Consider an ordered sequence of tokens,  $\tau = (\tau_1, \tau_2, ..., \tau_n)$ . A parse of this sequence provides, under one interpretation, a nested partitioning of the tokens, e.g. either

Figure 1: Representing the parse as a bracketing of the tokens.

or

## $\underline{ \text{The cat}} \text{ sat } \underline{ \text{on } \underline{ \text{the mat}}} \text{ .}$

Figure 2: Representing the parse as a collection of token sub-sequences.

Under another interpretation, the parse forms a tree of nodes, e.g.

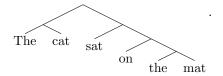


Figure 3: Representing the parse as a tree of nodes.

Thus, a parse tree can be characterised by a set of nodes and a set of relations or rules linking these nodes. The parse nodes are divided into leaf nodes and derived nodes. A *leaf* node represents all of the known information about a token, including the position of that token in the token sequence. A *derived* node represents a combination of a contiguous sub-sequence of leaf and/or derived nodes.

In order to define node contiguity, first observe from Figure 3 that each node  $\nu$  spans some contiguous sub-sequence of tokens, e.g.  $(\tau_i, \tau_{i+1}, ..., \tau_{i+k})$ . Let  $beta(\nu)$  be the index of the first token in the sub-sequence, e.g.  $\beta(\nu) = i$ , and let  $\varepsilon(\nu)$  be the index of the last token, e.g.  $\varepsilon(\nu) = i + k$ . Then the span of node  $\nu$  is defined as the set of token indices

$$\sigma(\nu) = \{ j \in \mathcal{I}_n \mid \beta(\nu) \le j \le \varepsilon(\nu) \},\,$$

where  $\mathcal{I}_n = \{1, 2, ..., n\}$ . Hence, for two arbitrary nodes  $\nu_1$  and  $\nu_2$  to be said to be non-overlapping, their spans must be mutually exclusive, i.e. obey  $\sigma(\nu_1) \cap \sigma(\nu_2) = \{\}$ . Furthermore, for  $\nu_1$  and  $\nu_2$  to be adjacent, their spans must further obey either  $\beta(\nu_1) = \varepsilon(\nu_2) + 1$  or  $\beta(\nu_2) = \varepsilon(\nu_1) + 1$ . Finally, an arbitrary sequence  $(\nu_1, \nu_2, ..., \nu_m)$  of nodes is contiguous if and only if

$$\beta(\nu_{i+1}) = \varepsilon(\nu_i) + 1, \forall i \in \mathcal{I}_{m-1}$$
.

Continuing now from previous remarks, a parse tree  $\mathcal P$  may be thought of as a set of node combination rules of the form

$$\nu_1 \ \nu_2 \dots \nu_m \xrightarrow{\rho} \nu$$
.

In order to be a valid rule, the sequence  $\pi(\rho) = (\nu_1, \nu_2, ..., \nu_m)$  of so-called *predecessor* nodes must be contiguous, such that the resulting derived node  $\delta(\rho) = \nu$  has span

$$\sigma(\nu) = \bigcup_{i=1}^{m} \sigma(\nu_i) ,$$

where  $\beta(\nu) = \beta(\nu_1)$  and  $\varepsilon(\nu) = \varepsilon(\nu_m)$ . In effect, the predecessor nodes partition the tokens spanned by  $\nu$  into mutually exclusive sub-sequences.

Taken together, the set  $\mathcal P$  of rules forms a parse of the whole token sequence, i.e.

$$\exists \rho \in \mathcal{P}, \ \sigma(\delta(\rho)) = \mathcal{I}_n \ .$$

Furthermore, the rules must partition the token sequence into a tree of nested sub-sequences, i.e.

$$\forall \rho_1 \in \mathcal{P}, \, \forall \rho_2 \in \mathcal{P} \setminus \{\rho_1\}, \, \sigma(\delta(\rho_1)) \bigcap \sigma(\delta\rho_2)) \neq \{\} \Rightarrow \sigma(\delta(\rho_1)) \subset \sigma(\delta(\rho_2)) \text{ or } \sigma(\delta(\rho_2)) \subset \sigma(\delta(\rho_1)).$$