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<b>Title</b>	Things are getting hairy: Enterobacteria bacteriophage vB_PcaM_CBB
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## Supplementary S4

**Table S1 - Phage CBB codon usage (CBB\_1 - CBB\_554) with tRNA gene codons highlighted**

aa F	Frequency	Codon	Occurrence	Codon Freq.	Freq./aa	tRNA occurrence
Ala	5.83%	GCA	3058	2.86%	49.06%	1
		GCC	313	0.29%	5.02%	
		GCG	584	0.55%	9.37%	
		GCT	2278	2.13%	36.55%	
Arg	4.09%	CGA	380	0.36%	8.69%	
		CGC	432	0.40%	9.88%	
		CGG	130	0.12%	2.97%	
		CGT	2587	2.42%	59.17%	1
		AGA	684	0.64%	15.65%	1
		AGG	159	0.15%	3.64%	
Asm	6.16%	AAC	2506	2.34%	38.06%	1
		AAT	4079	3.81%	61.94%	
Asp	6.65%	GAC	1525	1.43%	21.44%	1
		GAT	5588	5.22%	78.56%	
Cys	1.14%	TGC	287	0.27%	23.62%	1
		TGT	928	0.87%	76.38%	
Gln	3.60%	CAA	2750	2.57%	71.39%	1
		CAG	1102	1.03%	28.61%	
Glu	7.35%	GAA	6766	6.33%	86.13%	1
		GAG	1090	1.02%	13.87%	
Gly	5.83%	GGA	1316	1.23%	21.12%	1
		GGC	565	0.53%	9.07%	
		GGG	336	0.31%	5.39%	
		GGT	4015	3.75%	64.43%	
His	1.97%	CAC	593	0.55%	28.13%	1
		CAT	1515	1.42%	71.87%	
Ile	6.88%	ATA	1017	0.95%	13.83%	
		ATC	1620	1.51%	22.02%	1
		ATT	4719	4.41%	64.15%	
Leu	7.75%	CTA	1751	1.64%	21.13%	1
		CTC	264	0.25%	3.19%	
		CTG	715	0.67%	8.63%	
		CTT	1750	1.64%	21.12%	
		TTA	2521	2.36%	30.43%	2
		TTG	1284	1.20%	15.50%	1
Lys	6.58%	AAA	5314	4.97%	75.47%	1
		AAG	1727	1.61%	24.53%	
Met	2.35%	ATG	2512	2.35%	100.00%	3
Phe	4.58%	TTC	2109	1.97%	43.08%	2
		TTT	2786	2.60%	56.92%	
Pro	3.28%	CCA	1951	1.82%	55.70%	2
		CCC	86	0.08%	2.46%	
		CCG	329	0.31%	9.39%	
		CCT	1137	1.06%	32.46%	
Ser	6.63%	TCA	1883	1.76%	26.54%	1
		TCC	329	0.31%	4.64%	

		TCG	313	0.29%	4.41%	
		TCT	1836	1.72%	25.88%	1
		AGC	544	0.51%	7.67%	1
		AGT	2189	2.05%	30.86%	
Thr	6.41%	ACA	2141	2.00%	31.23%	1
		ACC	677	0.63%	9.87%	
		ACG	477	0.45%	6.96%	
		ACT	3561	3.33%	51.94%	
Trp	1.21%	TGG	1289	1.21%	100.00%	1
Tyr	4.89%	TAC	1752	1.64%	33.53%	1
		TAT	3473	3.25%	66.47%	
Val	6.85%	GTA	3095	2.89%	42.22%	
		GTC	365	0.34%	4.98%	1
		GTG	666	0.62%	9.08%	
		GTT	3205	3.00%	43.72%	1