

c.2Ktedonobacter_racemifer_DSM_44963,_NZ_ADVG01000002.1	23
4u.CONSENSUSStreptomyces_yunnanensis,_NZ_FRBK01000018.1	25
	26
	27
4c.2Planktothrix_rubescens,_NZ_KE734718.1	28
	29
	3)Cas14u.6Ktedonobacter_racemifer,_NZ_ADVG0
	30
4u.7Planktothrix_tepida_PCC_9214,_NZ_LN889780.1	32
	33
4u.7Streptomyces_rapamycinicus,_NZ_QYCY01000002.1	34
	4)Cas14h.CONSENSUSKtedonobacter_racemifer_
	40
	43
	45
	46
c.CONSENSUSKtedonobacter_racemifer,_NZ_ADVG01000001.1	5)Cas14c.2Ktedonobacter_racemifer_DSM_44963
	50
	52
	53























