## Queensland rainforest

We begin by loading the data with the two most prevalent species.

```
qld <- get_qld(prevalent = 2)
#> The chosen index corresponds to ep40.
configuration <- qld$configuration
window <- qld$window</pre>
```

```
par(mar = c(5, 4, 4, 13) + 0.1)
plot(configuration, window = window)
```

## Points in the configuration

