The value of arterial spin labelling in adults glioma grading: systematic review and meta-analysis

SUPPLEMENTARY MATERIALS

 $rTBF$

Supplementary Figure 1: rTBF for LGG patients' relative to the value for HGG patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have very different distribution (heterogeneity) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the LGG have significantly lower rTBF than the HGG $(-1.46, (-2.00, -0.91))$. The funnel plot is symmetric and does not show publication bias.

rTBF mean

Supplementary Figure 2: rTBF-mean for LGG patients relative to the value for HGG patients. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have very different distribution (heterogeneity) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the LGG have significantly lower rTBFmean than the HGG $(-1.53, (-2.26, -0.79))$. The funnel plot is symmetric and does not show publication bias.

SMD

Study

Supplementary Figure 3: rTBF-max for LGG patients' relative to the value for HGG patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have very different distribution (heterogeneity) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the LGG have significantly lower rTBFmax than the HGG $(-1.36, (-2.23, -0.49))$. The funnel plot is symmetric and does not show publication bias.

TBF

Supplementary Figure 4: TBF for LGG patients' relative to the value for HGG patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have very different distribution (heterogeneity) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the LGG have significantly lower TBF than the HGG $(-0.82, (-1.20, -0.45))$. The funnel plot is symmetric and does not show publication bias.

Supplementary Figure 5: TBFmean for LGG patients' relative to the value for HGG patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have small distribution around this line with small degree of heterogeneity. The pooled effect and their 95% CI (the diamond at the bottom) express that the LGG have significantly lower TBFmean than the HGG (–0.61, (–0.99, –0.23)). The funnel plot is symmetric and does not show publication bias.

Supplementary Figure 6: TBFmax for LGG patients' relative to the value for HGG patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have very different distribution (heterogeneity) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the LGG have significantly lower TBFmax than the HGG $(-0.96, (-1.53, -0.39))$. The funnel plot is symmetric and does not show publication bias.

 $rTBF$

Supplementary Figure 7: rTBF for grade-II patients' relative to the value for grade-III patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have low distribution (small heterogeneity degree) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the grade-II rTBF value about significantly lower than the that of the grade-III $(-1.39, (-1.89, -0.89))$. The funnel plot is symmetric and does not show publication bias.

Supplementary Figure 8: TBF for grade-II patients' relative to the value for grade-III patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have moderate distribution (moderate heterogeneity degree) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the grade-II has approximately significant lower TBF value than the grade-III (–0.90, (–1.85, 0.04)). The funnel plot cannot be produced due to the small study number.

Supplementary Figure 9: rTBF for grade-II patients' relative to the value for grade-IV patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have very large distribution (heterogeneity) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the grade-II rTBF value was significantly lower than the that of the grade-IV $(-2.07, (-3.38, -0.76))$. The funnel plot is symmetric and does not show publication bias.

Supplementary Figure 10: TBF for grade-II patients' relative to the value for grade-IV patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies represent very large distribution (heterogeneity) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the grade-II significantly lower TBF value than the grade-IV (-1.44, (-2.76, -0.12)). The funnel plot cannot be produced due to the small study number.

Supplementary Figure 11: rTBF for grade-III patients' relative to the value for grade-IV patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have large distribution (heterogeneity) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express that the grade-III significantly has lower rTBF value than the grade-IV $(-1.05, (-1.82, -0.27))$. The funnel plot is symmetric and does not show publication bias.

Supplementary Figure 12: TBF for grade-III patients' relative to the value for grade-IV patients'. In the forest plot, the dotted vertical line represents the pooled effect size point where the effect size in individual studies have low distribution (small heterogeneity degree) around this line. The pooled effect and their 95% CI (the diamond at the bottom) express a trend of lower rTBF value in grade-III than in grade-IV (–0.45, (–0.95, 0.05)). The funnel plot is asymmetric and does show publication bias.

Supplementary Table 1: Studies performed using PCASL

(*)study 18 did not mentioned the used ASL labelling method. Not available (NA).

Supplementary Table 2: Studies performed using CASL

Not available (NA).

Supplementary Table 3: Studies performed using PASL

Not available (NA); Proximal Inversion with Control of Off-Resonance Effects (PICORE); flow alternating inversion recovery (FAIR); quantitative STAR labeling of arterial regions (QUASAR).

Supplementary Table 4: Sensitivity, specificity, negative predictive values (NPV) and positive predictive values (PPV) of published ASL-derived biomarkers cut-off values for glioma grading

| HGG vs LGG | | | | | | | | |
|--|---|----------------------|---------|--------------------|--------------|------------|------------|------------|
| Author/vear | Study No. | ASL parameter | Cut-off | Sensitivity | Specificity | prevalence | PPV | NPV |
| Kim, H.S. et al.; 2007 | 5 | rTBFmean | 1.24 | 0.955 | 0.818 | 0.667 | 91.30 | 90.088 |
| Fudaba, H. et al.; 2014 Shen, N. et al.; 2016 | τ | rTBFmean | 2.562 | 0.652 | 0.778 | 0.719 | 88.243 | 46.661 |
| | τ | rTBFmax | 2.845 | 0.609 | 0.778 | 0.719 | 87.516 | 43.776 |
| | τ | rTBFmin | 2.017 | 0.739 | 0.667 | 0.719 | 85.0105 | 50 |
| | 7, (astrocytoma) | rTBFmean | 1.8 | 0.824 | 0.667 | 0.85 | 93.343 | 40.076 |
| | 7, (astrocytoma) | rTBFmax | 2.258 | 0.765 | 0.667 | 0.85 | 92.866 | 33.372 |
| | 7, (astrocytoma) | rTBFmin | 1.254 | 0.882 | 0.667 | 0.85 | 93.753 | 49.937 |
| | 9 | TBFmax | 52.21 | 0.889 | 0.826 | 0.5192 | 84.664 | 87.317 |
| | 9 | rTBFmax | 1.32 | 0.926 | 0.957 | 0.519 | 95.831 | 92.279 |
| Yang, X. et al.; 2016 | 11, (multiple TIs), (astrocytoma) | rTBFmean | 2.43 | $\mathbf{1}$ | 0.54 | 0.6511 | 80.229 | 100 |
| | 11, (single TI), (astrocytoma) | rTBFmean | 3.01 | 0.6 | 0.88 | 0.651 | 90.323 | 54.098 |
| | 11, (bolus arrival time (BAT)), (astrocytoma) | | 0.97 | 0.71 | 0.88 | 0.651 | 91.697 | 61.914 |
| Furtner, J. et al.; 2014 | 12 | | 1.48 | 0.85 | $\mathbf{1}$ | 0.788 | 100 | 64.220 |
| Cebeci, H. et al.; 2014 | 13 | rTBFmax | 2.1 | $\mathbf{1}$ | 0.92 | 0.606 | 95.057 | 100 |
| | 13 | rSImax | 2.19 | $\mathbf{1}$ | 0.92 | 0.606 | 95.057 | 100 |
| Kim, M J. et al.; 2008 | 14, (astrocytoma) | rTBFmax | 1.28 | 0.829 | 0.962 | 0.5738 | 96.707 | 80.691 |
| Canale, S. et al.; 2011 | 16, (oligodendroglioma) | rTBFmean | 1.8 | 0.88 | 0.6 | 0.762 | 87.562 | 60.976 |

Supplementary Table 5: Sensitivity, specificity, negative predictive values (NPV) and positive predictive values (PPV) of published ASL-derived biomarkers cut-off values between HGGs and LGGs

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