



# Analytical tool for *in vivo* magnetic resonance signals

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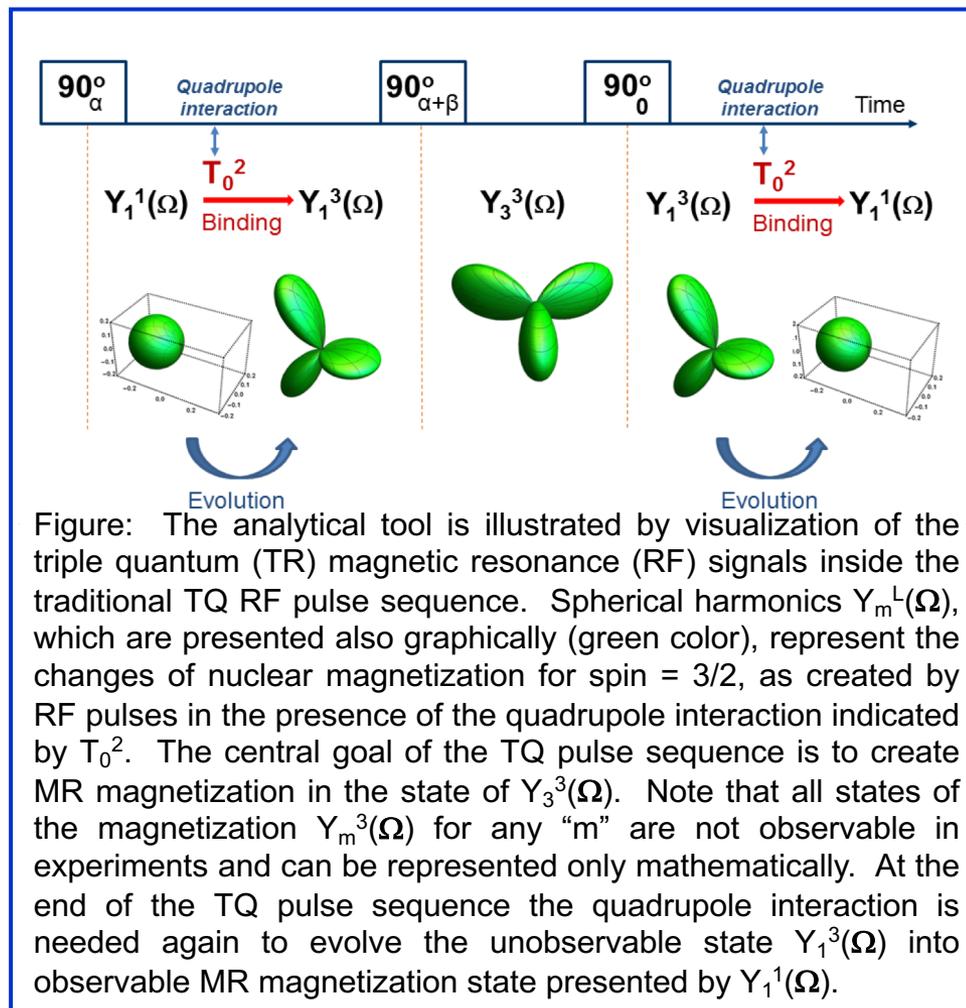
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Many scientists, including MagLab users, widely utilize triple quantum (TQ) magnetic resonance (MR) signals in their research. For *in vivo* experiments, such signals come mainly from sodium and potassium ions which interact with negatively charged groups of macromolecules, mainly proteins. MR signals appearing as a result of protein interactions indicate changes of intracellular ion content in vivo during diseases and drug administration. These signals can be efficiently extracted from other ions by using TQ effects. Note that the behavior of the MR magnetization in the presence of spin  $>1/2$  is not possible to explain by the simple vector model routinely used for proton MR.

The computer based tool is presented to describe and visualize the results of MR signal evolution *in vivo* in multi-pulse MR experiments. The system of theoretical equations describes MR signal evolution exactly without any approximation and needs to be constructed only once, which is then valid for any interval in the pulse sequence as well as for any spin value. The theoretical calculations are illustrated using the power of “Mathematica” software (Wolfram Inc.) for traditional TQ radio frequency (RF) pulse sequence (see Figure). The results of calculations can be visualized, noting that MR magnetization must remain unchanged during a parity transformation, i.e. changing the sign of all coordinates.



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