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4-14-2023

Creation of an N-terminal xCT Mutant Lacking Lysines for Use in Protein Turnover Studies

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Repository citation: Switzer, Alexandria, "Creation of an N-terminal xCT Mutant Lacking Lysines for Use in Protein Turnover Studies" (2023). 22nd Annual Celebration of Undergraduate Research and Creative Activity (2023). Paper 39. https://digitalcommons.hope.edu/curca_22/39 April 14, 2023. Copyright © 2023 Hope College, Holland, Michigan.

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Creation of an N-terminal xCT mutant lacking lysines for use in protein turnover studies Alexandria Switzer^{a,b,c}, and Leah Chase^{a,b,c} Departments of ^aChemistry and ^bBiology and ^cProgram of Neuroscience, Hope College, Holland, MI, 49423 Methods **Comparison of 6K xCT mutant amino** Background acid sequence to native Homo sapiens **(a) (b)** Homo sapiens solute carrier family 7 member 11 (SLC7A11), mRNA Mutant Strand Synthesis Miniprep Sequence ID: <u>NM_014331.4</u> Length: 9645 Number of Matches: 1 Perform thermal cycling to: Denature DNA template Range 1: 281 to 998 GenBank Graphics Vext Match Anneal mutagenic primers containing desired mutation Strand Identities Extend and incorporate primers with our 15/730(2%) 688/730(94%) Plus/Plus 1101 bits(596) exclusive Pfu-based DNA Polymerase Blend Total reaction time: 1 hour* AACGGGAGGCTGCCTTCCCTGGGCAACAGGGAGCCACCTGGGCAGGAGAGAGTGCAGCTG Sequencing and 2. Faster Dpn I Digestion of Template Sbjct 341 AACGGGAGGCTGCCTTCCCTGGGCAACAAGGAGCCACCTGGGCAGGAGAAAGTGCAGCTG Digest parental methylated and hemimethylated Analysis DNA with NEW Dpn I enzyme AGGAGGAGAGTCACTTTACTGAGGGGAGTCTCCATTATCATTGGCACCATCATTGGAGCA Total reaction time: 5 minute 401 AAGAGGAAAGTCACTTTACTGAGGGGAGTCTCCATTATCATTGGCACCATCATTGGAGCA 460 Figure 5. Nucleotide-blast for mutated xCT. A nonsynonymous SNP . Transformation \circ 6 K's are in the N-terminus of xCT: 4, 12, 30, 37, 41, 43. Four from $A \rightarrow G$ at positions 291, 315, 369, 390, 402, and 408 causes a Transform mutated molecules into competent cells for nick repair K's are conserved across species (4, 37, 41, 43). (Figure 1) missense mutation of Lys \rightarrow Arg in K4R, K12R, K30R, K37R, K41R, and Total reaction time: 1.5 hours K43R. * Based on a 5-kb plasmid; excludes ramping time. Figure 3. Schematic of overall experimental approach. (a) Process for inserting successive mutations in the Conclusions N-terminus of xCT (QuikChange Lightning Manual, 2015). We started with K37,41,43R and successively replaced Previous work in the Chase lab has shown the replacement of K's at the other 3 positions. (b) Following transformation, the samples were mini-prepped and sent to Eurofins for DNA sequencing. individual lysines with arginines does not lead to obvious changes the ubiquitination of a transporter **Sanger Sequencing Trace for N-terminus of xCT** Currently, all 6 N-terminal lysines in xCT have been replaced by Legend: PTM sites Phosphorylation arginines. Ubiquitylation • It is unknown if complete removal of lysines in the N-terminus Other CAGAAGGCCTGTTGTGTCCACCATCTCCAGAGGAGGTTACCTGCAGGGAAATGTTAACGGGAGG **(a)** will result in a lack of ubiquitination or change in transporter function or overall expression. • • • <u>IVRKPVVSTISKGGYLQGNVNGRLPSLGNKEPZGQEKVQLKRKVTI</u> 5 10 15 20 25 30 35 40 45 **Future Directions** • Transfect COS-7 cells (mammalian cells) with the 6K xCT mutant to study its level of expression and function in cells • Immunoprecipitate the 6K xCT from the cells and perform Western blotting to examine the uibquitination status of the 6K xCT mutant Examine the rate of turnover and cell surface expression of the **Research Question** 6K xCT mutant. • If we find that xCT is still being ubiquitinated, investigate what **(b**) other aspects of xCT may allow ubiquitin to bind to the protein. Hypothesis Acknowledgments Schaap Endowed Funds for Undergraduate Research • Amanda Gibson for creating the initial 3KR xCT mutant References

Function of xCT

- System x⁻ is an antiporter that exchanges intracellular glutamate for extracellular cystine
- Cystine is reduced to cysteine, which is necessary for the synthesis of glutathione (GSH)
- GSH is a reducing agent that mitigates oxidative stress in cells
- System x⁻_c consists of the light chain protein, xCT, that confers specificity, and the heavy chain protein, 4F2HC
- Trafficking of xCT is thought to modulate the oxidative stress response

Effects of Ubiquitin

- Ubiquitin tags on proteins typically leads to their internalization or degradation
- Ubiquitin reversibly attaches to lysine (K) residues
- Other proteins, such as epithelial sodium channel, are regulated by ubiquitin, with increased ubiquitination resulting in decreased membrane expression

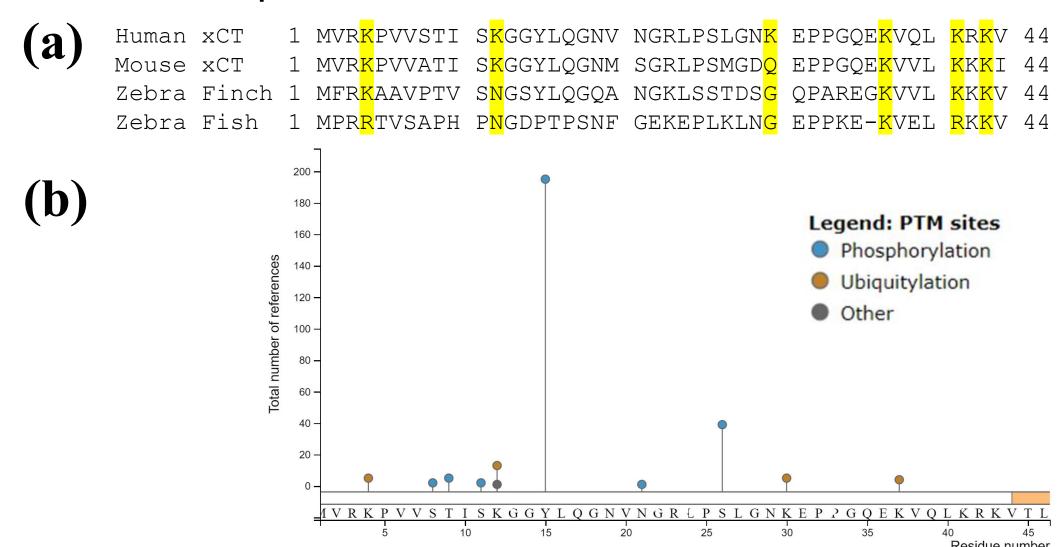


Figure 1. N-terminus xCT DNA sequences for four species. (a) A comparison of the DNA sequence of xCT for *Homo sapiens* compared to 3 other species. Most of these lysines are highly conserved across species. (b) Post-translational modifications of amino acids in the N-terminus of xCT identified through high-throughput screening (Phosphosite plus).

How does the lack of ubiguitination on the N-terminus of xCT affect the function of the protein?

Replacing all of the lysines in the N-terminus of xCT with arginines will lead to a loss of ubiquitination and an overall increase in the activity and overall expression of the transporter.

5
H ₂ N COOH
Lysine (K)

ОН	H ₂ N C
<)	Arginine (R

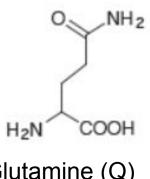
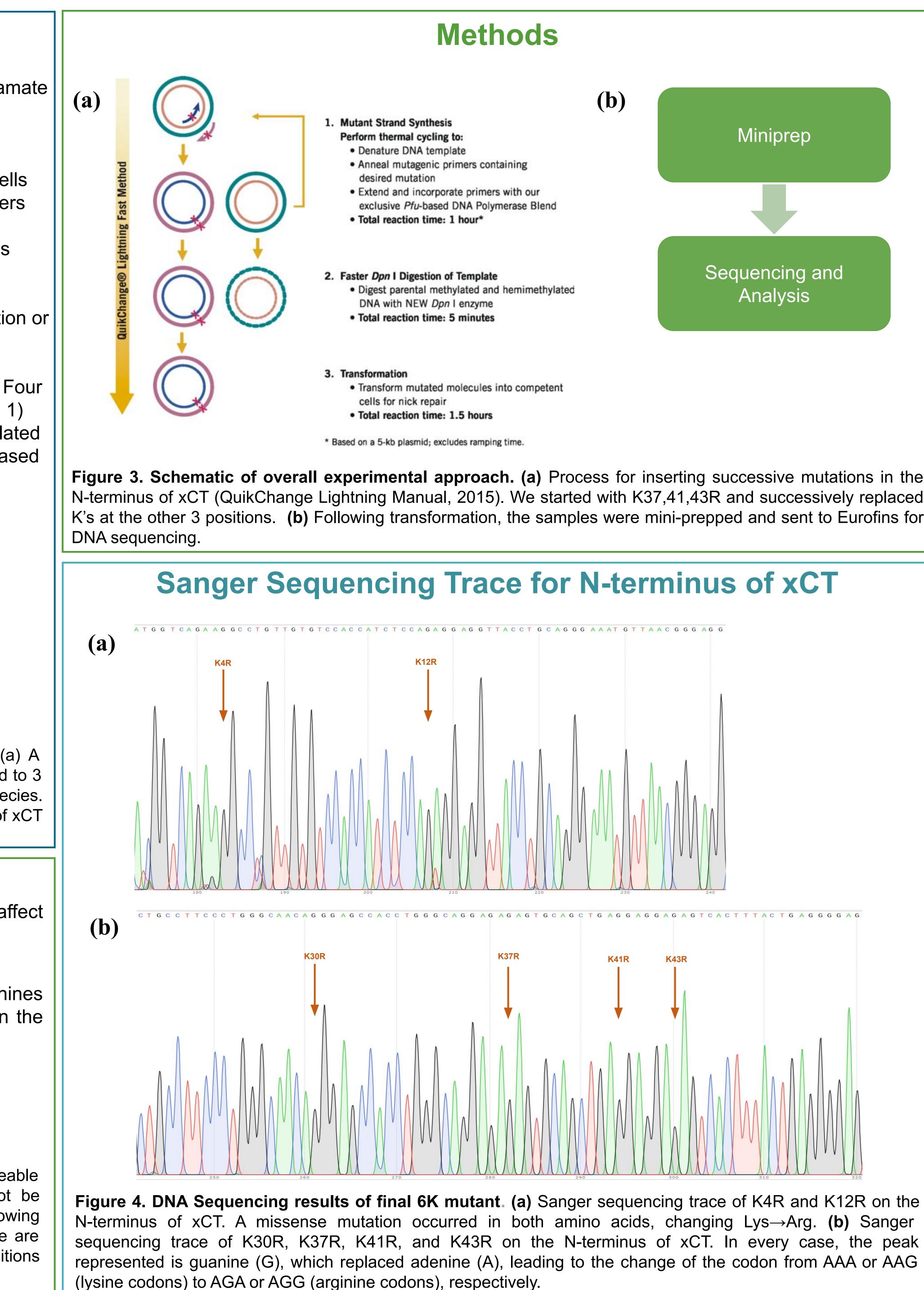


Figure 2. Mutagenesis strategy. Arginine and lysine are interchangeable because they are similar in size and charge, but arginine cannot be ubiquitinated, allowing for preservation of protein structure while allowing us to test the importance of ubiquitnation. Asparagine and glutamine are amino acids that are found in other species (besides humans) at positions for K. These can serve as acetylated mimetics of lysines.



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