

**Supplementary Table S4** The eight *Xanthomonas citri* pv. *citri* (Xcc) VNTRs used in this study found in the *X. citri* pv. *viticola* strains CFBP 7764 and LMG 965 (CFBP 7660) genome sequences and the obtained copy numbers after sequencing the PCR- amplified loci for both strains

VNTR loci	Xcc repeat sequence	Whole genome sequencing		Amplicon sequencing	
		Xcvt repeat sequence	Copy number predicted CFBP 7764 / LMG 965	Xcvt repeat sequence	Copy number obtained CFBP 7764 / LMG 965
XL1	TTCCCCA	TTCCCCA	4/4	TTCCCca (gc) (ag)	9/10
XL3	TTCCCGA	TTCCCGA	10/6	TTtCCGA/ TTcCCGA	13/9
XL4	CCCGAAT	CCCGAAT	10/6	CCCGAaT/ CCCGAtT	11/6
XL5	TTCCCGG	aTCCCGA	8/10	aTCCCGA /tTCCCGA	8/10
XL6	CCGATTC	CCGcTTC	24/24	CCGcTTC /CCGaTTC	24/24
XL8	CGGGAAT	CGGGAAT	2/3	CGGgAAT/CGGaAAT	5/13
XL13	CGGGAAT	CGTGAAT	8/8	CGTGAAT	7/8
XL15	CGAATCC	GGAATCG	10.4/5.4	CGAtTCC/ CGAaTCC	10/5