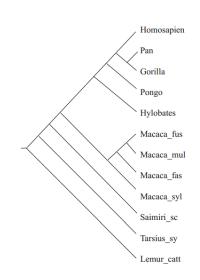
Averaging Phylogenetic Trees

Presenter: Suyi Wang

Phylogenetic Tree Space

- A phylogenetic n-tree is:
 - A Tree with n leaves
 - Leaves: represent distinguishable species
 - Interior vertices: #degree >= 3
 - Edges: weighted
 - Max: 2*n − 1 edges

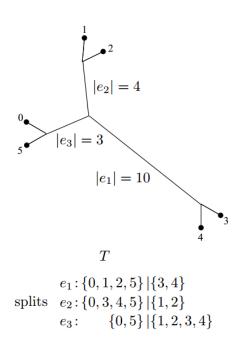
'Lemur_catta'	AAGCTTCATAGG
'Tarsius_syrichta'	AAGTTTCATTGG
'Saimiri_sciureus'	AAGCTTCACCGG
'Macaca_sylvanus'	AAGCTTCTCCGG
'Macaca_fascicul.'	AAGCTTCTCCGG
'Macaca_mulatta'	AAGCTTTTCTGG
'Macaca_fuscata'	AAGCTTTTCCGG
'Hylobates'	AAGCTTTACAGG
'Pongo'	AAGCTTCACCGG
'Gorilla'	AAGCTTCACCGG
'Pan'	AAGCTTCACCGG
'Homo_sapiens'	AAGCTTCACCGG

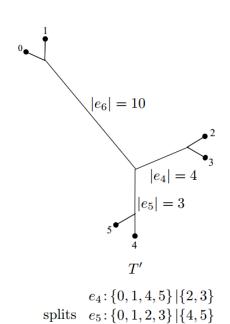


Phylogenetic Tree Space

• Splits:

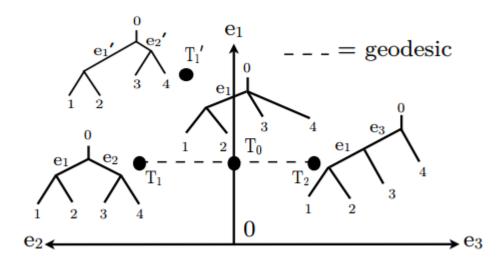
- Associates with an edge e
- A partition (Xe, Xe*) of leaves





Phylogenetic Tree Space

- Trees are uniquely defined by compatible set of splits [Semple 03]
 - Suggests a tree space
 - Axis length of edges
 - -2n-1 orthants



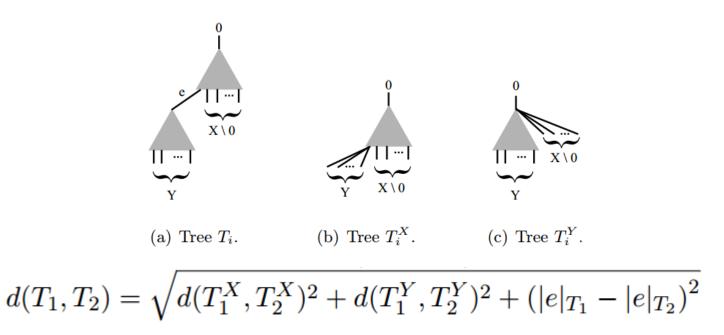
Geodesic distance

- Computing geodesic distances in tree space
 [Owen 07]
 - Shortest path from T1 to T2
 - Sum (sub path in each orthant)
 - Unique [Billera 01]

- Same orthant
 - Euclidean distance

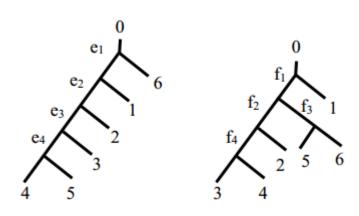
Geodesic distance

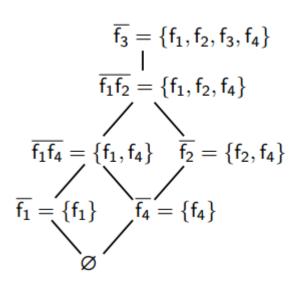
- Trees share a common edge
 - [Vogtmann 07] [Owen 08]



Geodesic distance

- Trees share no common edge
 - A path space: $K(\Sigma_1, \Sigma_2)$
 - Try all possible orthants series.
 - A dynamic programming





Averaging in tree space

• Tree space T={T1, T2, ..., Tr}

- Variance S(X, T)
 - Sum of squares distances from X to each tree in T

- Mean
 - X that minimize the variance

Averaging in tree space

- Iterative approach [Strum 03]
 - Random sample on T for k times {T1...Tk}
 - Current result u_i, i=[1, k]
 - $u_i+1 = 1/(k+1)$ from u_i to Ti
 - Converge to mean

- A descent method [Millier 12]
 - To be continued...