

# Package: InvasionCorrection (via r-universe)

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**Type** Package

**Title** Invasion Correction

**Version** 0.1

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**Description** The correction is achieved under the assumption that non-migrating cells of the essay approximately form a quadratic flow profile due to frictional effects, compare law of Hagen-Poiseuille for flow in a tube. The script fits a conical plane to give xyz-coordinates of the cells. It outputs the number of migrated cells and the new corrected coordinates.

**License** GPL-3

**Depends** lattice, stats, utils

**LazyData** TRUE

**RoxygenNote** 5.0.1

**NeedsCompilation** no

**Repository** CRAN

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correctByConicalPlane *Correct invasion data by conical plane*

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### Description

Correct z-component of a 3D collagen invasion essay. The correction is achieved under the assumption that non-migrating cells of the essay approximately form a quadratic flow profile due to frictional effects, compare law of Hagen-Poiseuille for flow in a tube.

### Usage

```
correctByConicalPlane(filename, nrfits = 1000, threshold = -30,  
  plot = FALSE, write_csv = TRUE)
```

### Arguments

filename	Name of data file in csv format. It should contain columns "Pos_X", "Pos_Y" and "Pos_Z".
nrfits	Numeric, Number of randomly chosen starting points for the optimization. Choose lower values for speeding up computational time. Choose higher values for more reliable optimization results.
threshold	Numeric, A threshold for counting cells as being invaded or not. When cells move towards negative z-direction, threshold should be negative.
plot	Boole, if TRUE exemplary 3D plots before and after the correction are plotted
write_csv	if TRUE resulting corrected values are saved as a csv file

### Value

Data.frame containing input positions, corrected z-positions as well as number and percentage of invaded cells.

### Author(s)

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