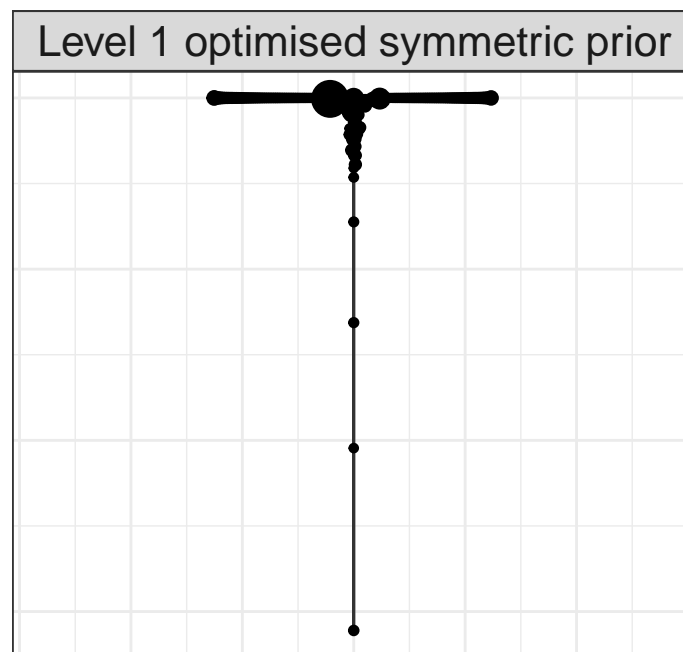
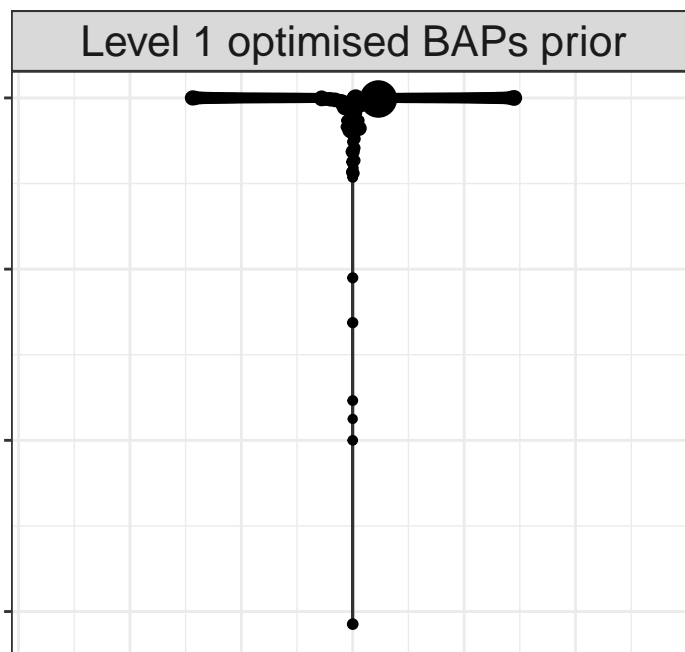
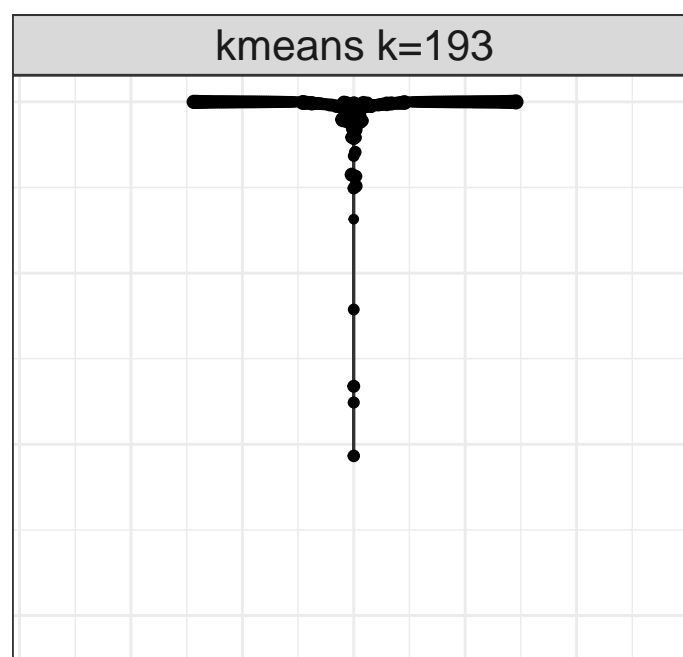
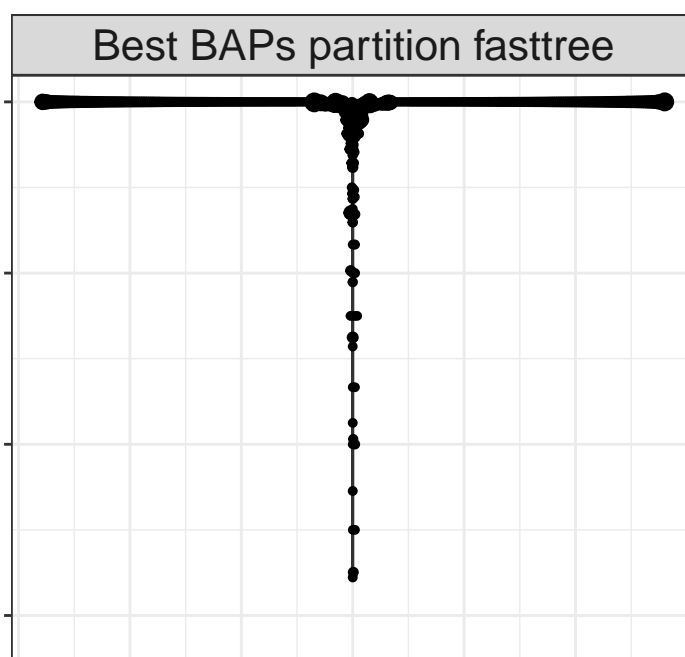


Proportion comprising the dominant HIV subtype



method

- Best BAPs partition fasttree
- kmeans k=193
- Level 1 optimised BAPs prior
- Level 1 optimised symmetric prior

cluster size

- 10000
- 20000
- 30000