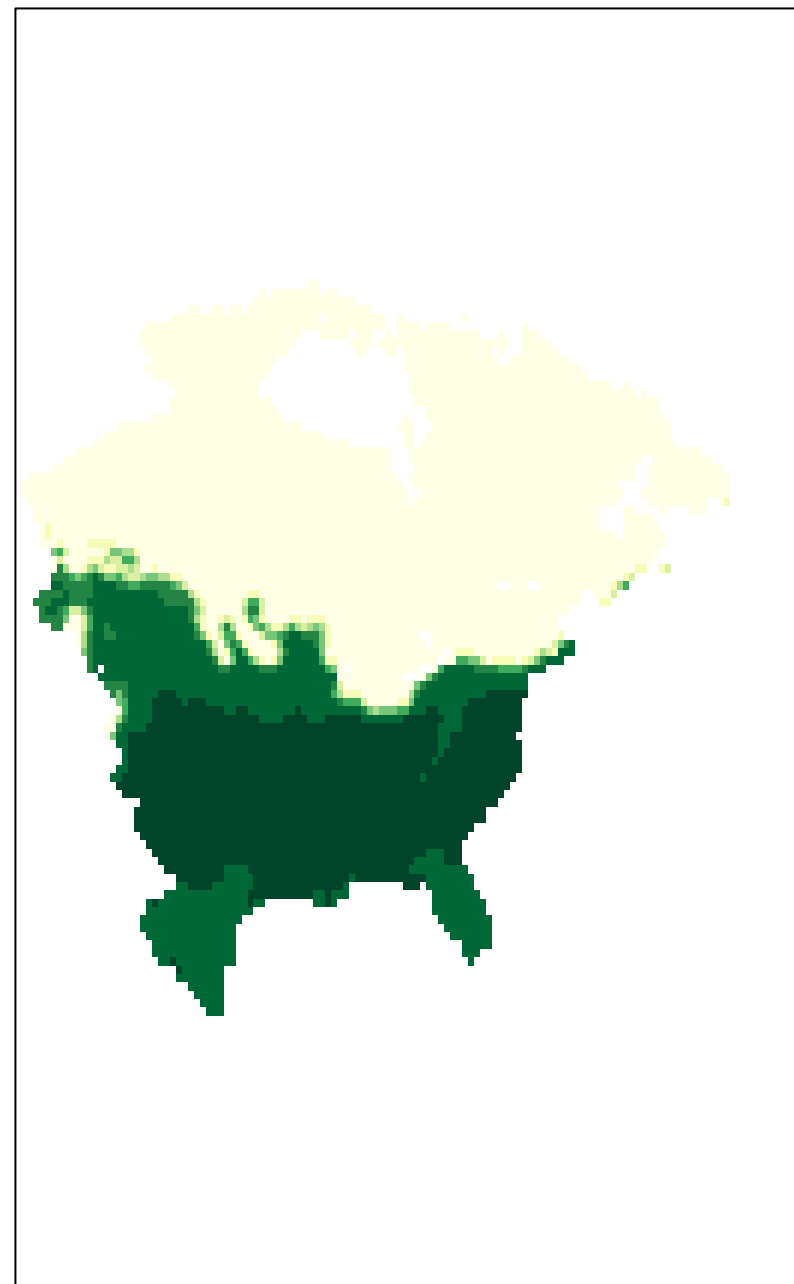


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SUM, X2000.ybp

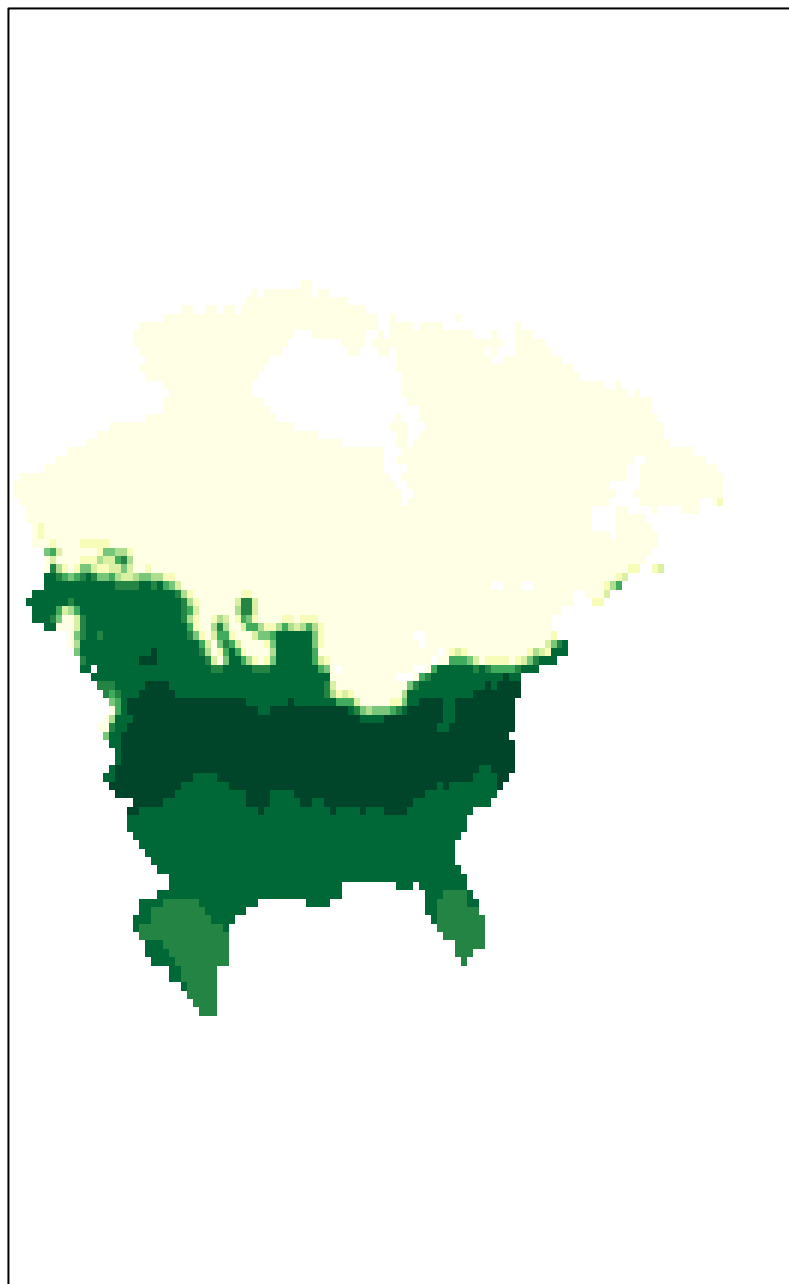


SUM, X2000.ybp

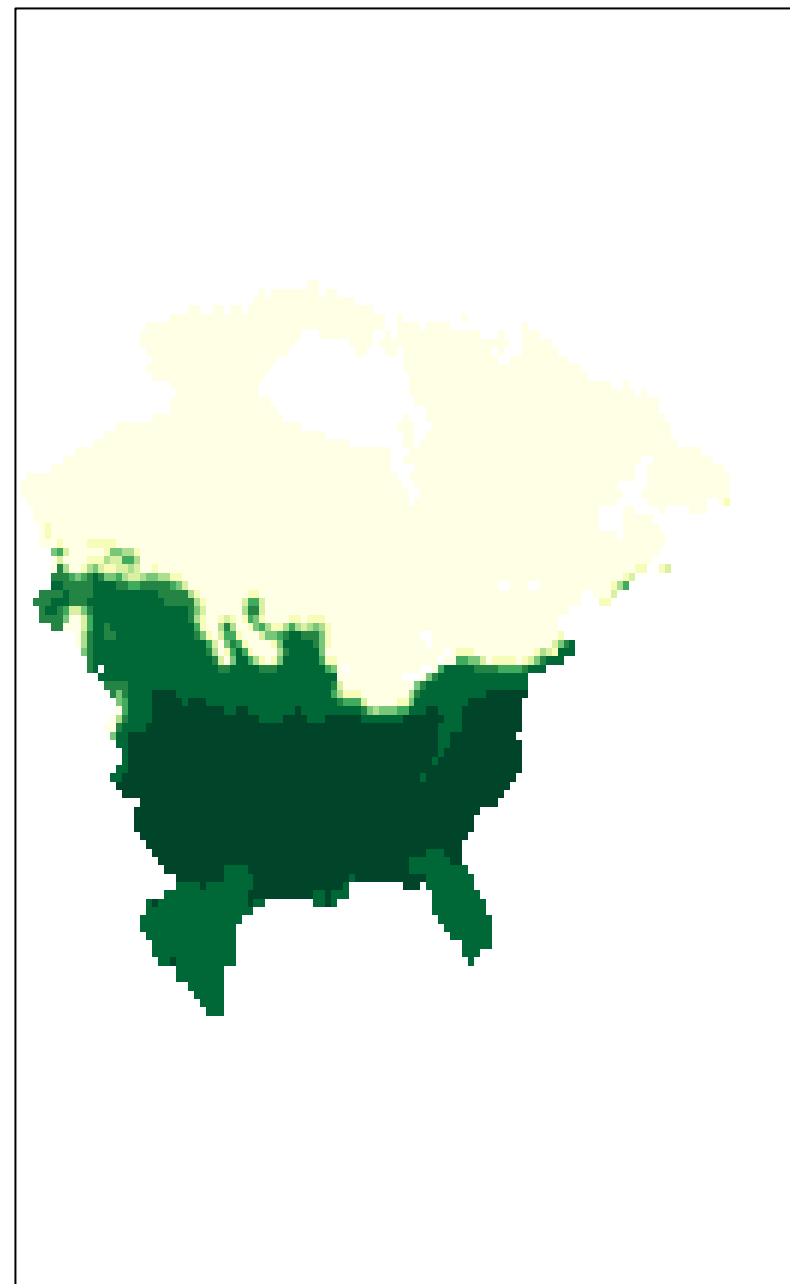


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X1000.ybp

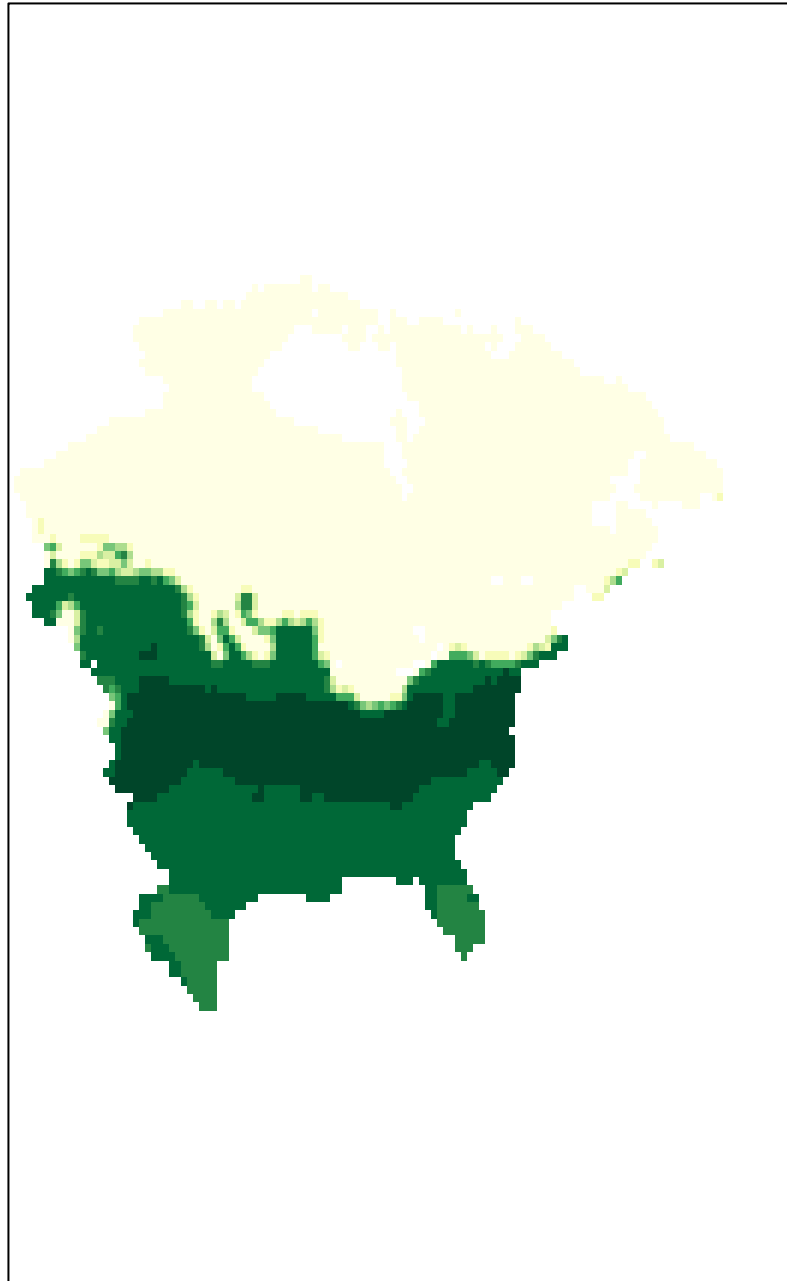


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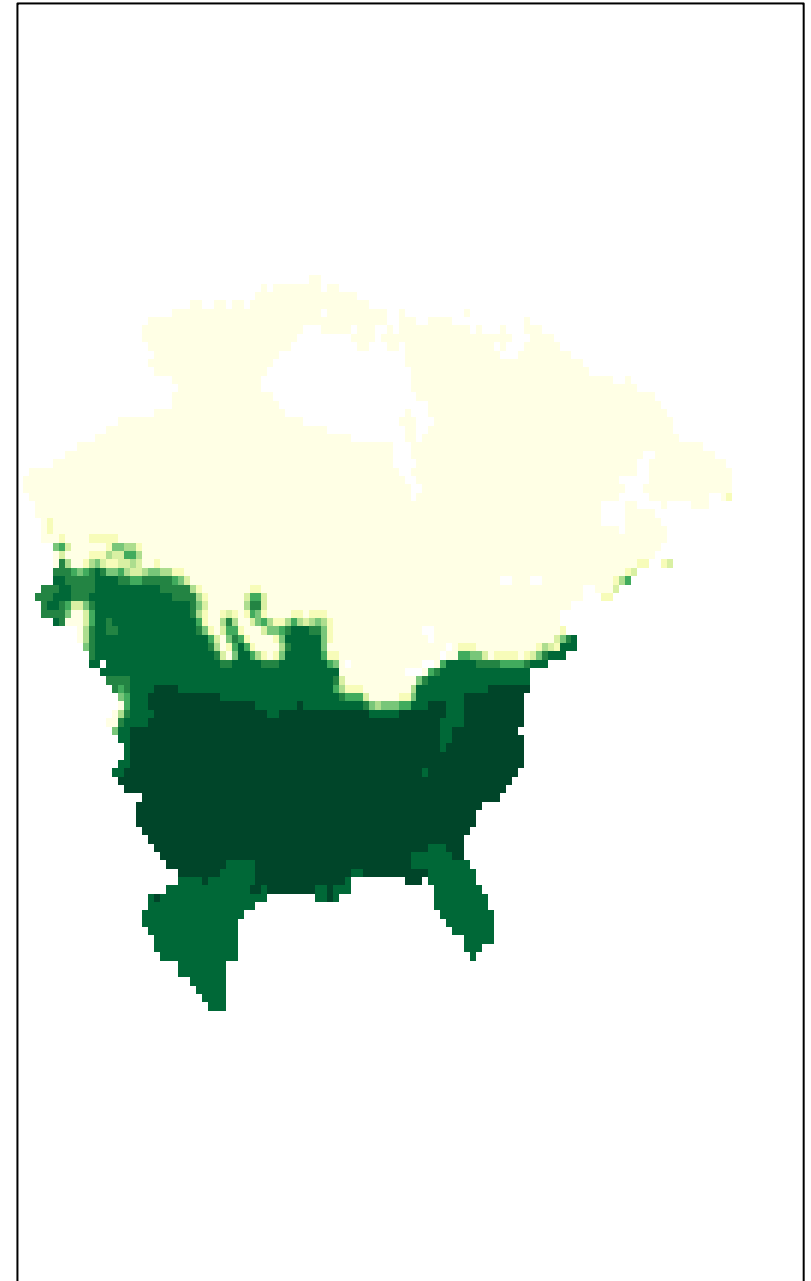


Species skipped = *Fraxinus greggii*, GCM = ecbilt

SUM, X0.ybp



SUM, X0.ybp



Species skipped = *Fraxinus greggii*, GCM = ecbilt