

In vivo Chloride Quantification with Partial Volume Corrected ^{35}Cl -MRI

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TARGET AUDIENCE Scientists and physicians interested in the field of non-proton MRI

PURPOSE Chloride plays a key role in many physiological processes [1]. Thus, a non-invasive determination of the chloride concentration by ^{35}Cl -MRI is desirable. However, the ^{35}Cl -nucleus experiences extremely fast transverse relaxation [2] and an *in vivo* signal which is reduced by six orders of magnitude compared to protons (^1H). Nevertheless, the feasibility of *in vivo* ^{35}Cl -MRI of human muscle and brain has recently been demonstrated [2]. For ^{35}Cl -MRI pulse sequences that enable ultra-short echo-times and high SNR efficiency such as 3D density adapted radial or twisted projection imaging [3] are necessary. Still, only resolutions of $(>6\text{mm})^3$ are achievable within acceptable acquisition times. Additionally, the applied acquisition schemes and fast T_2^* -relaxation lead to large full widths at half maximums (FWHM) of the *point spread functions* (PSF). The large voxel dimensions and the additional broadening of the PSF result in partial volume (PV) effects that decrease the accuracy of quantitative concentration measurements. For sodium (^{23}Na) MRI a partial volume correction (PVC) algorithm, based on a method for positron emission tomography by Rousset et al. [4], was presented that enabled improved quantification [5]. This method was now applied to *in vivo* ^{35}Cl -MRI.

METHODS A phantom was used to test the correction and quantification performance of the PVC. A silicone-caoutchouc cushion filled with 2% w/w agarose gel and different NaCl-concentrations (25-150 mmol/L) [6] was used for quantification of the signal (Fig 1.B). Imaging was conducted on a 7T MR system (Magnetom 7T, Siemens AG, Healthcare Sector, Erlangen, Germany), where image acquisition was performed with a density adapted projection pulse sequence (3D-DAPR [7]). For phantom imaging the following parameters were applied: TR/TE=150ms/0.5ms, 10000 projections, $\Theta=90^\circ$, nominal resolution: $(6\text{mm})^3$; $T_{\text{Acq}}=25\text{min}$ (Fig. 1.B). Additional data for a phase sensitive B_1 -correction [8] and a separate B_0 -correction were obtained.

In vivo quantification: PVC as described in the previous ^{23}Na -MRI approach [5] with relaxation weighting of the PSF (brain matter (BM): $T_{2s}^*=1.2\text{ ms}$, $T_{2l}^*=7\text{ ms}$ and $T_1=9.2\text{ ms}$ [2]; CSF: $T_2^*=T_1=35\text{ms}$) was applied to ^{35}Cl -data sets of three healthy volunteers with two experiments each. Data sets were acquired with a 3D-DAPR pulse sequence (TR/TE=125ms/0.4ms, 6000 projections, $(6.5\text{mm})^3$, $\Theta=90^\circ$, $T_{\text{Acq}}=12\text{min } 30\text{s}$, Fig. 2.A) with additional correction of B_1 - and B_0 -inhomogeneity, as in the phantom study. The chloride tissue concentration (CTC) was calculated for two separate CSF compartments (lateral ventricles (CSF_l) and sulci (CSF_s), same expected CTC) and for BM. Signal of the reference cushion (Fig. 2.B) was also PV-corrected to prevent underestimation of the calibration. Structural information of the cushion was obtained with a proton 3D-GRE (TR/TE=8.1ms/4.88ms, $\Theta=10^\circ$, $(1\text{mm})^3$) and segmented manually. Correction behavior of the two CSF compartments with the calculated difference ΔCSF between CSF_l and CSF_s was used as an intrinsic correction control.

RESULTS Performance of phantom PVC and calibration was directly verified with known ^{35}Cl -concentrations of eleven reference tubes (Fig.1.B). The magnitude of discrepancy of obtained to expected concentration values before and after correction was calculated and plotted (Fig. 1.A). A mean reduction of discrepancy of 16% is seen after correction. *In vivo* data showed a cutback of mean ΔCSF from 35.8% to 5.4% and an upward correction for both CSF values (Tab.1). Brain matter CTC was shifted downward from 32 mmol/L to 27 mmol/L (-15.6%).

DISCUSSION The PV effects were reduced by a PVC algorithm that allowed a strongly improved quantification for ^{35}Cl -MRI. For phantom measurements, heavy PV influence was seen (mean discrepancy 27%). True values were recovered, quantified and verified with known concentration values. Results of the phantom study showed correction capability of the algorithm and quantification capacity of the reference cushion. For *in vivo* measurements the intrinsic correction control indicates good correction behavior: CSF CTC values were shifted closer together and the theoretical ^{35}Cl -concentration for both CSF compartments was met (99-110 mmol/L). BM CTC was corrected downward as expected and was lower than previously reported values [2], where PV effects were not considered. However, the calculated CTC might be underestimated due to the fast transverse relaxation of ^{35}Cl in tissue.

CONCLUSION Correction capability of a previously introduced PVC algorithm for ^{23}Na -MRI [5] was demonstrated for ^{35}Cl -MRI where much stronger PV influence due to larger voxel sizes is expected. The correction and quantification approach allowed absolute ^{35}Cl -quantification in the human brain, where PV-bias was strongly reduced.

REFERENCES

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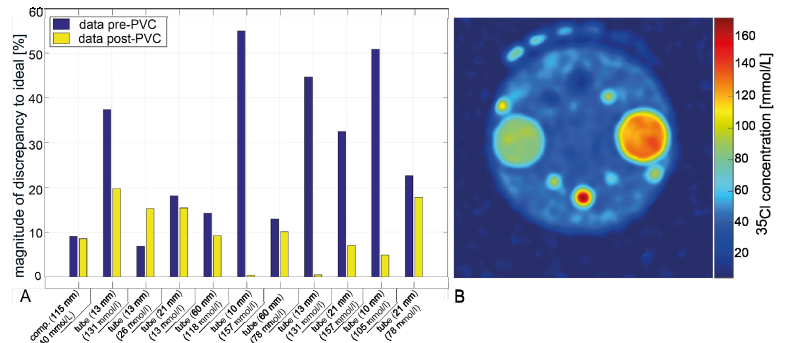


Fig.1: (A) Plotted magnitude of discrepancy for phantom data before (blue) and after correction (yellow), an improvement in quantification accuracy is seen for post-correction data. (B) Used ^{35}Cl -data $((6\text{mm})^3$, TR=150ms, TE=0.5ms) for quantification.

Tab. 1: Mean values \pm std. of CTC for *in vivo* experiment

	CSF _o [mmol/L]	CSF _i [mmol/L]	mean diff. ΔCSF [%]	BM [mmol/L]
Pre-PVC [mmol/L]	35 \pm 2	54 \pm 4	35.8 \pm 2.7	32 \pm 3
Post-PVC [mmol/L]	98 \pm 7	100 \pm 4	5.4 \pm 3.1	27 \pm 3

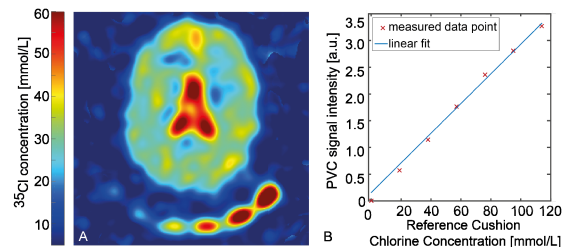


Fig. 2: (A) Used ^{35}Cl *in vivo* data $((6.5\text{mm})^3$, TR/TE=125ms/0.4m) with signal of reference cushion (zoomed FOV 228x228x228 mm). (B) result of fit (blue) with PV-corrected data of calibration cushion (red).