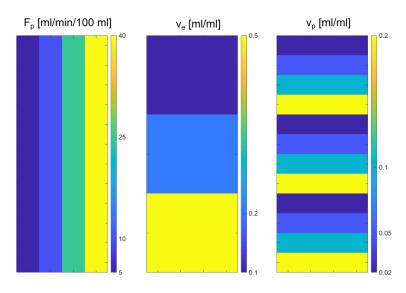
## dce\_2CXM\_ValidationDataSet

This is the documentation for the two compartment exchange model (2CXM) digital reference object (DRO). 144 concentration time curves with different parameter combinations of  $F_p$ , PS,  $v_p$  and  $v_e$  were simulated were simulated using the 2CompFlowExch model in JSIM (http://www.physiome.org/jsim/). Parameter values used for curve simulation were

$$F_p \in \{5, 10, 25, 40\} \frac{ml}{\min 100ml} \qquad v_p \in \{0.02, 0.05, 0.1, 0.2\}$$

$$PS \in \{0.0, 5, 15\} \frac{ml}{\min 100ml}$$
  $v_e \in \{0.1, 0.2, 0.5\}$ 

Temporal resolution is 0.5 s over a total of 661 time points. The arterial input function was taken from the QIBA\_v6\_Tofts DRO (<a href="https://sites.duke.edu/dblab/qibacontent/">https://sites.duke.edu/dblab/qibacontent/</a>)



Within the DRO, curves are arranged in a 3D+t image of spatial dimensions <40,120,3>, where each  $<10\times10\times1>$  block of voxels contains one curve type with a combination of the above listed parameter values . Within the 3D volume,  $F_p$  varies along the x-axis,  $v_e$  along the y-axis,  $v_p$  on a subset of the y-axis (in alternating form) and PS along the z-axis, as illustrated in the schematics.

The AIF was added as <40, 20, 3> block on the bottom of the image, leading to a final DCE MRI data set of dimension <40, 140, 3, 661>.

The 4D concentration image is converted to signal intensities via relative signal enhancement using a factor of k=1

$$C(t) = k \cdot \frac{S(t) - S_0(t)}{S_0(t)} \rightarrow S(t) = \frac{1}{k} \cdot C(t) \cdot S_0(t) + S_0(t)$$

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Debus C and Floca R, Ingrisch M, Kompan I, Maier-Hein K, Abdollahi A, Nolden M, MITK-ModelFit: generic open-source framework for model fits and their exploration in medical imaging - design, implementation and application on the example of DCE-MRI (arXiv:1807.07353)

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