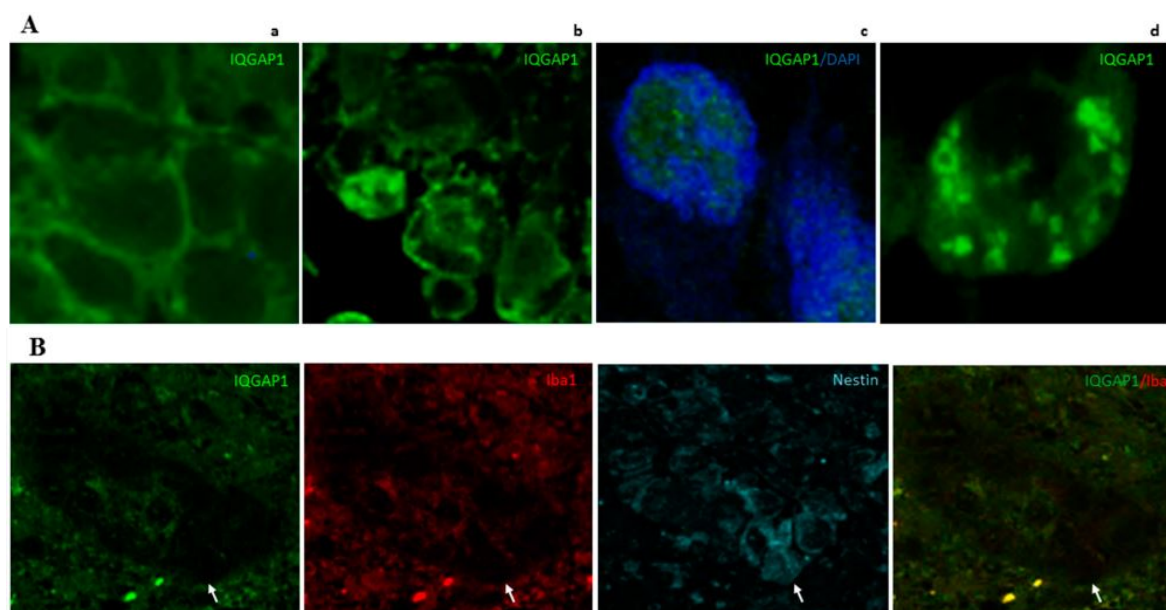


## Supplementary Materials: IQGAP1 in Podosomes/Invadosomes is Involved in the Progression of Glioblastoma Multiforme Depending on the Tumor Status

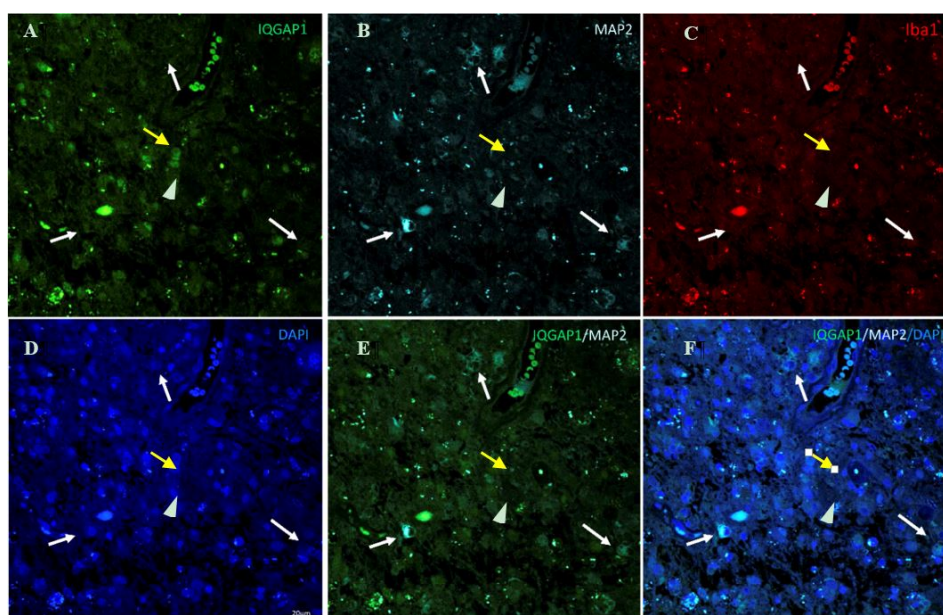
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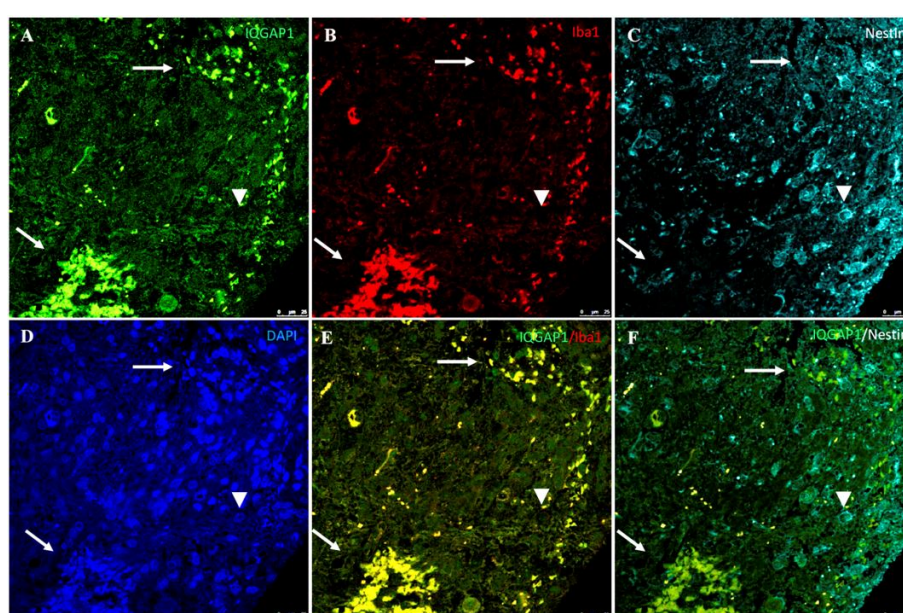
**Supplemental Figure S1.**

**A:** Higher magnification of panels A-D of Figure 1. IQGAP1 protein in GBM detected in (a) plasma membrane; (b) cell protrusions, plasma membrane and cytosol; (c) nucleus. (d) cell with podosome-like structures highly positive for IQGAP1.

**B:** Higher magnification of panels K-O of Figure 2. Triple immunolocalization of IQGAP1 protein (green), Iba1 (red) and nestin (cyan). Arrow points to a nestin<sup>+</sup>/IQGAP1<sup>-</sup>/Iba1<sup>-</sup> cell flanked by nestin<sup>+</sup>/IQGAP1<sup>+</sup>/Iba1<sup>+</sup> cells.



**Figure S2.** Triple immunolocalization of IQGAP1 protein (green), the microglia/macrophage marker Iba1 (red) and the Microtubule-associated protein 2 (MAP2, cyan) in GBM tissue sections. White arrows point to Map2<sup>+</sup> neurons IQGAP1<sup>+</sup>/Iba1<sup>-</sup>. Yellow arrows points to a cell IQGAP1<sup>+</sup>/Iba1<sup>+</sup>. Arrowhead points a cell IQGAP1<sup>+</sup>/nestin<sup>+</sup>. Bar: 20µm.



**Figure S3.** Triple immunolocalization of IQGAP1 protein (green), the microglia/macrophage marker Iba1 (red) and the CSC marker nestin (cyan). In this highly vascularized area of a GBM tissue section many nestin<sup>+</sup> cells are observed (arrowhead). Note the massive presence of macrophages (Iba1<sup>+</sup>/IQGAP1<sup>+</sup>/nestin<sup>+</sup>) in the periphery (arrows). Scale bar: 25µm.

**Table S1.** IQGAP1/PCNA and IQGAP1/GFAP colocalization. Mean and standard deviation of values obtained with the ImageJ plug-in “Manders’ coefficients”.

	<i>n</i>	R	Ch1/Ch2	M1	M2	Ch1 Thresh, Ch2 Thresh
Ch1 = GFAP Ch2 = IQGAP1	27	0.59 ± 0.05	0.995 ± 0.02	0.969 ± 0.02	0.971 ± 0.01	1;255
Ch1 = PCNA Ch2 = IQGAP1	20	0.677 ± 0.13	0.750 ± 0.29	0.851 ± 0.14	0.739 ± 0.20	1;255

<b>Ch1 = Iba1</b>						
<b>Ch2 = IQGAP1</b>	14	0.84 ± 0.04	0.801 ± 0.15	0.954 ± 0.04	0.835 ± 0.09	1;255

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*n* = samples analyzed. **R** = Manders Overlap coefficient. Value Range: 0–1, with 0 = low colocalization; 1 = high colocalization; **Ch1/Ch2** = red/ green pixel ratio. Ratio ≈ 1 allows the use of Mander's Overlap coefficient (R); **M1, M2** = Manders Colocalization coefficients for channel 1 (M1) and channel 2 (M2). Value range: 0–1, with 0 = no colocalization 1 = all pixels colocalize; **Ch1 thresh, Ch2 thresh** = ch1 and ch2 threshold.