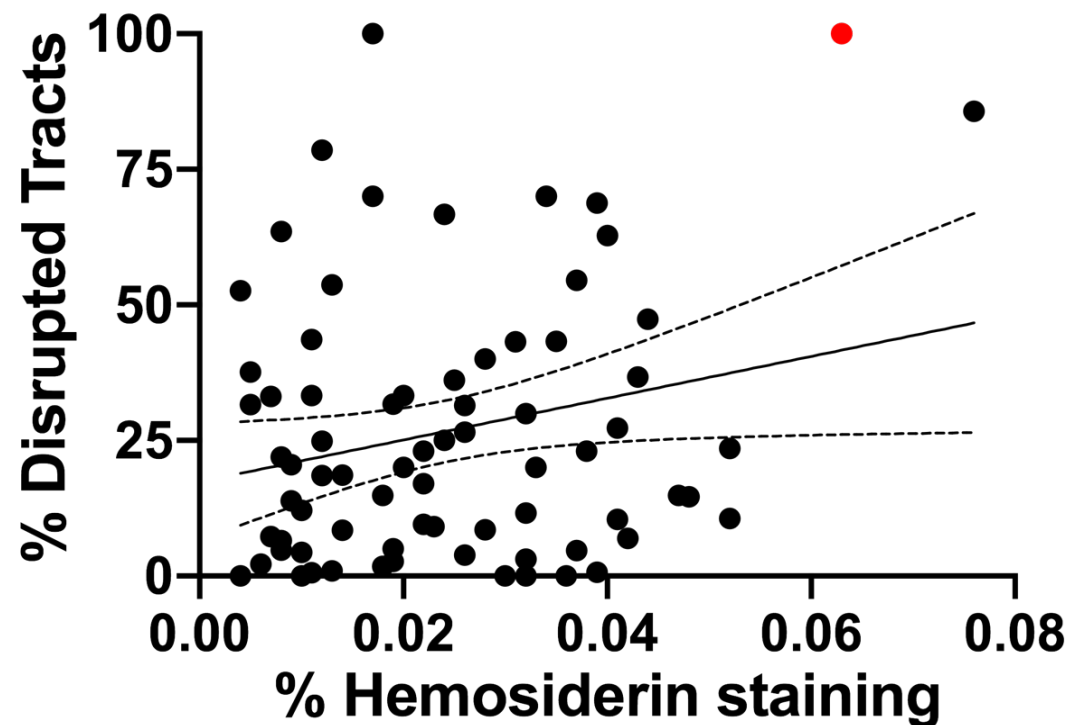


Variables	% Disrupted Tracts
# APP swellings/ μm^2	9.1 [3.7, 14.5]
<i>p-value</i>	0.0012
NF-H % area	-10.4 [-16.1, -4.8]
<i>p-value</i>	0.0005
GFAP % area	-1.6 [-7.0, 3.7]
<i>p-value</i>	0.5435
IBA1 % area	1.4 [-5.6, 8.5]
<i>p-value</i>	0.6851
Heme % area	4.2 [-1.0, 9.5]
<i>p-value</i>	0.1031
LFB intensity	-3.3 [-9.9, 3.4]
<i>p-value</i>	0.3282



Supplemental Table 4. Multiple Linear Regression Analysis without one of two ROIs driving Heme correlation indicated in red. Degrees of freedom = 66. Coefficient [Confidence Interval]; APP = amyloid precursor protein, NF-H = neurofilament, GFAP = glial fibrillary acidic protein, IBA1 = ionized calcium binding adaptor molecule 1, Heme = hemosiderin, LFB =Luxol fast blue.