37 Participants enrolled
D0 samples, Pv (+): n = 37
ssPCR(+): n = 35

2-year follow up after treatment
Month 1: weekly
Month 2-4: monthly

No Pv recurrences: 9 participants

Pv recurrences: 28 participants
Total episodes, Pv (+): n = 142
ssPCR(+): n = 140

1 Pv sample was not analysed by MS genotyping

1st year (n=66)
2nd year (n=76)

MS genotyping
D0 + recurrences
n = 174 Pv samples

Overall genetic diversity analysis (n=136)
Monoclonal/complete allele data (n=125)
Monoclonal/incomplete allele data (n=11)
Polyclonal/complete+incomplete (n=28)
Negative PCR amplification (n=10)

Identified haplotypes (n=162)

Figure 2