## Supercharged Green Fluorescent Proteins as Bimodal Reporter Genes for CEST and Optical Imaging

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Target Audience: Researchers and clinicians who are interested in molecular and cellular MRI especially in non-invasive monitoring of gene expression and gene therapy.

*Purpose*: The overall goal is to modify the optical reporter gene green fluorescent protein (GFP) into an MRI-based reporter gene based on previous studies on the lysine rich protein (LRP)<sup>1</sup> and GFP<sup>2</sup>.

Methods: E. coli optimized genes encoding to wild type GFP (wt) and its superpositively charged variants (+36 and +48), achieved by modifying the solvent-exposed amino acids to lysine or arginine<sup>3</sup>, were obtained from Dr. David R. Liu (Harvard University, Cambridge, MA<sup>3-5</sup>). The proteins were expressed in BL21 (DE3) E. coli after induction in Magic Media<sup>TM</sup> and purified using cobaltbased immobilized metal affinity chromatography. The expression and purity was determined by SDS-polyacrylamide gel electrophoresis (SDS-PAGE). Pure proteins were dialyzed against PBS and their CEST-MRI characteristics were measured using a

modified RARE sequence (TR/TE=6000/9.4 ms) including a magnetization transfer module (B<sub>1</sub>=4.7 µT/4000 ms). The absolute water resonant frequency shift was measured using a modified WASSR method, with the same parameters as in CEST imaging except TR=1.5 sec and B<sub>1</sub> saturation pulse=0.5 µT/250 ms. Mean CEST spectra were derived from a ROI for each sample after B<sub>0</sub> correction for each voxel using MatLab. MTR asymmetry (MTR<sub>asym</sub>)=100×(S<sub>-Δω</sub> - $S_{+\Delta\omega}$ )/  $S_0$  was computed at different offsets of  $\Delta\omega$ .

Results: Fig. 1a shows fluorescence images of solutions containing the wt, +36, and +48 GFP variants. The high purity of the three proteins was validated by SDS-PAGE (Fig. 1b). The CEST MRI characteristics of the three examined GFPs (wt, +36, and +48) showed that both superpositive GFPs (+36 and +48) generated higher MTR<sub>asym</sub> values as compared to wt-GFP at both 1.5 ppm (guanidine protons of arginine) and 3.3 ppm (amide protons) (Fig. 1d-f). The +48 GFP generated higher MTR<sub>asym</sub> values at the amide frequencies (3.3 ppm) than the +36 mutant. Fig. 1e-f display MTR<sub>asym</sub> maps that reveal the correlation between the protein charge and the CEST contrast.

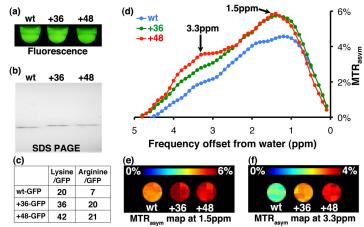


Fig 1. Characterization of wt-GFP and its superpositive mutants +36-GFP and +48-GFP. a) Fluorescence. b) SDS-PAGE of purified GFPs. c) Number of lysine and arginine amino acids in each GFP (total 239 amino acids). d) MTR<sub>asym</sub> plots. e) MTR<sub>asym</sub> maps (Δω=1.5 ppm). e) MTR<sub>asym</sub> maps ( $\Delta\omega$ =3.3 ppm).

Discussion: Superpositively charged GFPs are highly resistant to aggregation and retain high fluorescence levels even after being boiled and cooled<sup>3</sup>. Here we show that the +36 and +48 GFPs mutants generate high CEST MRI contrast. Both examined mutants contain an equal number of arginine residues (20 and 21, respectively) and therefore generated the same MTR<sub>asym</sub> value from the guanidine exchangeable protons at  $\Delta\omega$ =1.5 ppm. In contrast, as +48-GFP mutant has more lysines as compared to +36-GFP (42 and 36, respectively), it generated higher MTR<sub>asym</sub> values at the resonance of the amide protons at  $\Delta\omega$ =3.3 ppm.

Conclusion: Superpositively charged GFP, a lysine/arginine rich fluorescent protein, is an excellent reporter gene for CEST MRI while retaining its capability as an optical reporter gene. Supported by MSCRFF-0103-00.

References: 1. A. A. Gilad et al., Nat Biotechnol 25, 217 (2007). 2. C. J. Perez-Torres, C. A. Massaad, S. G. Hilsenbeck, F. Serrano, R. G. Pautler, Neuroimage 50, 375 (2010). 3. M. S. Lawrence, K. J. Phillips, D. R. Liu, J Am Chem Soc 129, 10110 (2007). 4. J. J. Cronican et al., ACS Chem Biol 5, 747 (2010). 5. B. R. McNaughton, J. J. Cronican, D. B. Thompson, D. R. Liu, Proc Natl Acad Sci *USA* **106**, 6111 (2009).