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Supplementary S4 Table S1 - Phage CBB codon usage (CBB_1 - CBB_554) with tRNA gene codons highlighted

aa F	Frequency		Occurrence			tRNA occurance	
Ala	5.83%		3058				1
Ald	5.65%	GCA					
			313				
		GCG	584				
•	4.000/	GCT	2278		36.55%		_
Arg	4.09%		380		8.69%		
		CGC	432				
		CGG	130		2.97%		_
		CGT	2587	2.42%	59.17%		1
		AGA	684		15.65%		1
		AGG	159		3.64%		
Asm	6.16%		2506				1
		AAT	4079	3.81%	61.94%		_
Asp	6.65%		1525		21.44%		1
		GAT	5588				
Cys	1.14%		287	0.27%	23.62%		1
		TGT	928	0.87%	76.38%		
Gln	3.60%		2750		71.39%		1
		CAG	1102	1.03%			
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%		
lle	6.88%	ATA	1017	0.95%	13.83%		
		ATC	1620	1.51%	22.02%		1
		ATT	4719	4.41%	64.15%		
Leu	7.75%	СТА	1751	1.64%	21.13%		1
		стс	264	0.25%	3.19%		
		CTG	715	0.67%	8.63%		
		СТТ	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%		2109	1.97%	43.08%		2
		TTT	2786		56.92%		_
Pro	3.28%		1951	1.82%	55.70%		2
	0.2070	CCC	86		2.46%		-
		CCG	329		9.39%		
		ССТ	1137	1.06%	32.46%		-
Ser	6.63%		1883		26.54%		1
	0.0370	TCC	329		4.64%		-
		100	529	0.51/6	7.04/0		

		TCG	313	0.29%	4.41%	
		тст	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