

Towards Short-TE MR Spectroscopic Imaging: Spectral Decomposition and Removal of Baseline Signals

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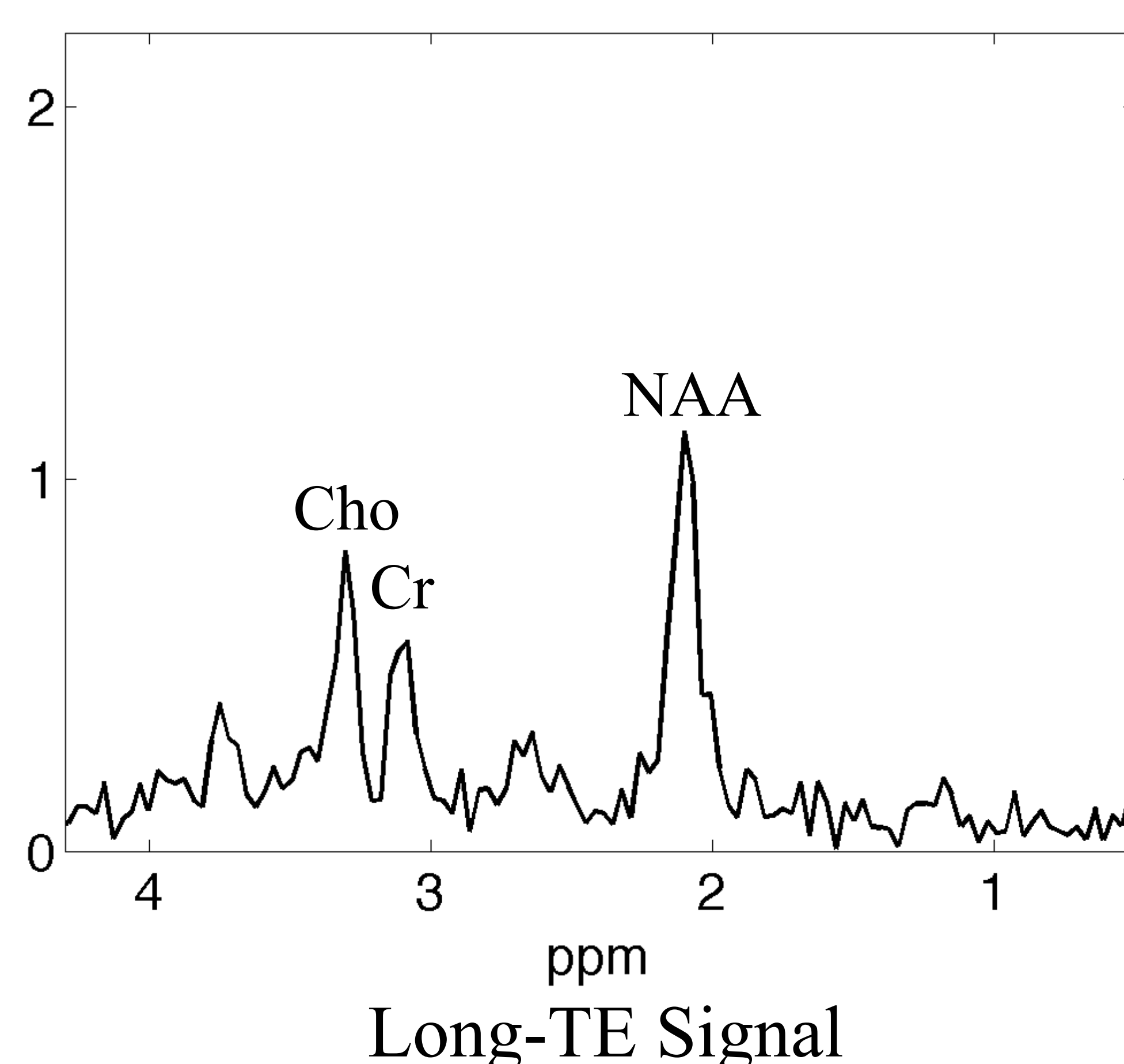
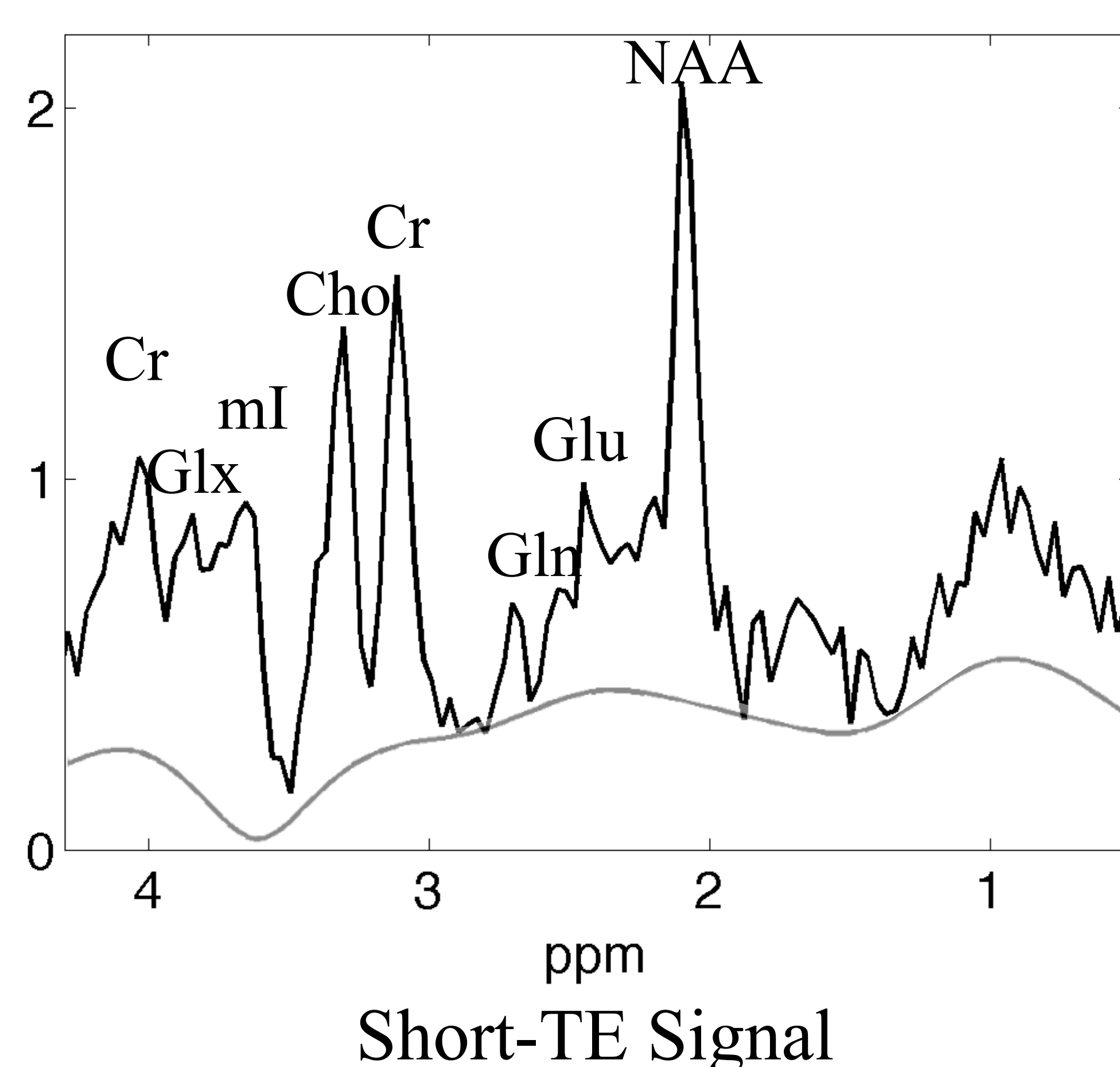
INTRODUCTION

Magnetic resonance spectroscopic imaging (MRSI)

- A powerful tool for mapping metabolite levels in vivo
- Challenge: very low SNR

Short echo-time (TE) MRSI

- ✓ High intrinsic SNR
- ✓ Rich metabolic information
- ✗ Significant baseline signals
- ✗ Difficult quantification problem



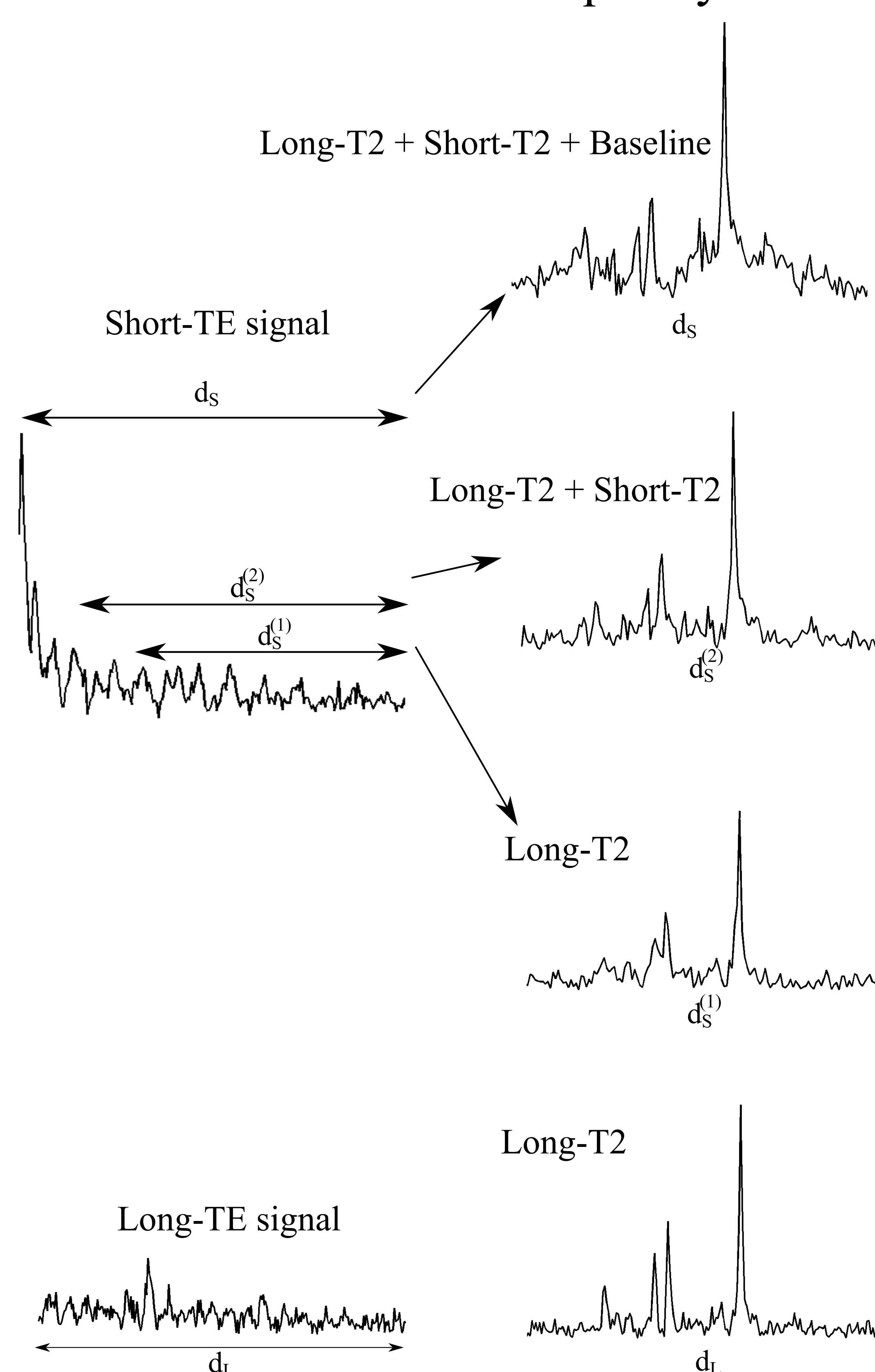
To address these issues, we exploit the **temporal separability** of short-TE MRSI signals to decompose them into three components

- *Long- T_2 metabolites*
- *Short- T_2 metabolites*
- *Baselines*

PROPOSED METHOD

Temporal Separability

Time Domain Frequency Domain



Signal model of MRSI data

$$d(t) = \sum_{n=1}^N c_n e^{-\frac{t}{T_{2,n}}} e^{-i2\pi f_n(t-T_E)} g(t-T_E) + \xi(t)$$

- $g(\cdot)$ -- signal decay caused by field inhomogeneity

Proposed method

- Estimate *long- T_2 metabolites* using the variable projection method^[1]

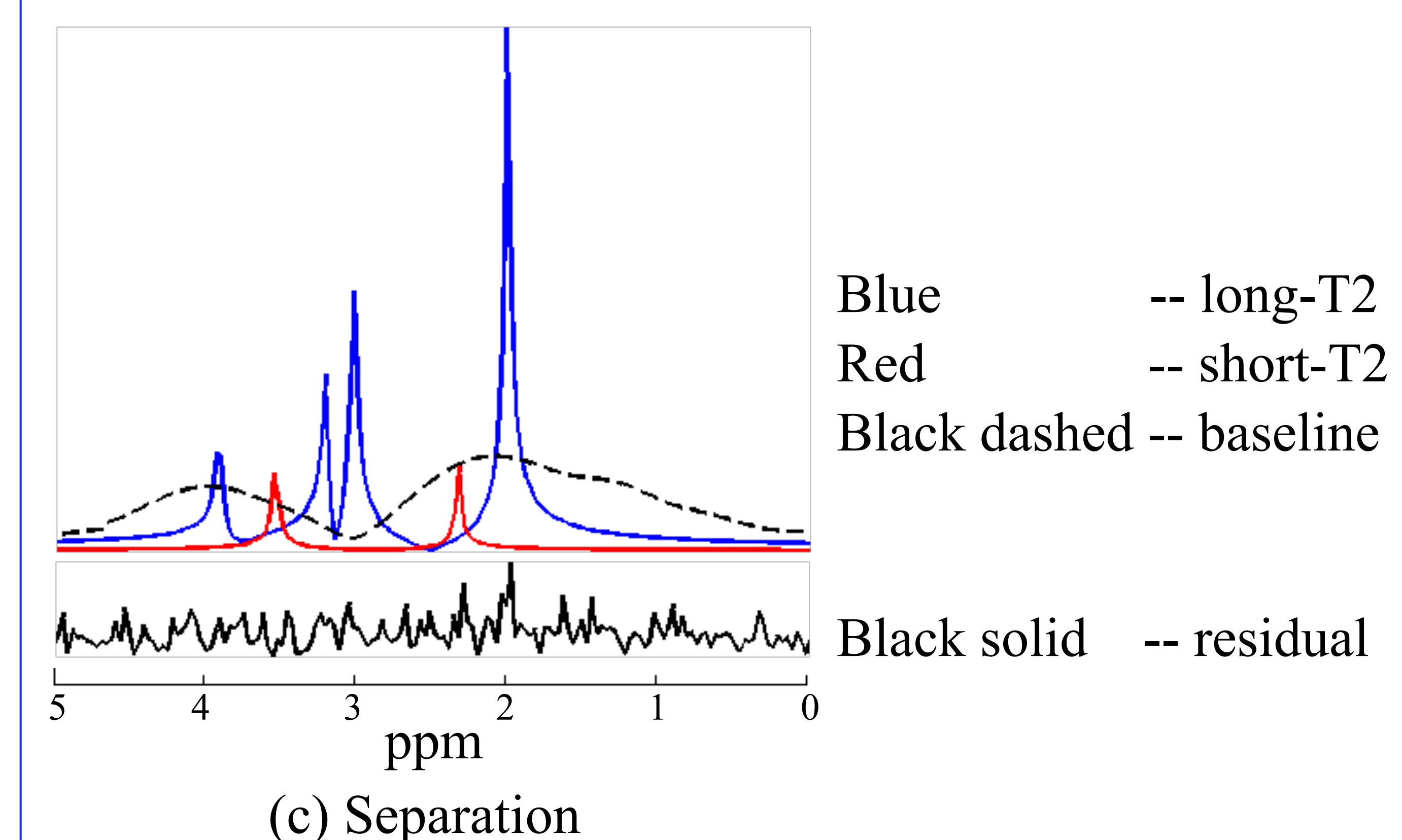
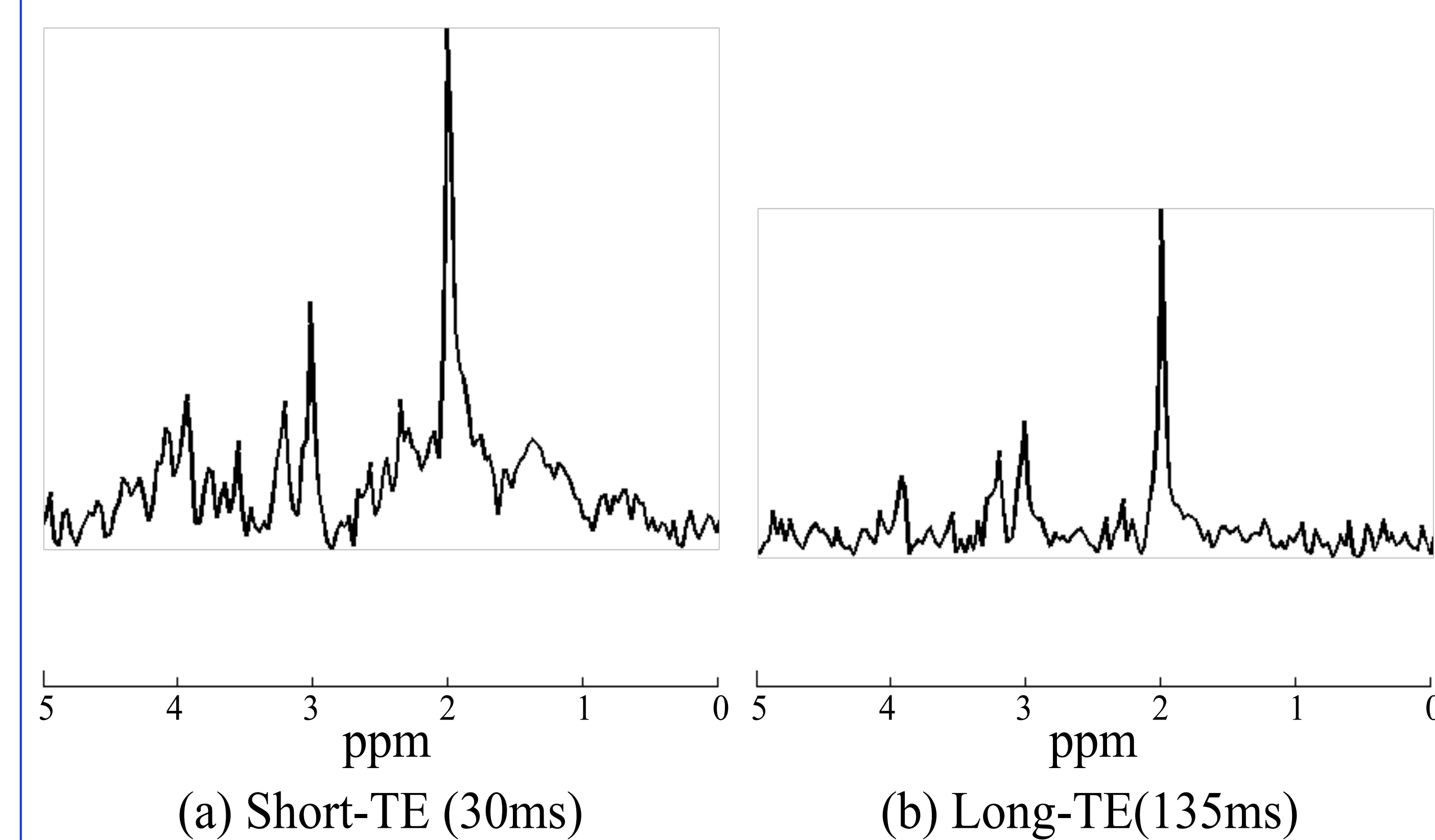
$$\min_{\alpha_L, c_L} \left\| \begin{bmatrix} d_S^{(1)} \\ d_L \end{bmatrix} - \begin{bmatrix} \Phi_S^{(1)}(\alpha_L) \\ \Phi_L(\alpha_L) \end{bmatrix} c_L \right\|_2$$

- Estimate *short- T_2 metabolites* using HSVD^[2]
- Estimate *baselines* using smooth splines fitting

RESULTS

In vivo human brain data

- $TE_1=30$ ms, $TE_2=135$ ms
- Water and lipid signals are already removed



CONCLUSION

A novel method has been proposed for processing short-TE MRSI data, which may improve the practical utility of short-TE MRSI experiments.

REFERENCE

- [1] G. H. Golub, *et al.*, SIAM J Numer Anal 1973; 10 (2): 413-432.
- [2] H. Barkhuijsen, *et al.*, J Magn Reson 1987; 73: 553-557.