



(d)

Genome binner (% contamination)	Predicted bins (% completeness)			
	>50%	>70%	>90%	
CONCOCT (CAMI)	<10%	129	129	123
	<5%	124	124	118
CONCOCT 2	<10%	239	232	196
	<5%	226	222	189
MaxBin 2.0.2 (CAMI)	<10%	277	274	244
	<5%	254	252	224
MaxBin 2.2.4	<10%	274	271	236
	<5%	249	247	216
MetaBAT (CAMI)	<10%	173	152	126
	<5%	159	140	118
MetaBAT 2.11.2	<10%	427	417	361
	<5%	414	404	353
Metawatt 3.5 (CAMI)	<10%	408	387	338
	<5%	396	376	330
Metawatt 3.5.3	<10%	64	52	25
	<5%	51	44	22
MyCC (CAMI)	<10%	189	182	145
	<5%	166	159	127
Binsanity 0.2.5.9	<10%	9	9	9
	<5%	6	6	6
Binsanity-refine 0.2.5.9	<10%	206	204	192
	<5%	183	181	171
COCACOLA	<10%	88	87	75
	<5%	69	69	60