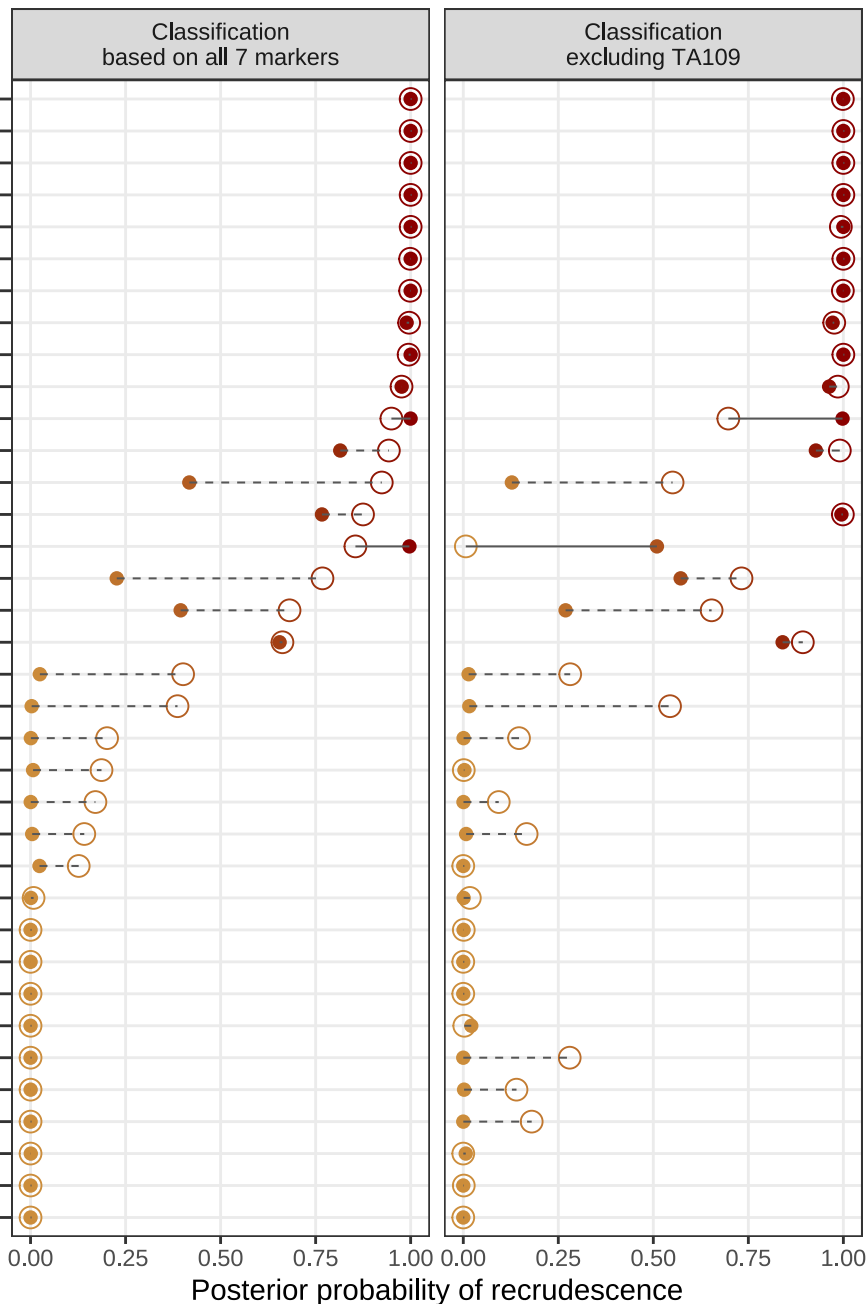
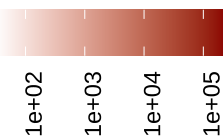


Patient ID

	D0 MOI	DX MOI
ZQ21-030	3	1
ZL21-304	4	2
ZL21-269	2	2
ZL21-263	2	2
LL21-089	2	1
ZL21-203	2	3
BD21-041	2	2
ZL21-245	3	1
LL21-054	2	3
ZQ21-103	5	1
ZQ21-077	3	1
ZL21-233	2	2
LL21-061	3	1
LL21-077	2	5
ZL21-260	5	3
BD21-053	4	4
LL21-094	4	3
ZQ21-085	3	3
ZL21-289	4	2
ZL21-242	2	3
LL21-008	2	4
ZL21-227	2	2
BP21-280	4	2
ZQ21-046	2	3
BP21-245	3	3
ZL21-209	3	3
ZQ21-094	2	2
ZQ21-031	2	1
ZL21-291	2	1
LL21-086	2	2
LL21-038	2	3
LL21-037	3	2
LL21-027	3	1
BP21-233	3	1
BD21-099	3	1
BD21-040	2	1

Parasitemia (per  $\mu$ L)



Model

- Probability of at least one recrudescence clone (M1) under PfRecur (genotyping error rate: 0.05, per-clone marker-wise detection prob: 0.9)
- Posterior probability of recrudescence under CDC model