## Table S5. Sequencing of the *hsvtk* regions of the counter-selection escapees arose from dP sensitive strain MG1655 $\triangle lacZ$ :: $hsvTK-km^r$ .

dP sensitive clone was cultured overnight and plated on dP-selection plate from which nine clones were picked, and sequenced their *hsvtk* coding regions.

Clone	Nucleoside substitution	Amino Acid substitution
number		
1	T865C	F289L
2	A472C	T158P
3	A472C	T158P
4	865TTCACCCTG> 865TTCACCCTG - TTCACCCTG	<sup>289</sup> FTL> <sup>289</sup> FTL-FTL
5	A196C	Т66Р
6	A164C	D55A
7	865 TTCACCCTG> 865 TTCACCCTG - TTCACCCTG	<sup>289</sup> FTL> <sup>289</sup> FTL-FTL
8	A683C	D228A
9	A196C	T66P