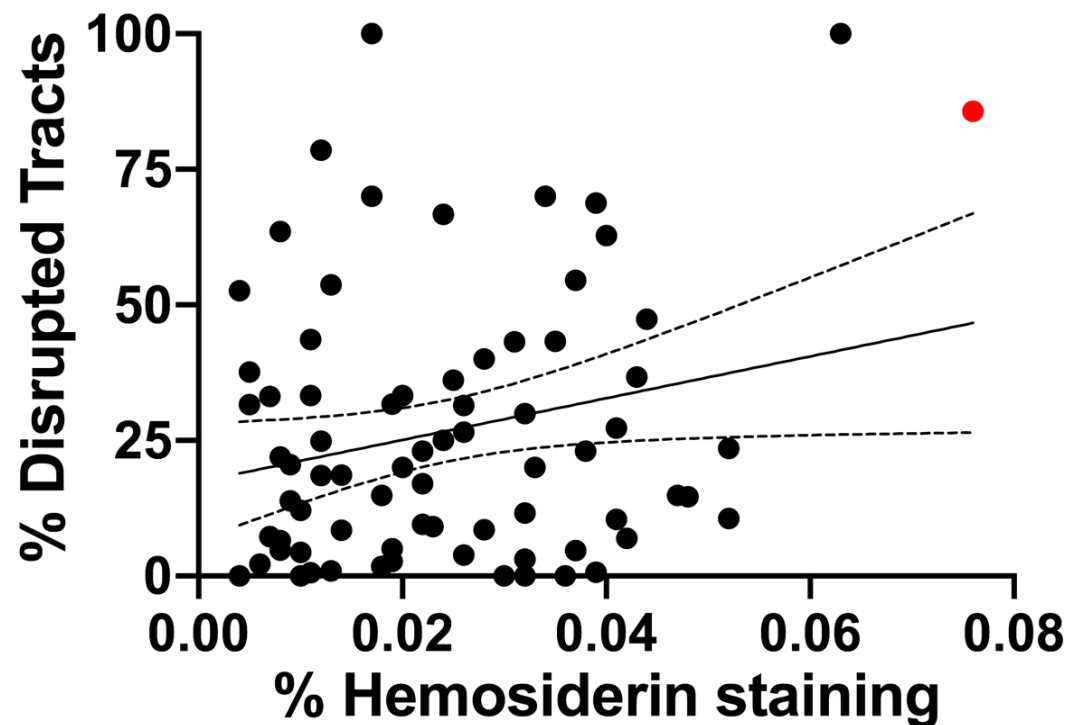


Variables	% Disrupted Tracts
# APP swellings/ μm^2	9.5 [4.0, 14.9]
<i>p-value</i>	0.0009
NF-H % area	-11.4 [-17.0, -5.7]
<i>p-value</i>	0.0002
GFAP % area	-1.9 [-7.4, 3.6]
<i>p-value</i>	0.4899
IBA1 % area	0.4 [-6.4, 7.2]
<i>p-value</i>	0.9016
Heme % area	4.8 [-0.9, 10.5]
<i>p-value</i>	0.0999
LFB intensity	-2.9 [-9.5, 3.8]
<i>p-value</i>	0.3976



Supplemental Table 3. 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.