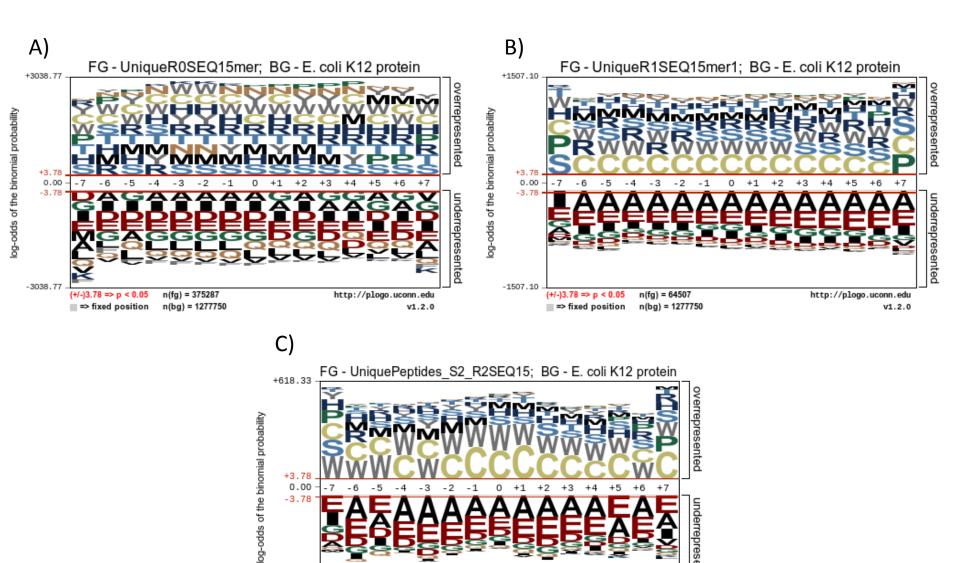
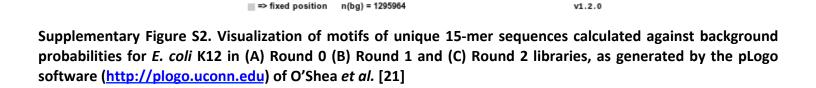


Supplementary Figure S1. Amino acid representation within all unique sequences within the library excluding (A) and including (B) the effects of frequency of sequence occurrence, for Round 0 (blue), Round 1 (orange) and Round 2 (grey).





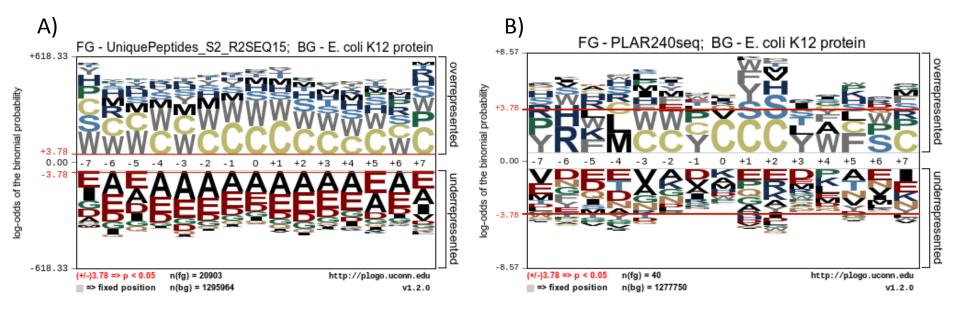
n(fg) = 20903

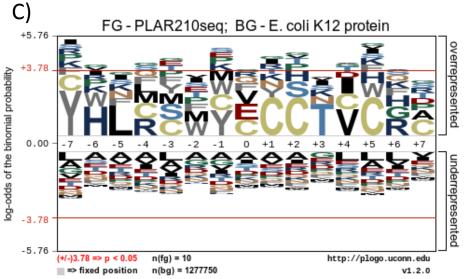
underrepresentec

http://plogo.uconn.edu

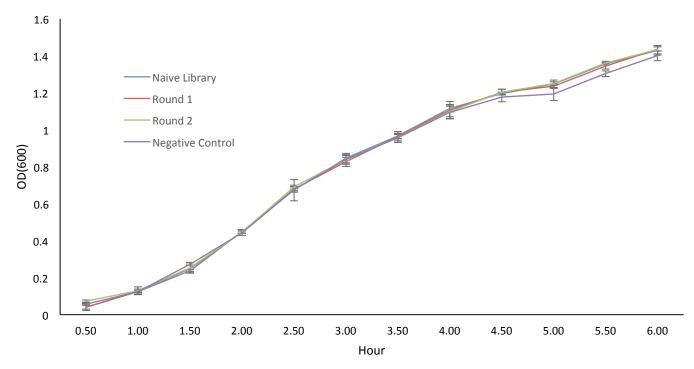
0.00 -3.78

-618.33





Supplementary Figure S3. Visualization of motifs of unique 15-mer sequences calculated against background probabilities for *E. coli* K12 in Round 2 library (A) all 15-mers (B) 40 most frequently occurring 15-mers and (C) 10 most frequently occurring 15-mers, as generated by the pLogo software (http://plogo.uconn.edu) of O'Shea *et al.* [21]



Supplementary Figure S4. Growth curves over 6 hours for the naive library, both sorting rounds, and the negative control after a 1:100 dilution from overnight cultures in fresh media.

		Valid Inserts	Frame Shift	Stop Codon	Empty Scaffold	Total # of Sequences
Round 0	Total	466,476	331,377	255,658	158,791	1,212,302
	Unique	396,641	313,601	224,685		
Round 1	Total	591,400	350,783	299,598	176,237	1,418,018
	Unique	67,885	208,322	42,017		
Round 2	Total	770,853	279,786	201,774	152,954	1,405,367
	Unique	21,931	133,599	7,966		

Supplementary Table S1. Raw count of total and unique sequences for all sequence categories of each library.

		0 cysteines	1 cysteine	2 cysteines	3 cysteines	4 cysteines
Round 0	raw count	252165	116024	24846	3275	309
	decimal	0.635751	0.292516	0.062641	0.008257	0.000779
Round 1	raw count	42299	20649	4305	564	65
	decimal	0.623098	0.304176	0.063416	0.008308	0.000958
Round 2	raw count	12344	7809	1497	263	18
	decimal	0.562856	0.356071	0.06826	0.011992	0.000821

Supplementary Table S2. Raw count and decimal of total sequences in library containing 0-4 cysteines, round by round.