What can pollen tell us about palæo-climate?

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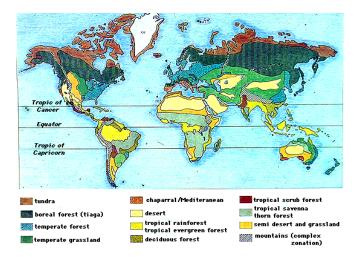
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Obligatory picture of biomes





Source: http://www.ucmp.berkeley.edu/exhibits/biomes/index.php



Source: http://www.life.illinois.edu/bio100/lectures/s97lects/04Ecosystems/BiomeMap.gif

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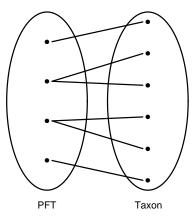
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- Reconstructed biomes (including measures of uncertainty) allow us to critique the performance of climate simulators 'out of sample', especially the new generation which will be run with dynamical vegetation models.
- 3. They may also allow us to calibrate climate simulators and vegetation models (i.e. to constrain the values of the uncertain parameters).

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But mainly because ...

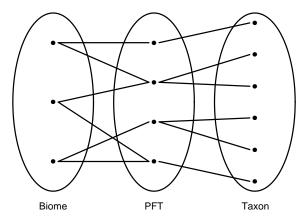
4. It's interesting and challenging!

Biomes, PFTs, and pollen taxa



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Biomes, PFTs, and pollen taxa



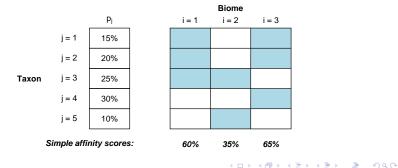
Defines a relationship between biomes and taxa, such that \mathcal{J}_i is the set of taxa that can be reached from biome *i*.

Current approach: Affinity score

Biomisation is estimating the biome from a pollen assemblage $\{x_j : j = 1, ..., n\}$. The dominant method is to maximise the affinity score:

$$\operatorname{Aff}(i) := \sum_{j \in \mathcal{J}_i} \hat{p}_j \qquad \hat{p}_j := \{\max(0, p_j - \theta)\}^\gamma$$

where p_j is the proportion of taxon j in the assemblage, and typically $\theta = 0.5\%$ and $\gamma = 1/2$.



Current approach: Affinity score

Some observations

- 1. The affinity score for biome j will be relatively high if and only if this biome contains the well-represented taxa.
- 2. The choice of $\gamma = 0.5$ down-weights taxa with large proportions, which is probably a crude adjustment for differential rates of productivity and dispersal.
- 3. But $\gamma = 0.5$ over-weights the contribution from the large number of taxa with small proportions, and so $\theta = 0.5\%$ is need to squash these.

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Some observations

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Statistical concerns

Is $\max_i \operatorname{Aff}(i)$ a good estimator? For example, is there a reasonable underlying statistical model from which it follows that the affinity score is the likelihood function?

- ► *If so,* we can quantify uncertainty and do hypothesis tests.
- If not, what confidence do we have in our reconstructions?

The affinity score is not a likelihood function



This follows by considering two biomes, *i* and *i'*, for which $\mathcal{J}_i \supset \mathcal{J}_{i'}$.

- Consider the set of pollen taxa that are in J_i but not in J_{i'}, *J* := J_i \ J_{i'}. Suppose that the counts are zero for all the taxa in J. In this case Aff(i) = Aff(i').
- 2. But one would think that getting zero counts in \mathcal{I} was improbable if the biome was *i*, but probable if the biome was *i'*. Therefore, statistically, we would want L(i) < L(i'), where *L* is the likelihood function.
- Extending this argument, if there were small numbers of counts in J then Aff(i) > Aff(i'), but we would still want L(i) < L(i') if these could be attributed to a background process.

Statistical model (sketch)

Aleatory model

- 1. The pollen grains on the microscope slide follow independent Poisson processes with rates θ_j that depend on the biome;
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Epistemic model

- 1. The biome rates θ_j are zero if $j \notin \mathcal{J}_i$, and otherwise Gamma $(\alpha_{ij}, \beta_{ij})$;
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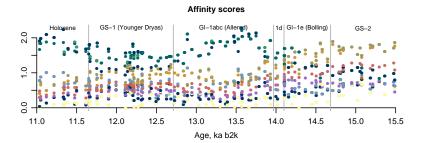
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Special vague case

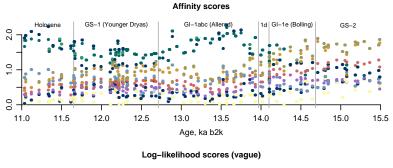
Set $\alpha_{ij} = \alpha_{ij}^{\lambda} = 1$. All of the β 's can then be set, in the very simple case, according to a single rate $\kappa \in (0, 1)$ which represents the expected proportion that comes from the background. One intuitive tuning parameter!

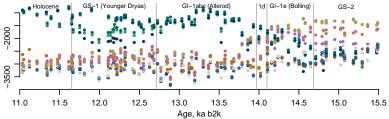


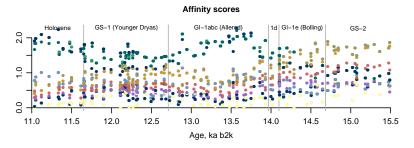




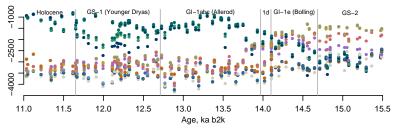
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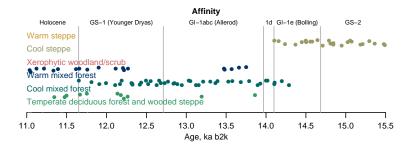






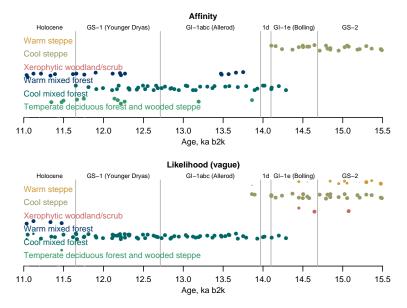


Another visualisation, Monticchio



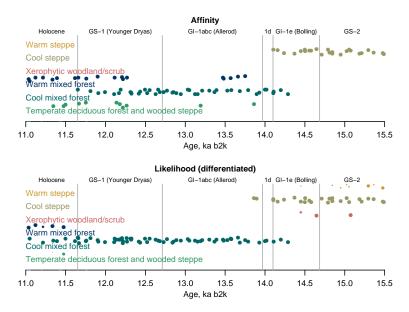
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Final observations

- Statistics is about doing sensible things when faced with uncertainty. In particular, how to make good estimates, like what the biome was at site x and time t BP.
- A crucial aspect of the statistical approach is its clarity. One is obliged to make explicit statements of one's judgements, that can be discussed and challenged.
- The benefit is the development, one hopes, of a consensus, and the much richer inference that is possible with more structured judgements.

Many thanks to: Prof Judy Allen, Dr Heather Binney, Dr Philip Brohan, Prof Caitlin Buck and the SUPRAnet project, Prof Mary Edwards, Prof Brian Huntley, Prof Jim Smith.