

1 **Title**

2 Intravital imaging of glioma border morphology reveals distinctive cellular dynamics and
3 contribution to tumor cell invasion

4

5 **Authors**

6 Maria Alieva ^{1,2*#}, Verena Leidgens ^{1,3#}, Markus J. Riemenschneider ⁴, Christoph A. Klein ⁵,

7 Peter Hau ^{3*\$} and Jacco van Rheenen ^{1,6\$}

8

1 **Supplementary Figures**

2 **Supplementary Fig 1 Tumor infiltration patterns in human glioblastoma samples.**

3 Right column shows glioblastoma with typically *diffuse cell infiltration* in the surrounding
4 brain tissue (right column). Note that there is a decreasing gradient in cell density from the
5 upper right to the lower left of the images. Captured is a cortical infiltration zone (**CI**) with
6 interspersed neurons. GFAP staining in the illustrated case hardly differentiates between
7 infiltrative tumor cells and reactive astrocytes. A relevant number of tumor nuclei stain for the
8 p53 tumor suppressor protein. Single tumor cell infiltration in the lower left of the image (**CI**)
9 is highlighted by staining with the proliferation marker Ki-67 (MIB-1). Left column shows a
10 tumor with a more circumscribed growth pattern with an appearance resembling the *well-*
11 *defined tumor boarder* pattern in the model (red arrow). In the illustrated case solid tumor
12 tissue (**T**) and surrounding non-neoplastic brain tissue (**B**) can be clearly differentiated. This is
13 obvious also in the immunohistochemical stains with reactive GFAP expression and virtually
14 absent p53 and MIB-1 labeling in the brain tissue (**B**). Note that the MIB-1 stain, however,
15 highlights finger-shaped protrusions (black arrow) into the surrounding non-neoplastic tissue
16 corresponding to the *invasive margin* pattern in the model. The finger-shaped protrusions can
17 be also identified in the corresponding H&E image.

18

19 **Supplementary Fig 2 Velocity of different border configurations in individual mice.**

20 Quantification of cell velocity at indicated border and tumor core configurations. Data from
21 individual mice and BTIC-10 and -12 is plotted in separate graphs. The data is shown as mean
22 \pm S.E.M.

23

24 **Supplementary Fig 3 Intraparenchymal and invasive tumor margin.**

1 Representative 3D reconstructed tile-scan showing perivascular and intraparenchymal *invasive*
2 *margin*. Shown are H2B expressing BTICs in green, collagen fibers in blue, blood vessels in
3 red. Scale bar = 300 μm .

4

5 **Supplementary Fig 4 Comparison between the intraparenchymal and invasive tumor**
6 **margin.**

7 (A) Quantification of cell velocity, percentage of motile (cell velocity > 2 $\mu\text{m}/\text{hour}$) and static
8 cells, speed of motile cells and persistence of motile cells for the the intraparenchymal and
9 invasive tumor margin. Data is shown as mean \pm S.E.M., n = 4 mice (BTIC-10 and BTIC-12
10 lines). The data is shown as mean \pm S.E.M, n = 4 mice (BTIC-10 and BTIC-12 lines).
11 * $p < 0.05$, ** $p < 0.01$, *** $p < 0.0001$, one-way ANOVA with Tukey's post hoc test. B)
12 Representative still images from a time-lapse movie showing migrating tumor cells from
13 perivascular and intraparenchymal *invasive margin*. White lines highlight individual tumor cell
14 tracks. Scale bar = 30 μm . (C) Percentage of invading and receding cells for each condition.
15 (D) Quantification of the Y-axes displacement for invading and receding cells of each
16 condition. n = 7 (BTIC-10 and BTIC-12). (E) Quantification of direction correlation between
17 perivascular and intraparenchymal *invasive margin*. n = 4. ^{ns} $p > 0.05$. One sample Student's *t*,
18 hypothetical value = 0.

19

20 **Supplementary Fig 5 Direction of cell migration at different positions.**

21 Representative wind rose plots showing distinct direction of cell migration at individual
22 positions from the different border configurations. Direction of a leaflet represents the
23 percentage of cells migrating in that direction. Color scale indicates migration speed.

24

1 **Supplementary Fig 6 Linear regression model for cell spreading into the parenchyma.**

2 (A) Linear regression model for cell spreading with the corresponding coefficients for each
3 border configuration type. Variables are explained in Table2. (B) Histograms showing the
4 frequency distribution of the incidence of invasion within a position for the *invasive margin*
5 and the *diffuse margin*.

6

7 **Supplementary Fig 7 Intracranial imaging window**

8 Descriptive image showing the cranial imaging window in the living mouse in overview and
9 magnification.

10

11 **Supplementary tables**

12 **Supplementary Table 1 Summary of Mixed-effect Linear Regression Model for cell**
13 **spreading perpendicular to the tumor border**

14

15 **Supplementary Table 2 List of simulation parameters**

16

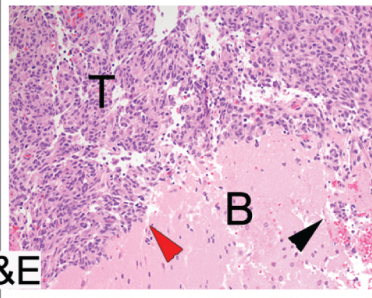
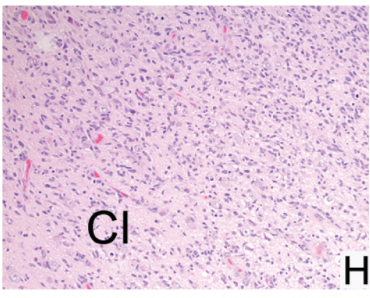
17 **Supplementary movie**

18 **Supplementary Movie 1 Invasive front cell dynamics.**

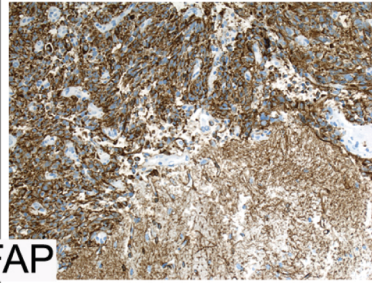
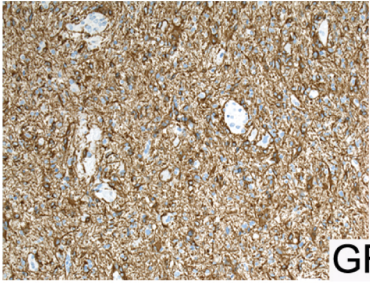
19 Intravital imaging video showing BTIC-10 H2B-Dendra2 cells migrating in opposite directions
20 at the *invasive margin*. Colored arrows follow individual cells of the *invasive margin*. Top:
21 tumor core; bottom: brain parenchyma. Scale bar represents 30 μm .

diffuse cell infiltration

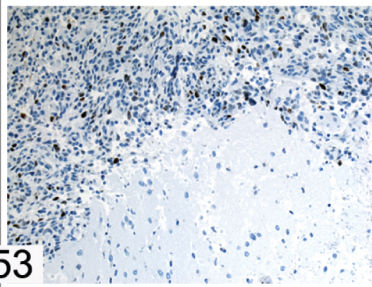
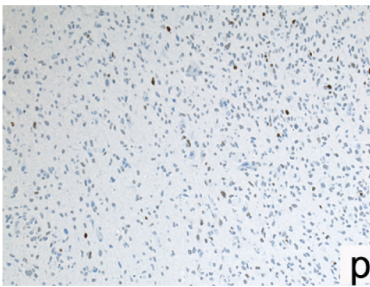
well-defined tumor boarder
invasive margin



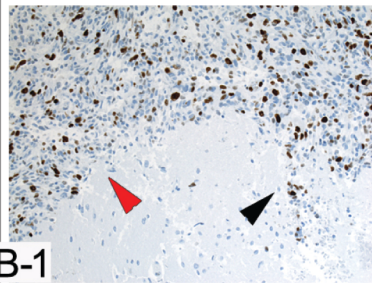
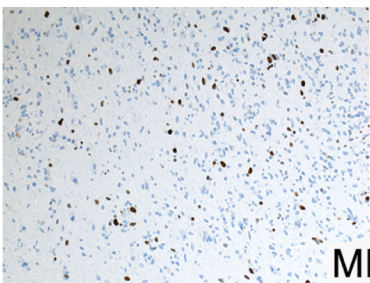
H&E



GFAP

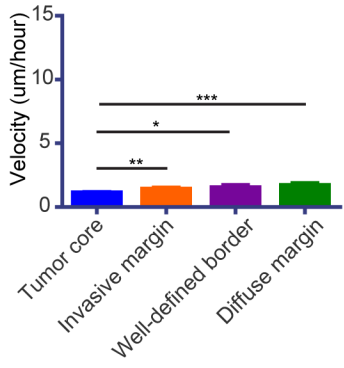


p53

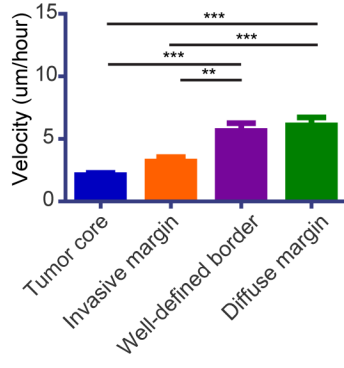


MIB-1

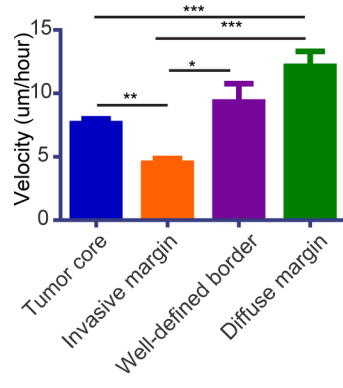
Mouse 1 BTIC 12



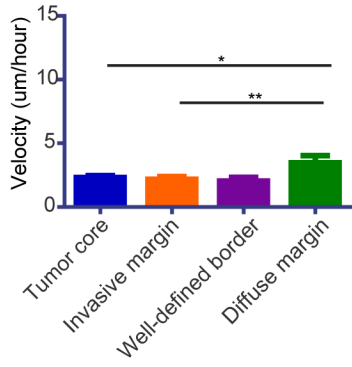
Mouse 2 BTIC 12



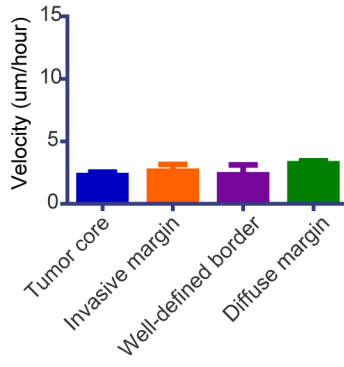
Mouse 3 BTIC 12



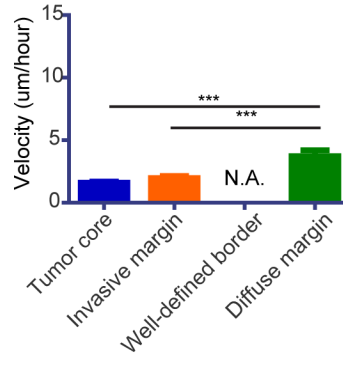
Mouse 4 BTIC 10



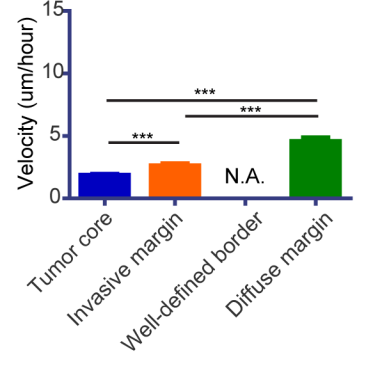
Mouse 5 BTIC 10



Mouse 6 BTIC 10



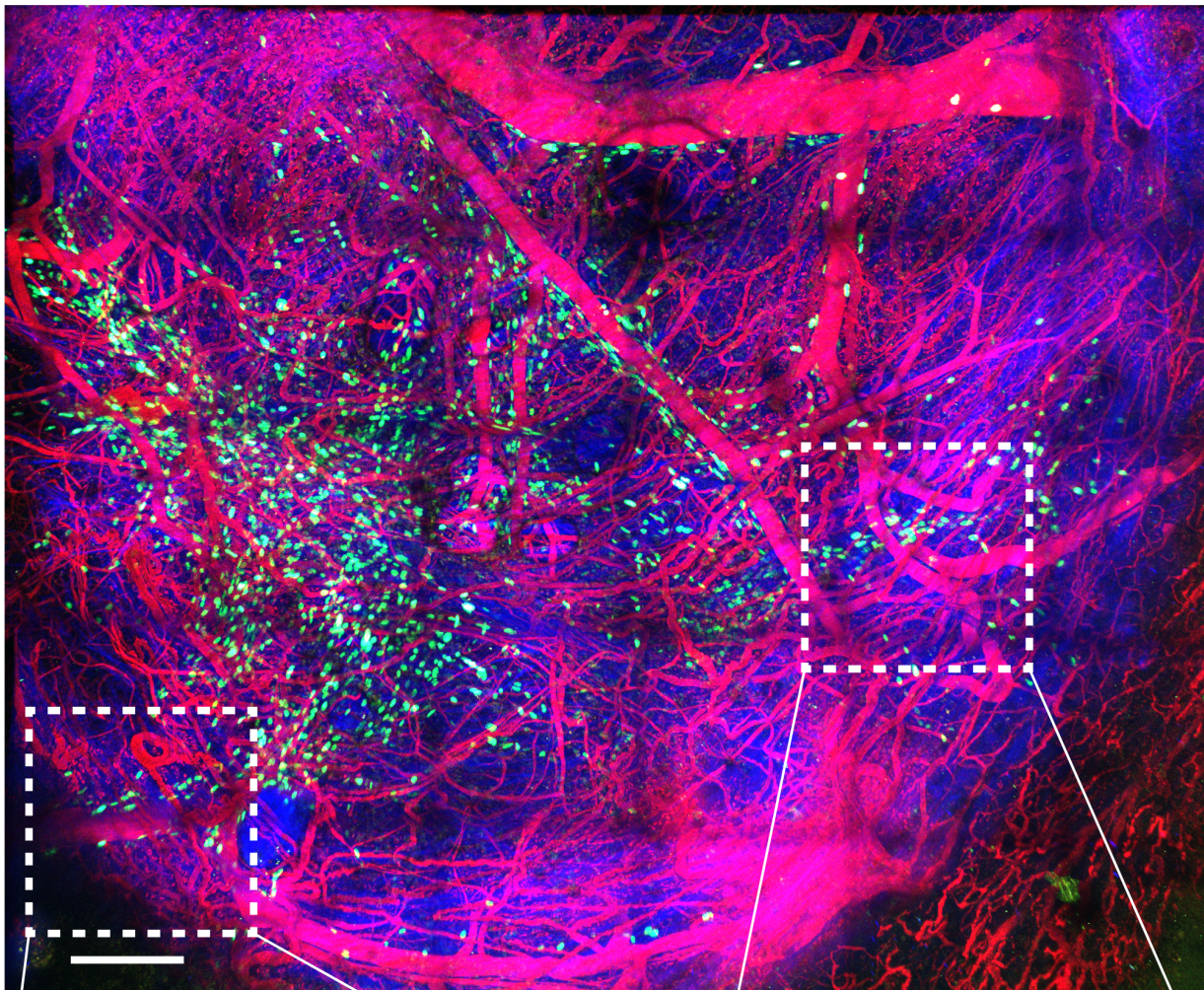
Mouse 7 BTIC 10



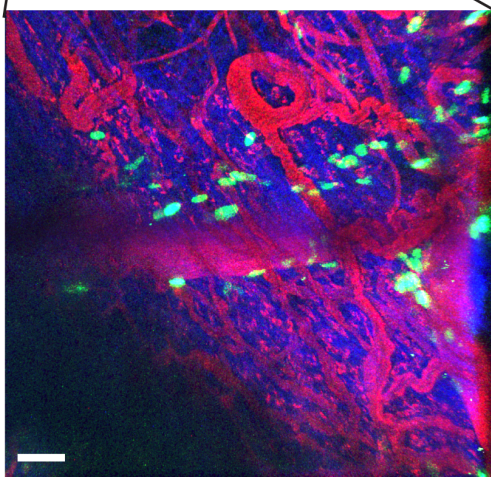
H2B Dendra2 BTIC-10 glioma cells

Collagen fibers

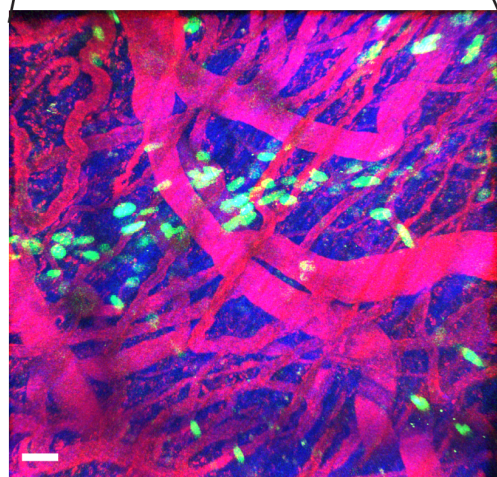
Blood vessels



300um

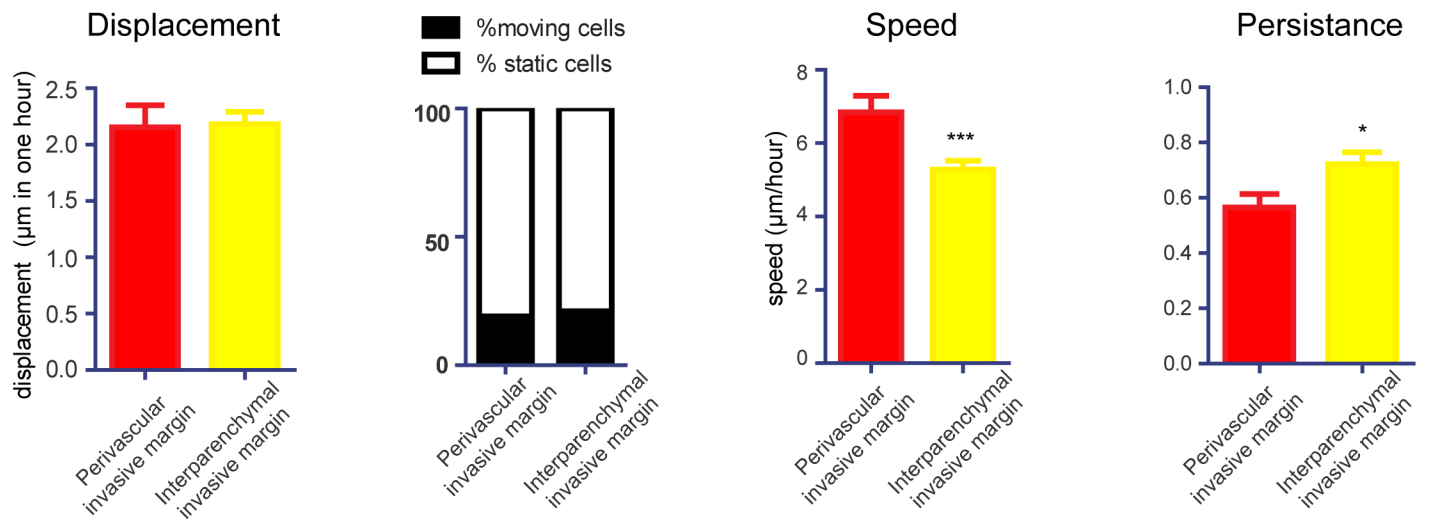


Perivascular invasive margin

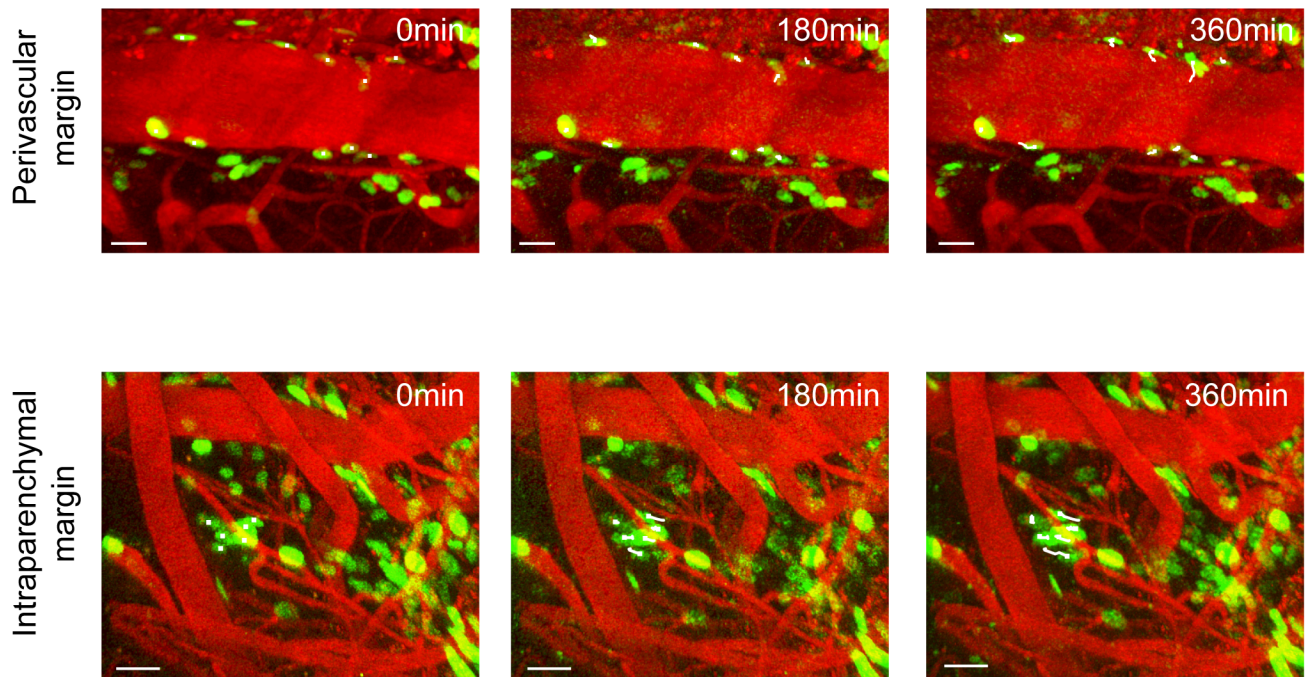


Intraparenchymal invasive margin

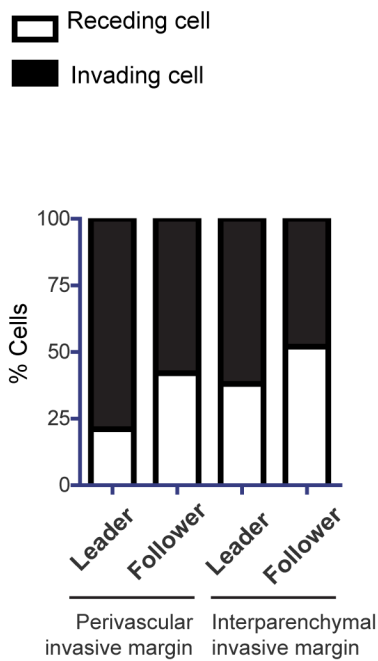
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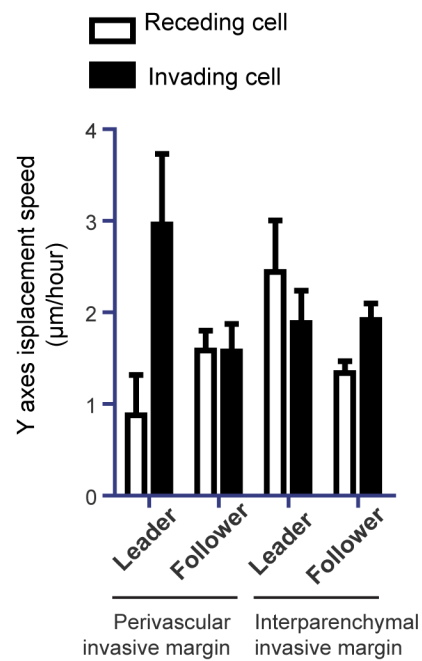
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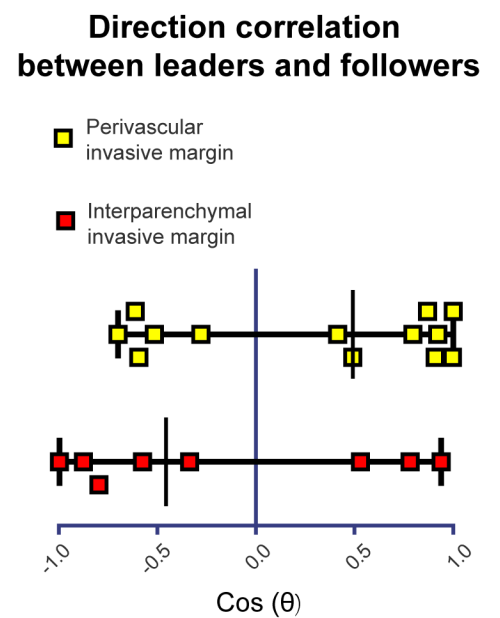
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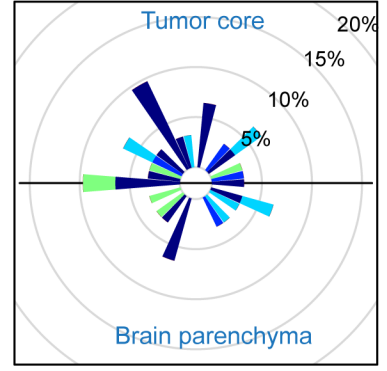
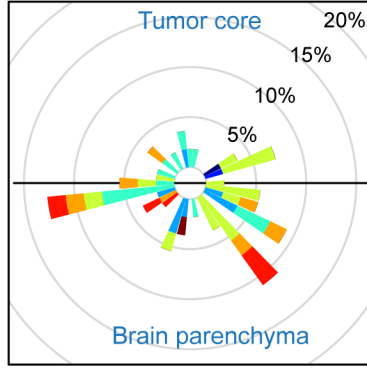
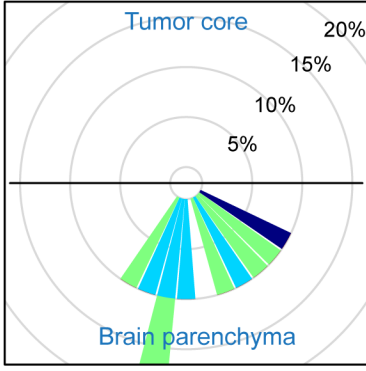
d



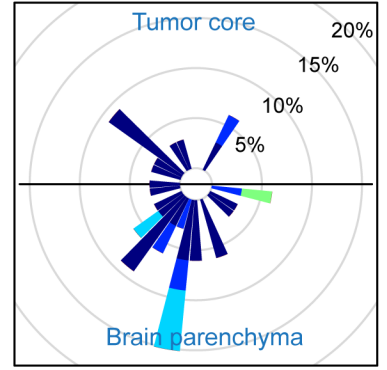
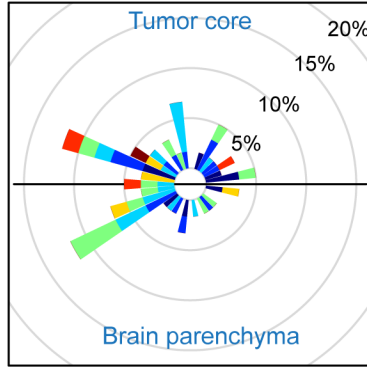
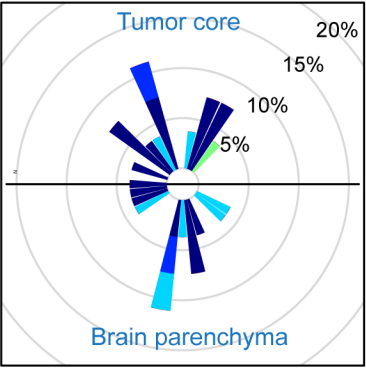
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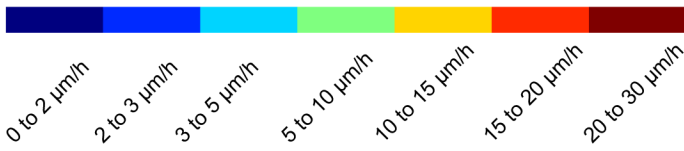
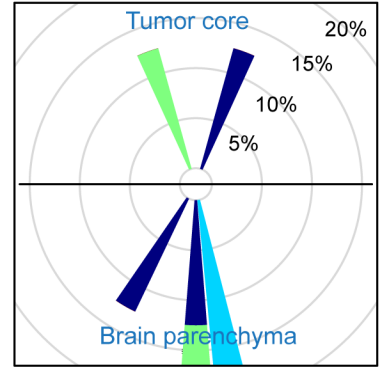
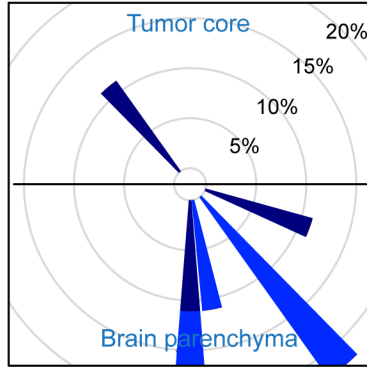
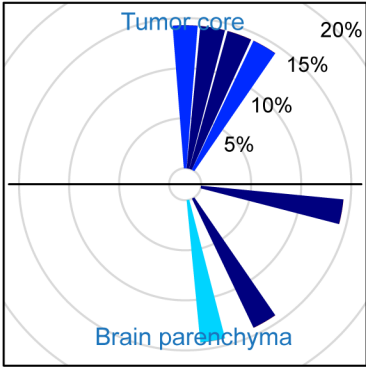
Diffuse margin



Well-defined border



Invasive margin



a

$$\text{COM } y_{\text{border}} = 0.36 + 0.86V_{\text{inv}} - 1.69(V_{\text{inv}} \times F_{\text{inv}})$$

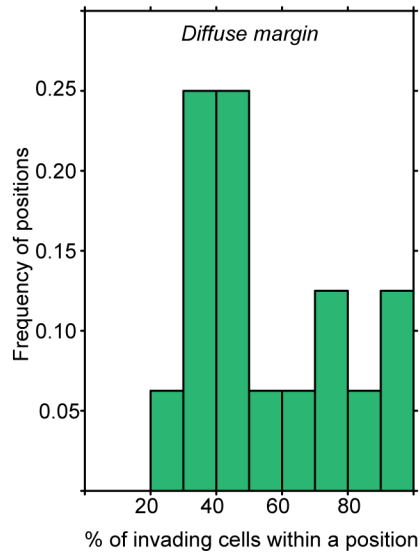
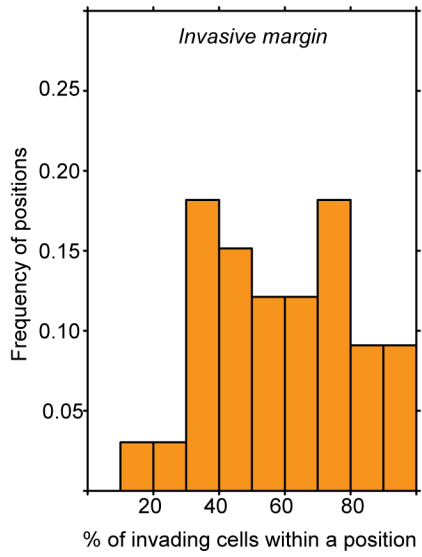
$$\text{COM } y_{\text{invasive}} = 0.36 + 0.62V_{\text{inv}} - 1.69(V_{\text{inv}} \times F_{\text{inv}})$$

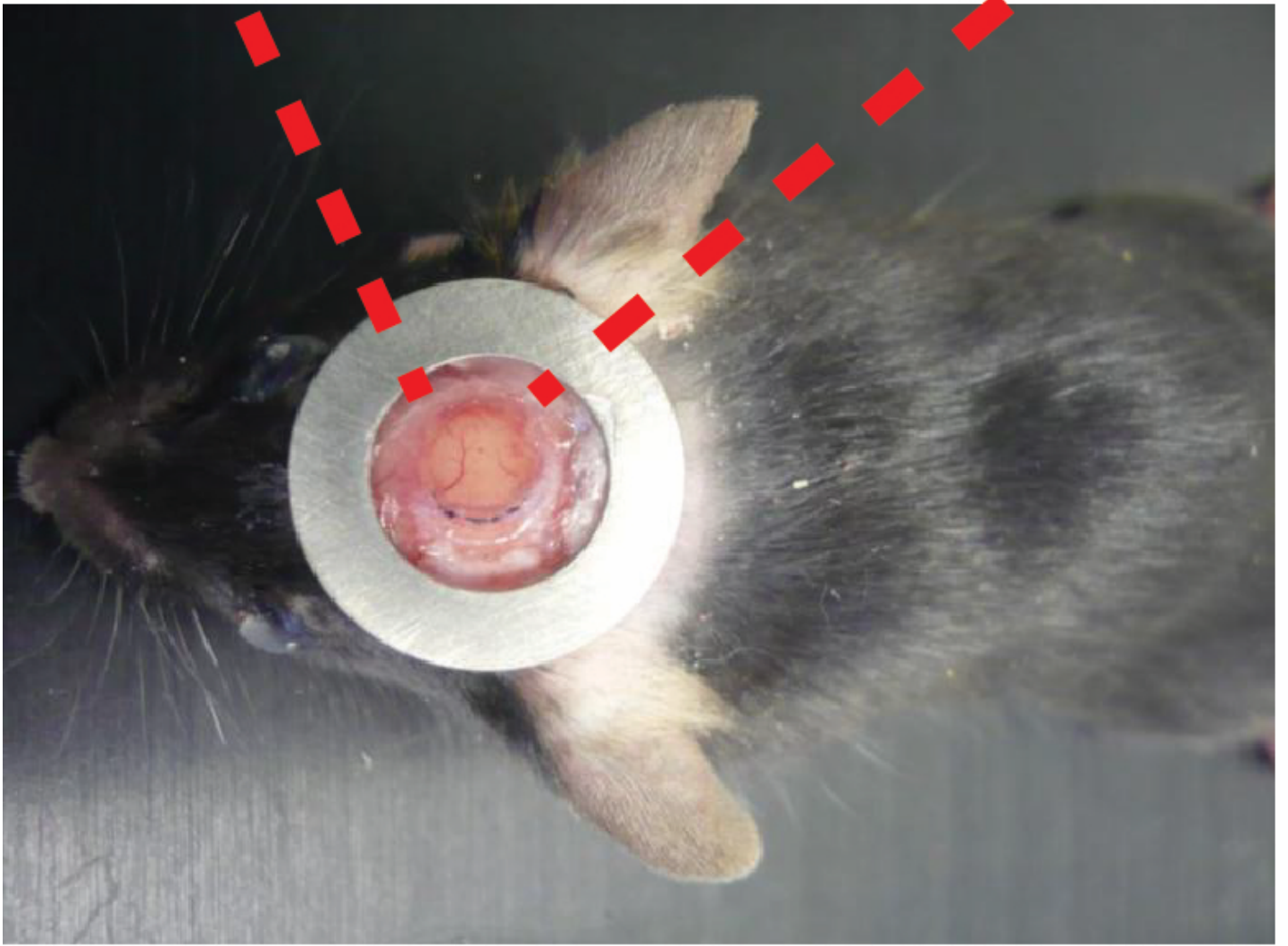
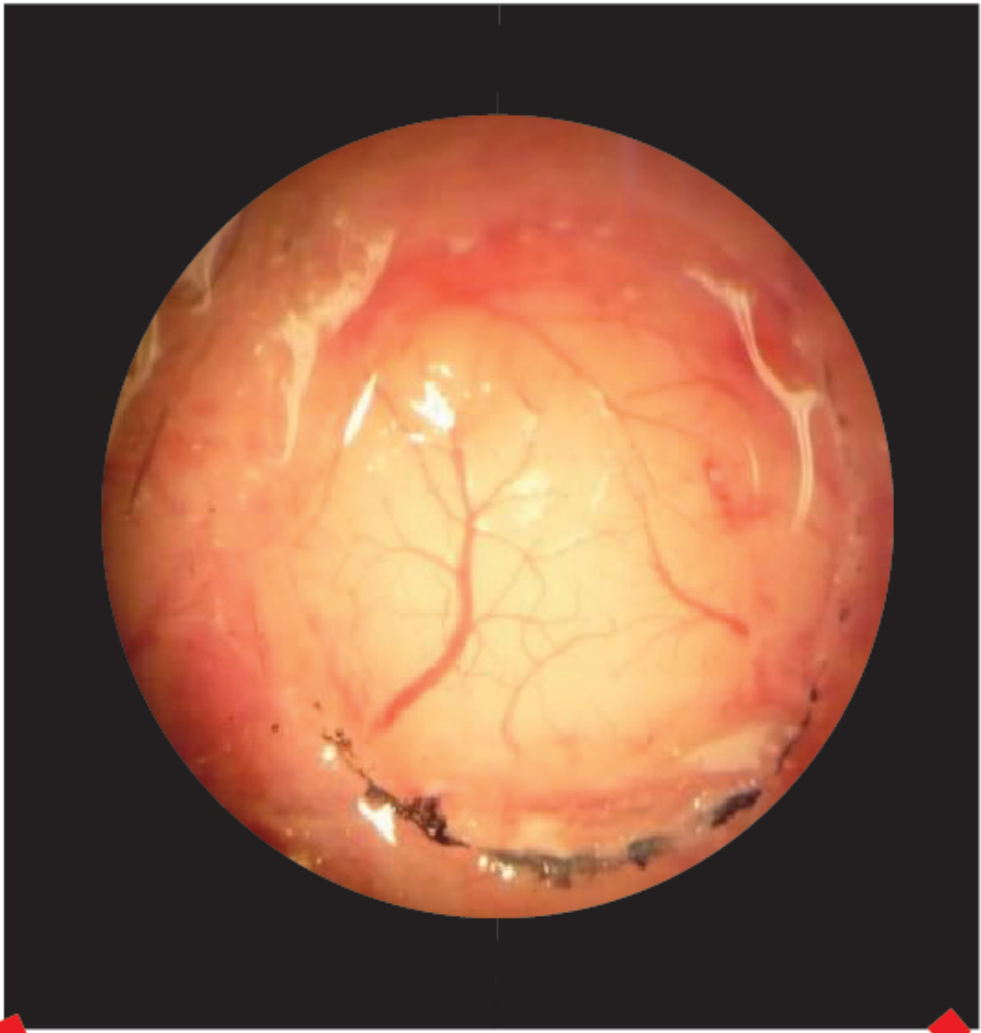
$$\text{COM } y_{\text{diffuse}} = 0.36 + 0.64V_{\text{inv}} - 1.69(V_{\text{inv}} \times F_{\text{inv}})$$

$$R^2 = 0.933$$

$$P < 0.001$$

b





Formula:

COM $y \sim \text{Vinv:T} + \text{Finv:Vinv} + (1 | \text{BTIC})$

Random effects:

Groups	Variance	Std.Dev.
BTIC	0.003019	0.05495
Residual	0.131593	0.36276

Fixed effects:

	Estimate	Std.	Error t value
(Intercept)	0.38738	0.10306	3.759
Vinv:Tborder	0.84942	0.08389	10.126
Vinv:Tinvasive	0.62006	0.06715	9.233
Vinv:Tdiffuse	0.64303	0.05678	11.324
Vinv:Finv	-1.68903	0.07633	-22.128

Parameter	Definition
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COMy	The difference in the centre of mass along the Y axes between its initial value and that at the end of the experiment per unit of time (h) within a position
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Vinv	Absolute velocity perpendicular to the tumor border of invading cells within a position
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Vret	Absolute velocity perpendicular to the tumor border of retreating cells within a position
------	---

Finv	Frequency of invading cells within a position
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T	Type of border configuration
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BTIC	Type of BTIC cell line
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