Branch number, with a bootstrap value of 1 and containing 3 isolates or more. See figure below	Maximum nb of years between to isolates sampling date	Nombre de SNP max	Name of the associated clone
1	7	504	
2	23	796	
3	10	150	
4	4	60	
5	***2***	***13***	***C1***
6	***9***	***34***	***B1***
7	23	401	
8	***5***	***48***	***A1***
9	***3***	***10***	***A1-sub***
10	5	168	
11	***5***	***26***	***A4***
12	***18***	***93***	***A3***
13	3	31	
14	***23***	***112***	***A2***
15	3	37	

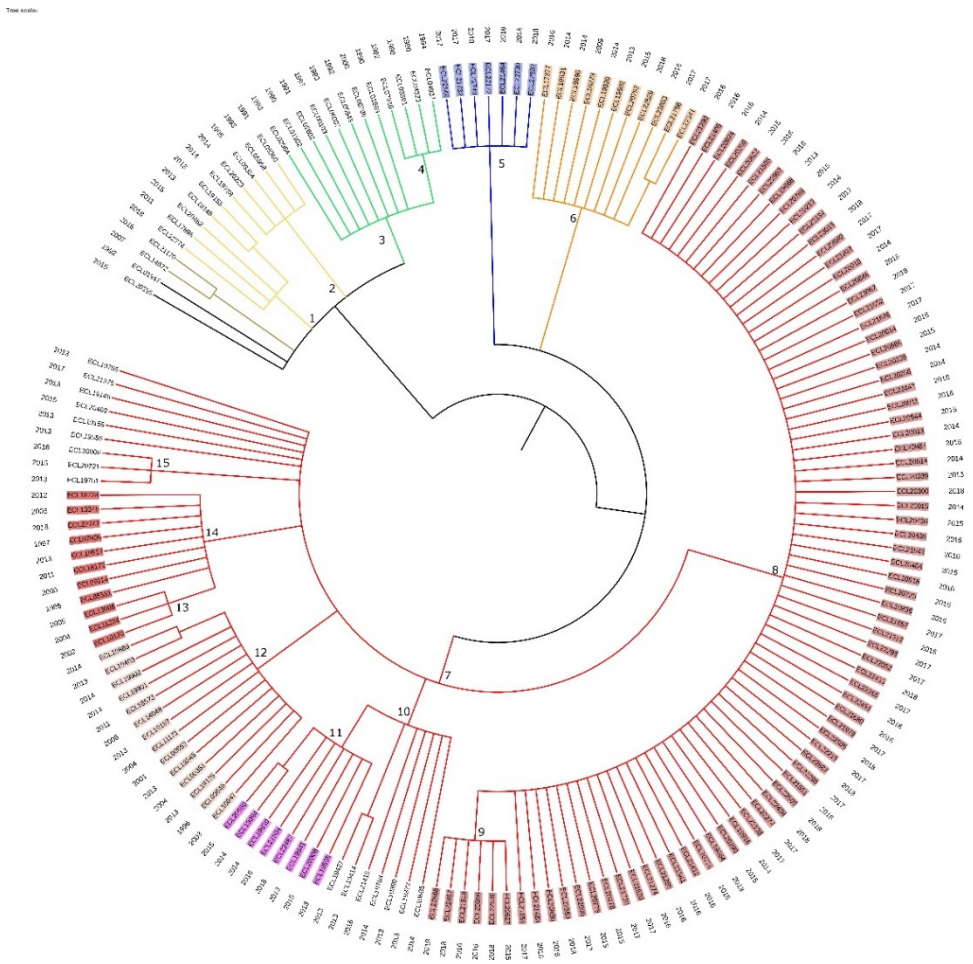

Figure S1 : Number of maximun SNPs in relation to the number of years between two isolates within each branch tree with a bootstrap value of one and composed of 3 isolates and more. The bolded branches in the table are considered as clones. In this phylogenetic analysis the number of SNP per year should not exceed 10 (MxP = $3x10^{-6}$ x 3 366 150).