

plasmid: Statistics of Functional Gene Prediction Analysis by Barcode					
Barcode	Total Predicted Proteins	Hypothetical Proteins	Not EggNog Annotations	Proportion Hypothetical	Proportion Not EggNog Annotated
barcode30	7,049	5,453	2,343	77.4%	33.2%
barcode31	1,177	1,040	506	88.4%	43.0%
barcode32	2,813	2,236	924	79.5%	32.8%
barcode33	3,081	2,561	1,146	83.1%	37.2%