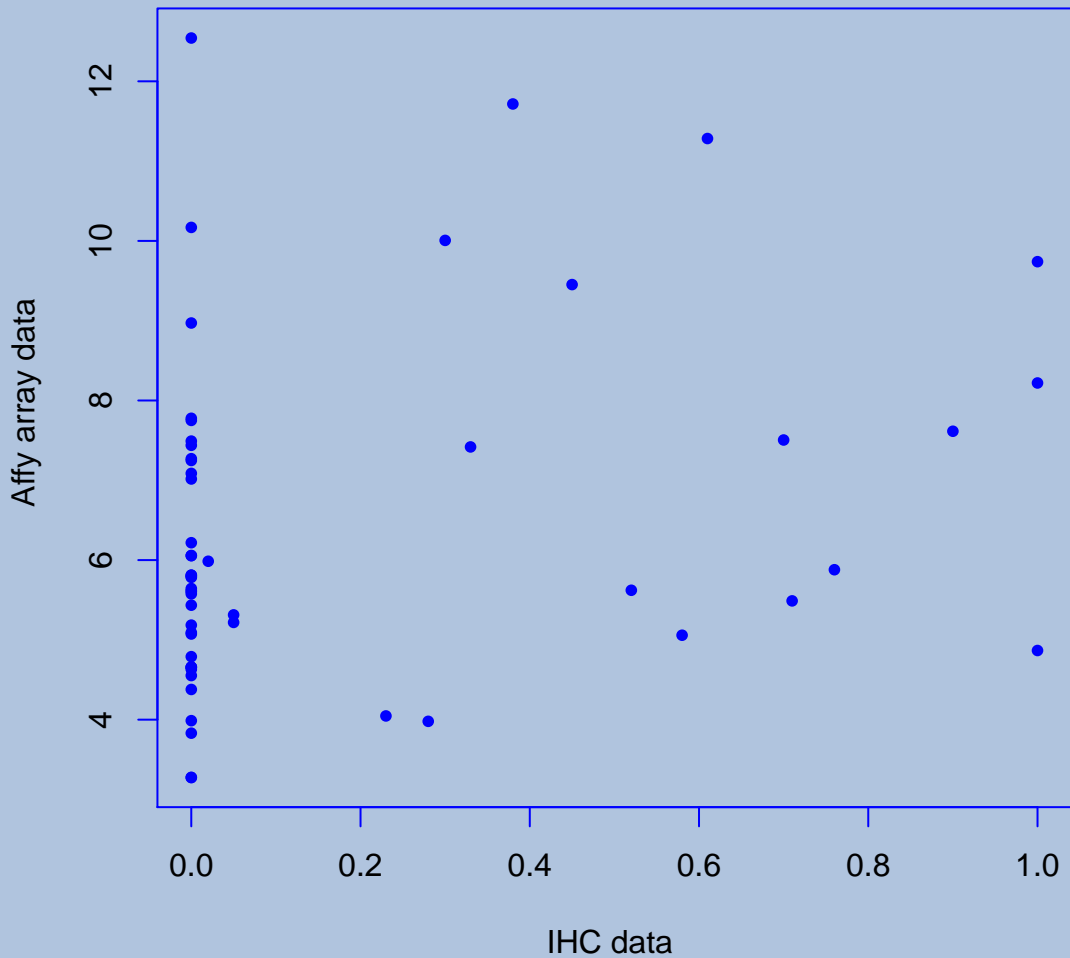
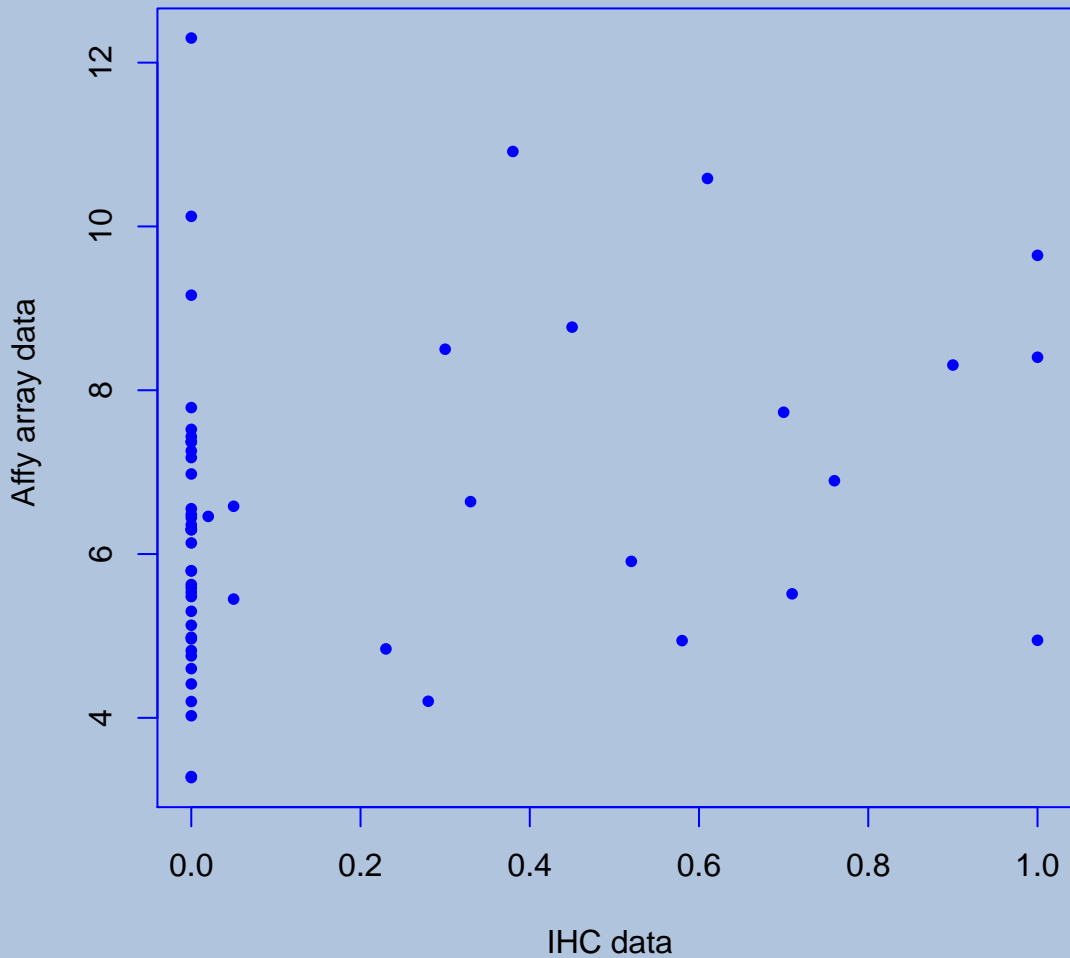


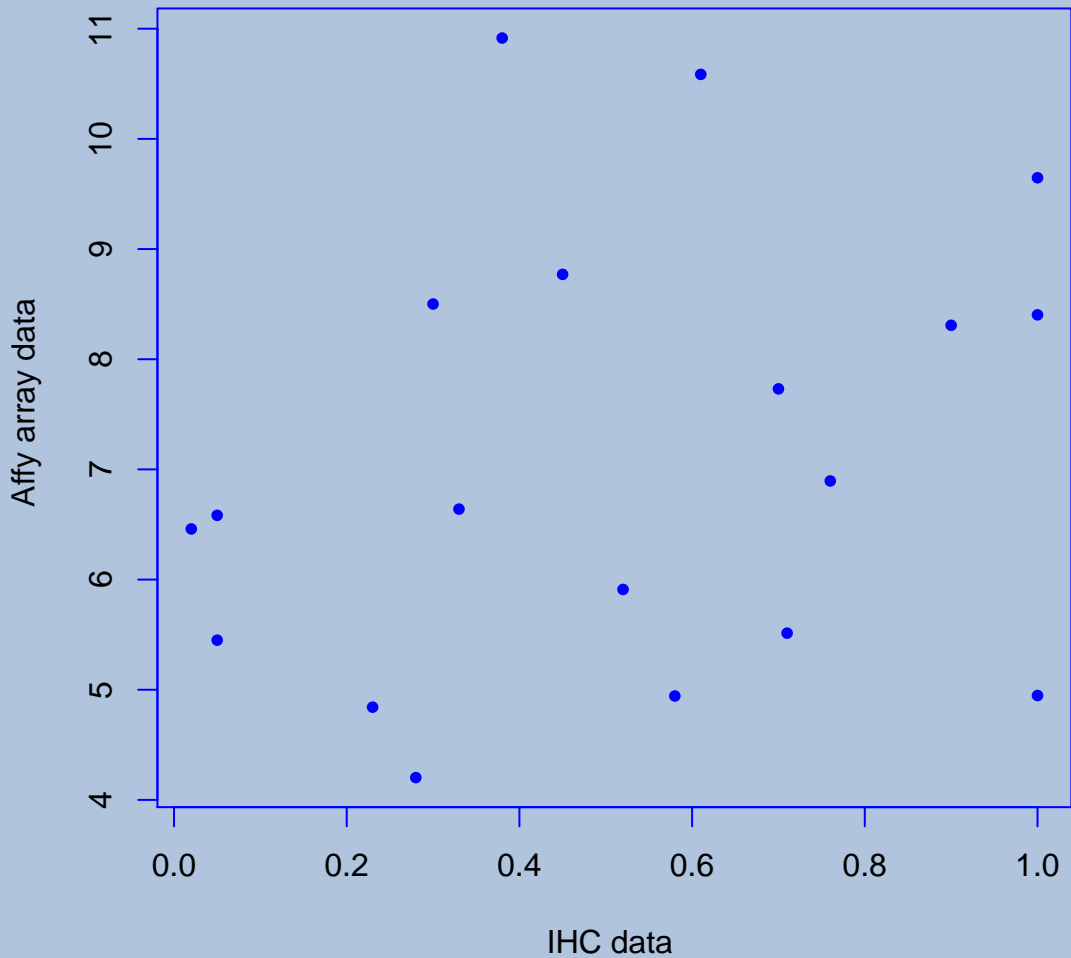
Endothelial_CD31 , median
spearman = 0.25 , pearson = 0.28



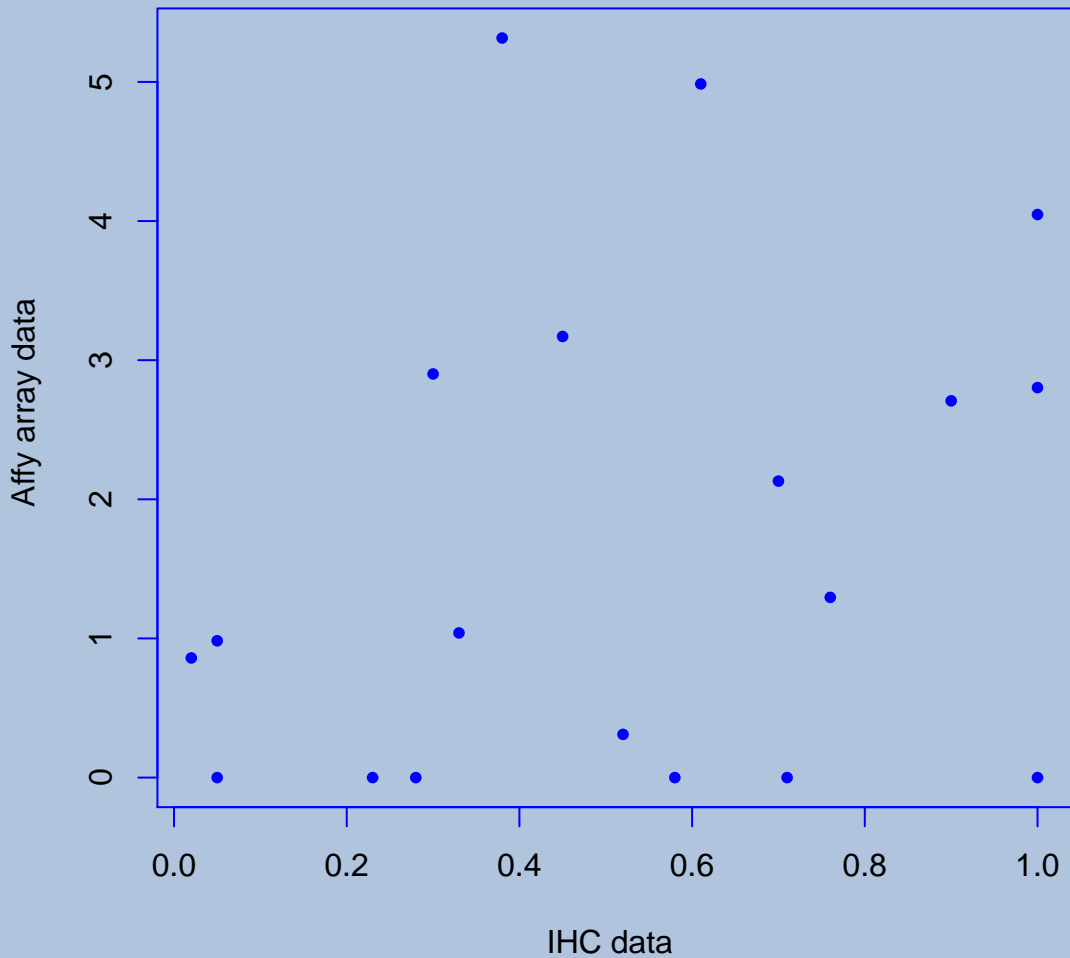
Endothelial_CD31 , mean
spearman = 0.26 , pearson = 0.28



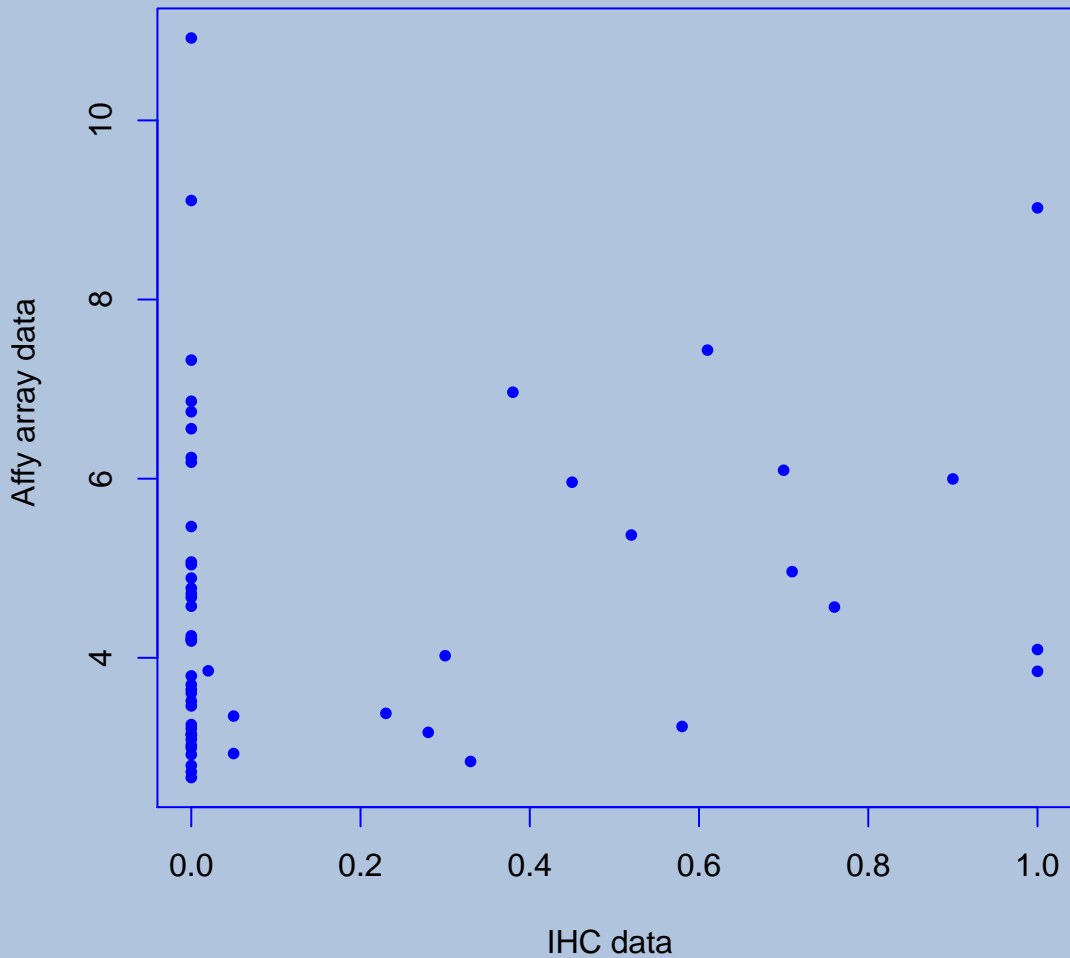
Endothelial_CD31 , no_zeros
spearman = 0.31 , pearson = 0.26



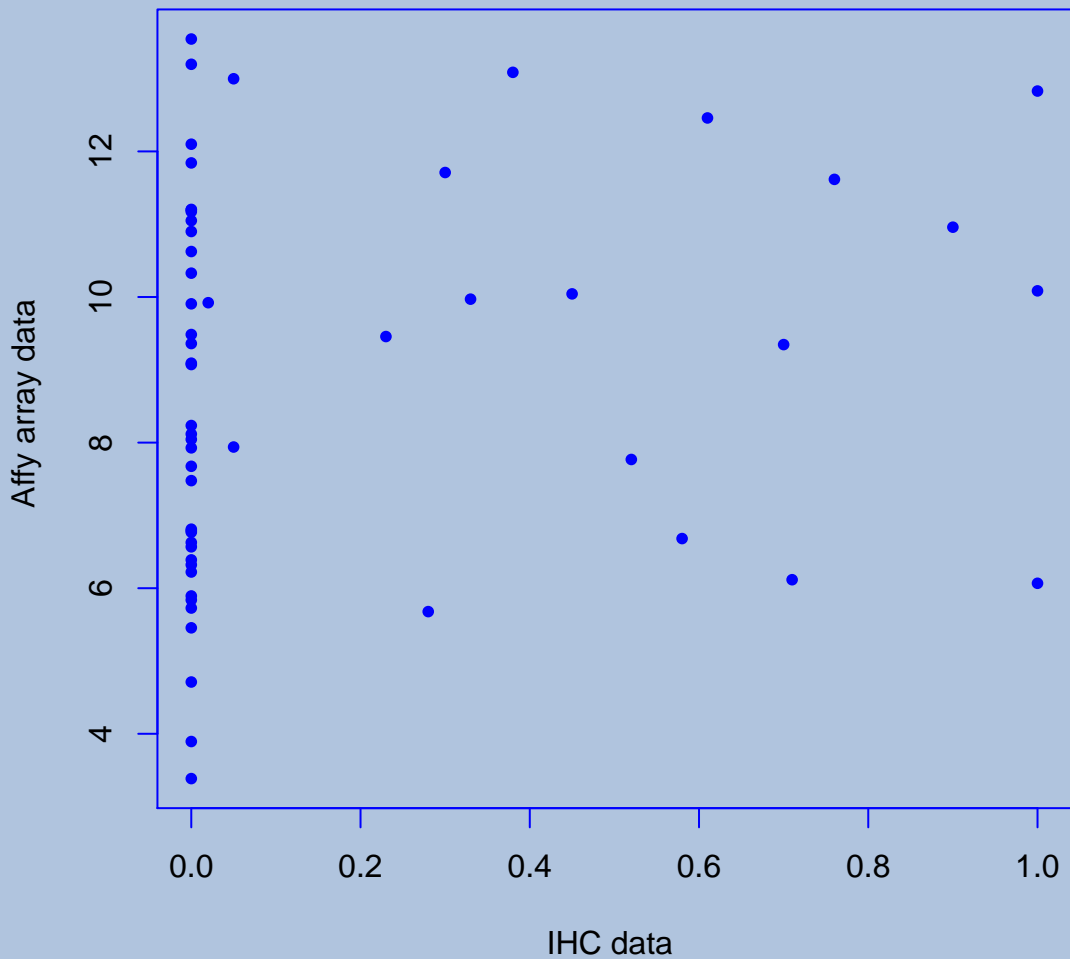
Endothelial_CD31 , xform
spearman = 0.27 , pearson = 0.27



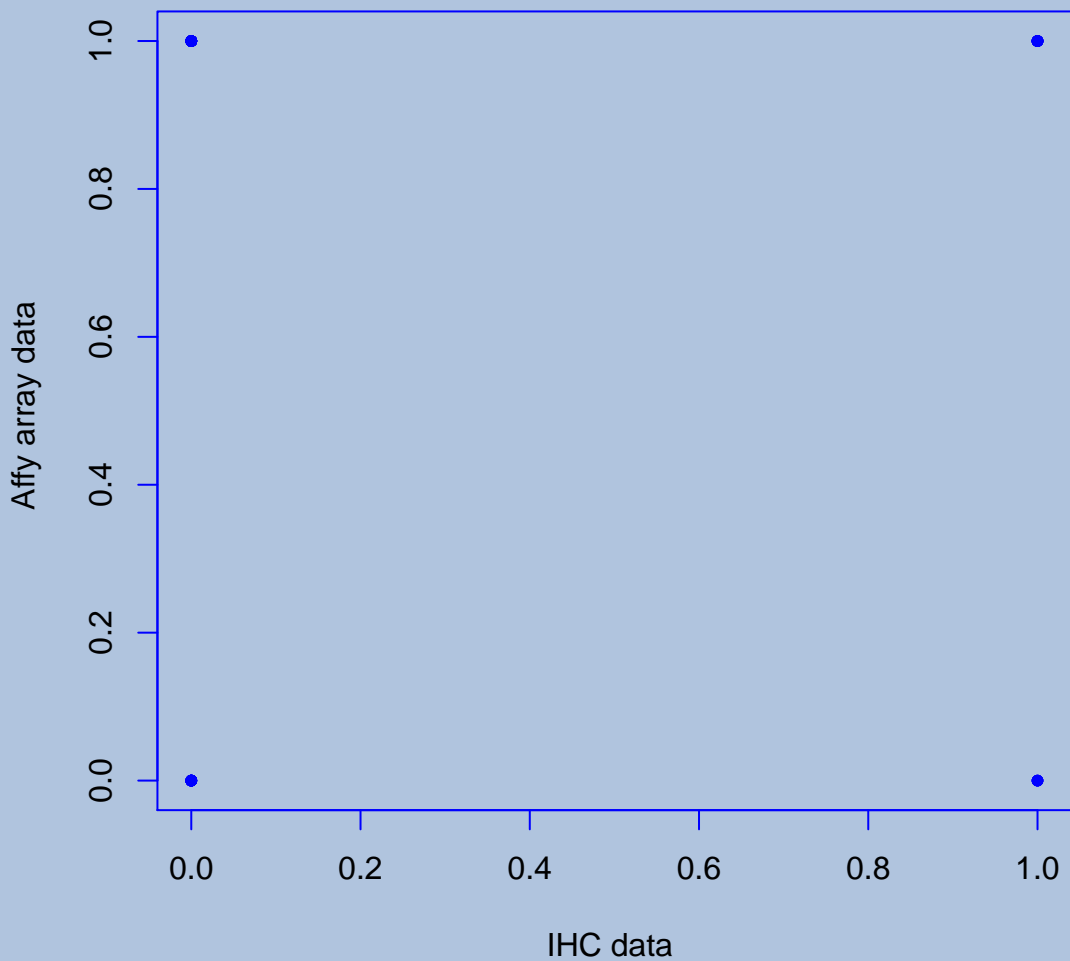
Endothelial_CD31 , min
spearman = 0.14 , pearson = 0.20



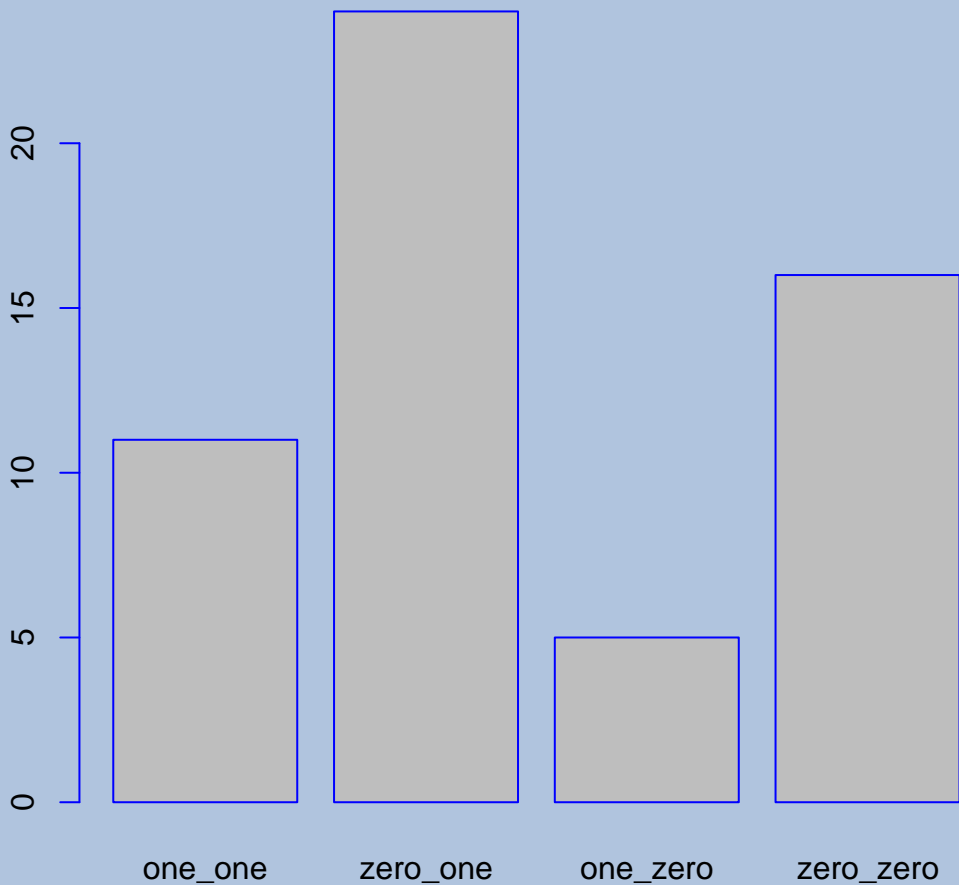
Endothelial_CD31 , max
spearman = 0.22 , pearson = 0.19



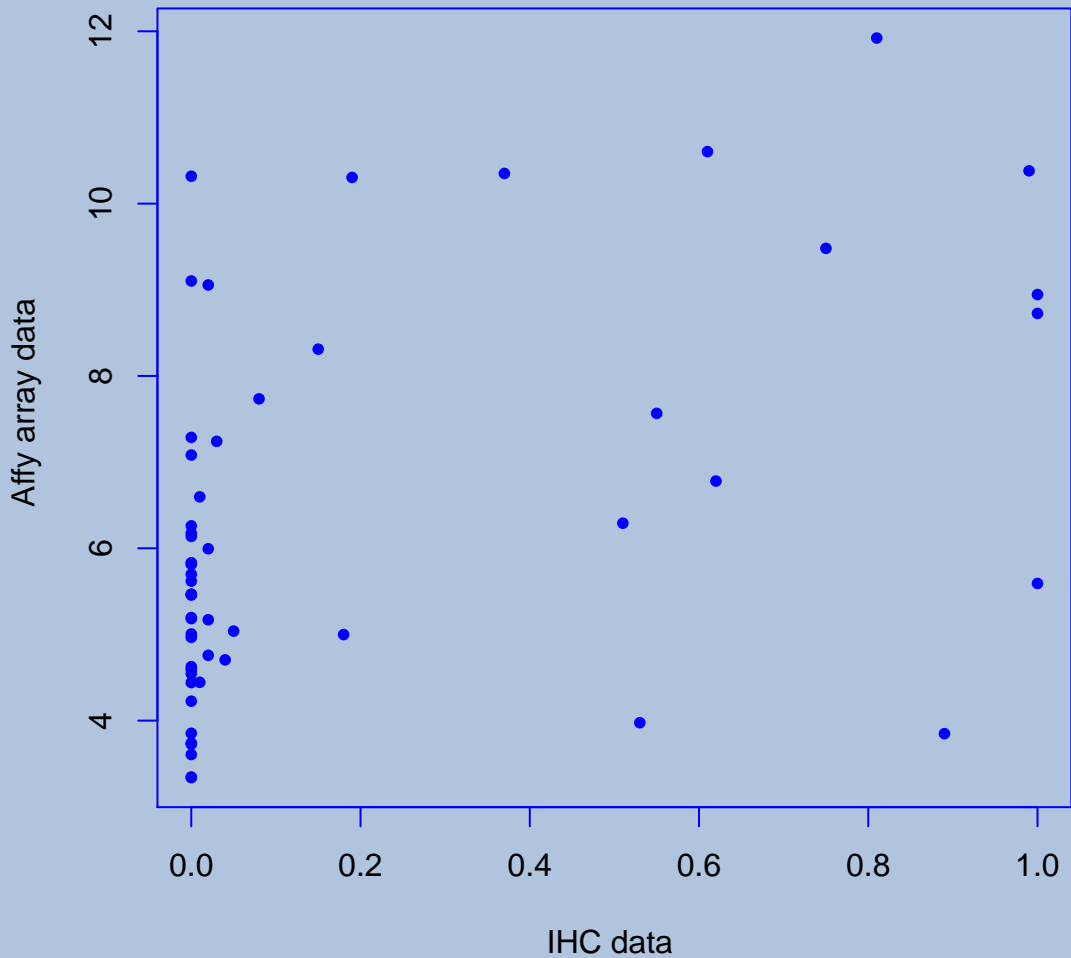
Endothelial_CD31 , digitized
spearman = 0.08 , pearson = 0.08



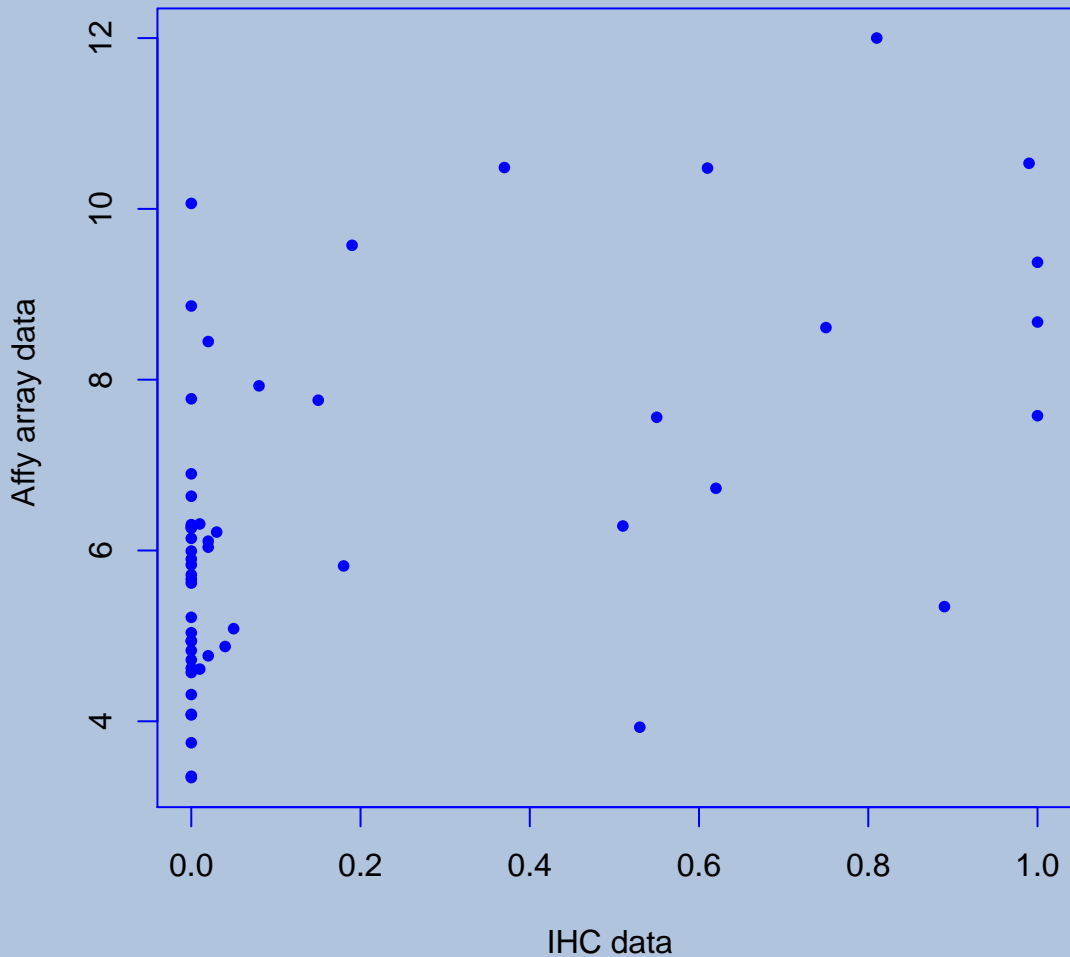
Endothelial_CD31



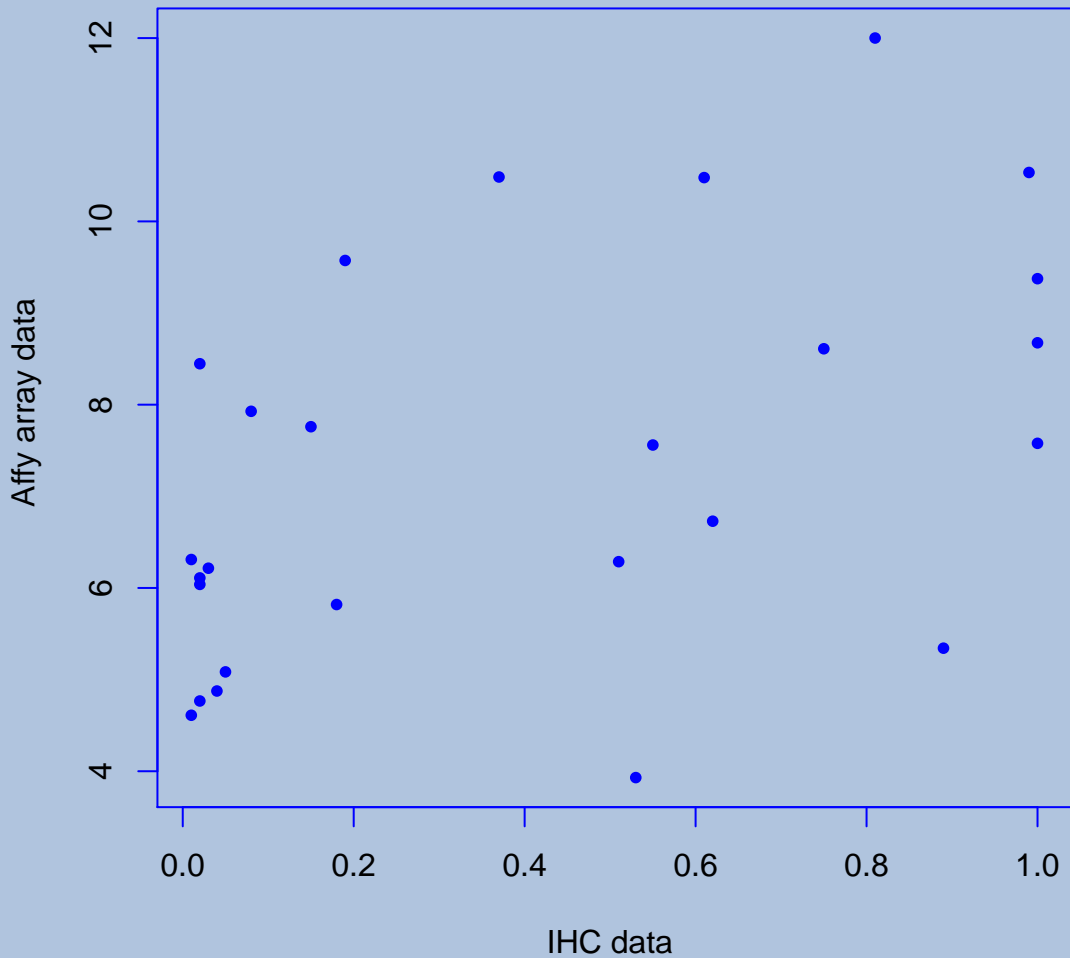
Luminal_CD26 , median
spearman = 0.46 , pearson = 0.47



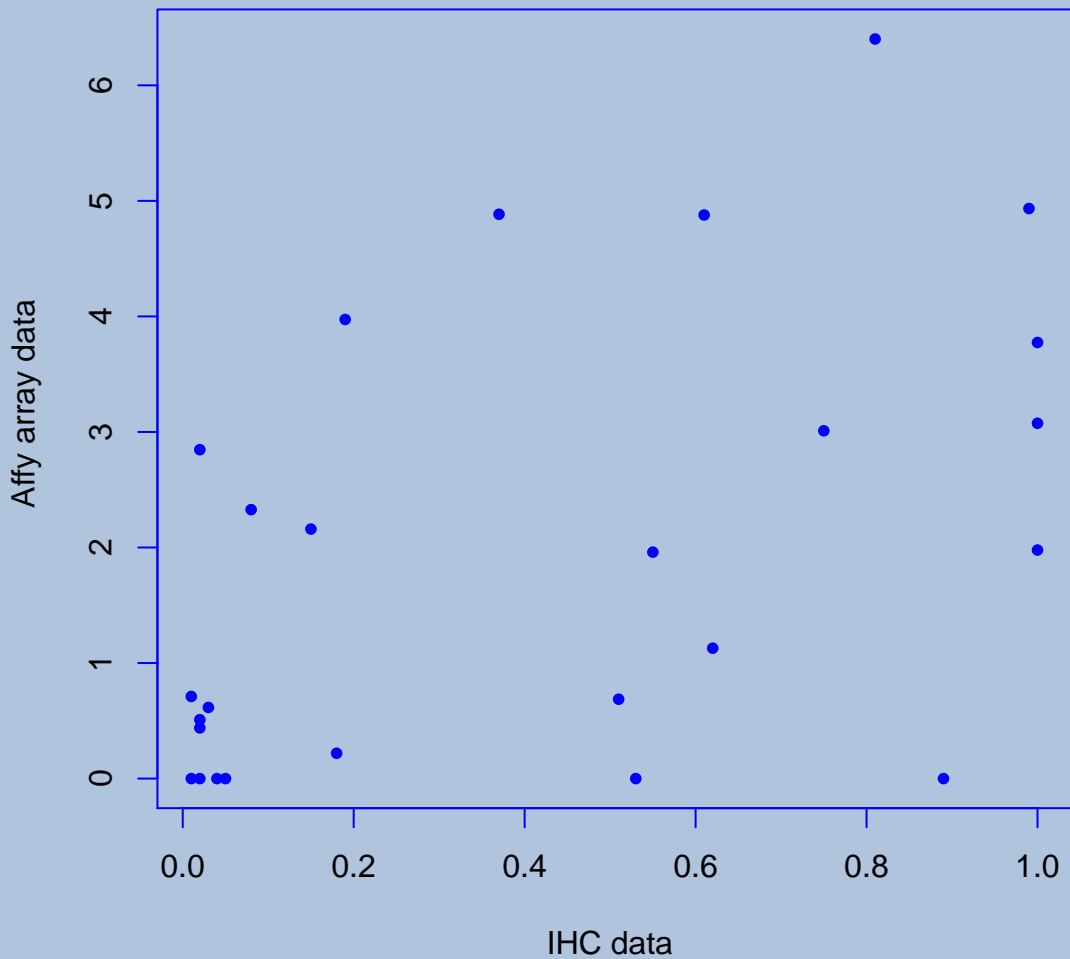
Luminal_CD26 , mean
spearman = 0.51 , pearson = 0.55



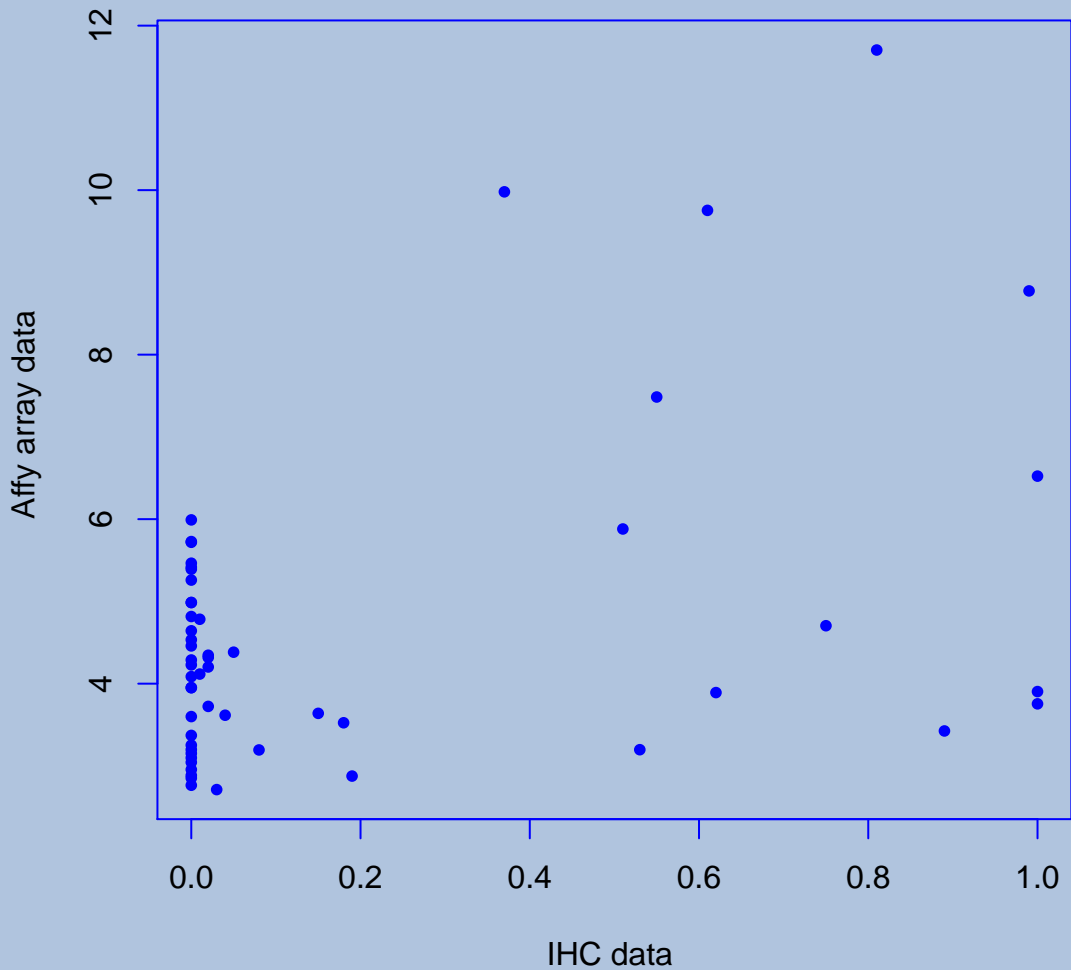
Luminal_CD26 , no_zeros
spearman = 0.52 , pearson = 0.49



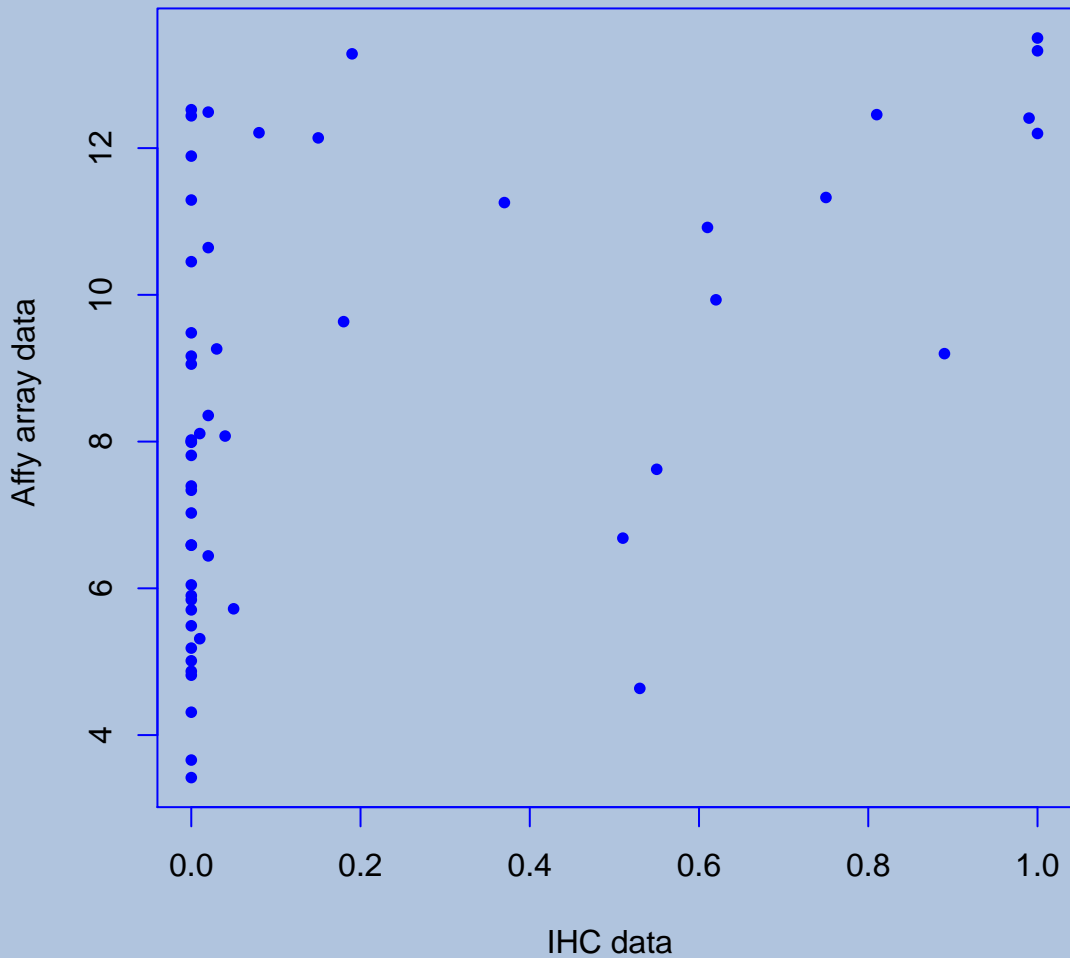
Luminal_CD26 , xform
spearman = 0.51 , pearson = 0.50



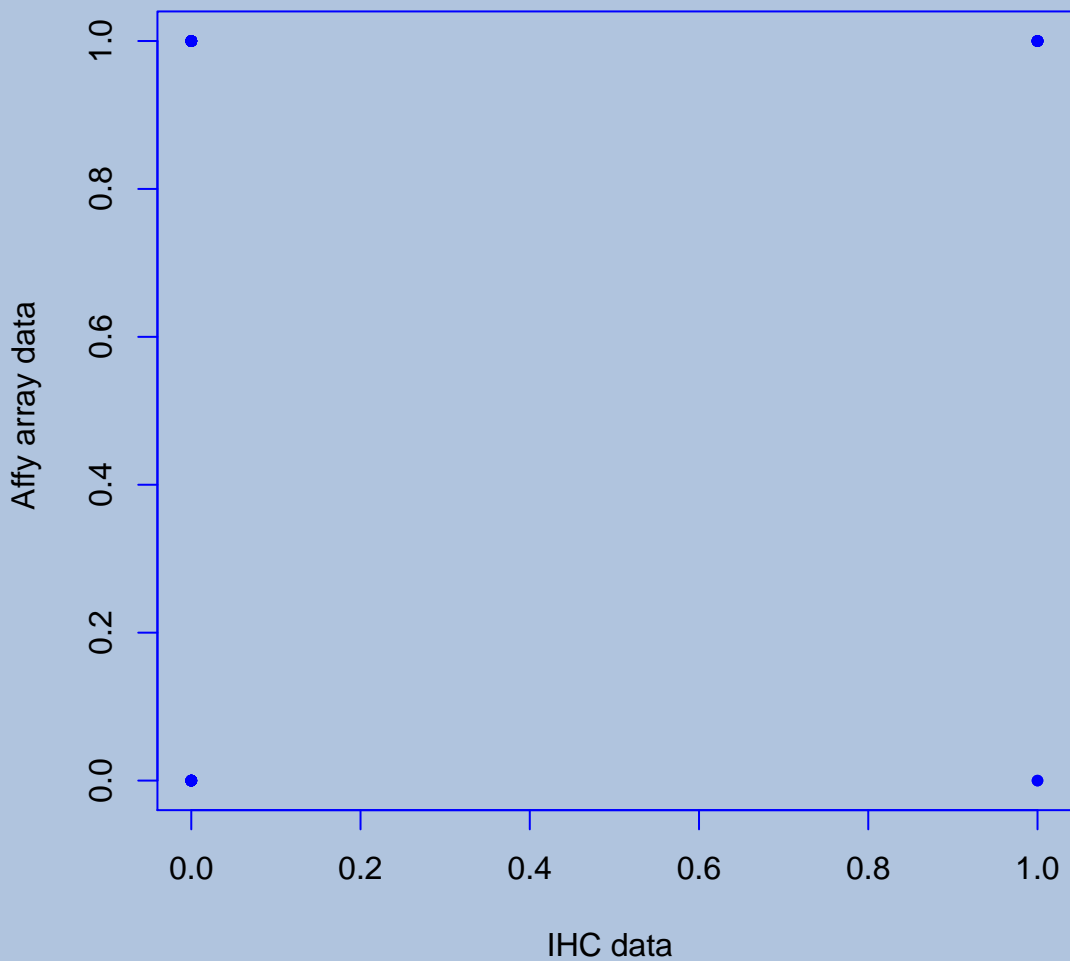
Luminal_CD26 , min
spearman = 0.16 , pearson = 0.43



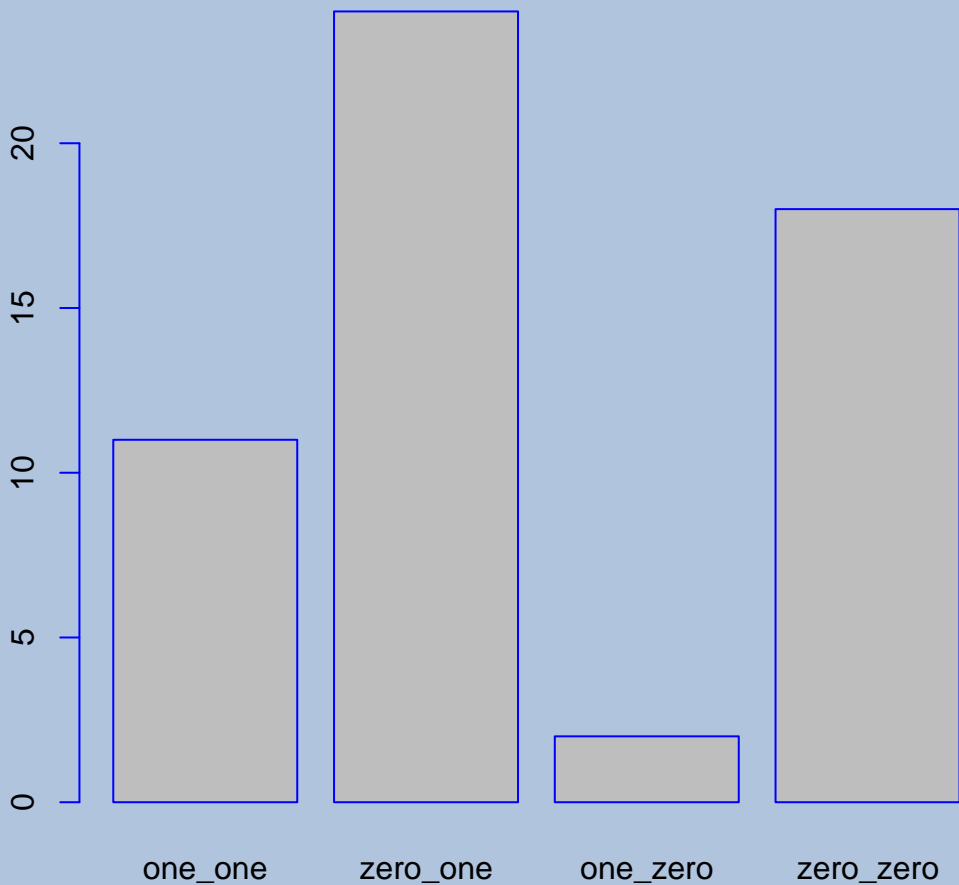
Luminal_CD26 , max
spearman = 0.51 , pearson = 0.48



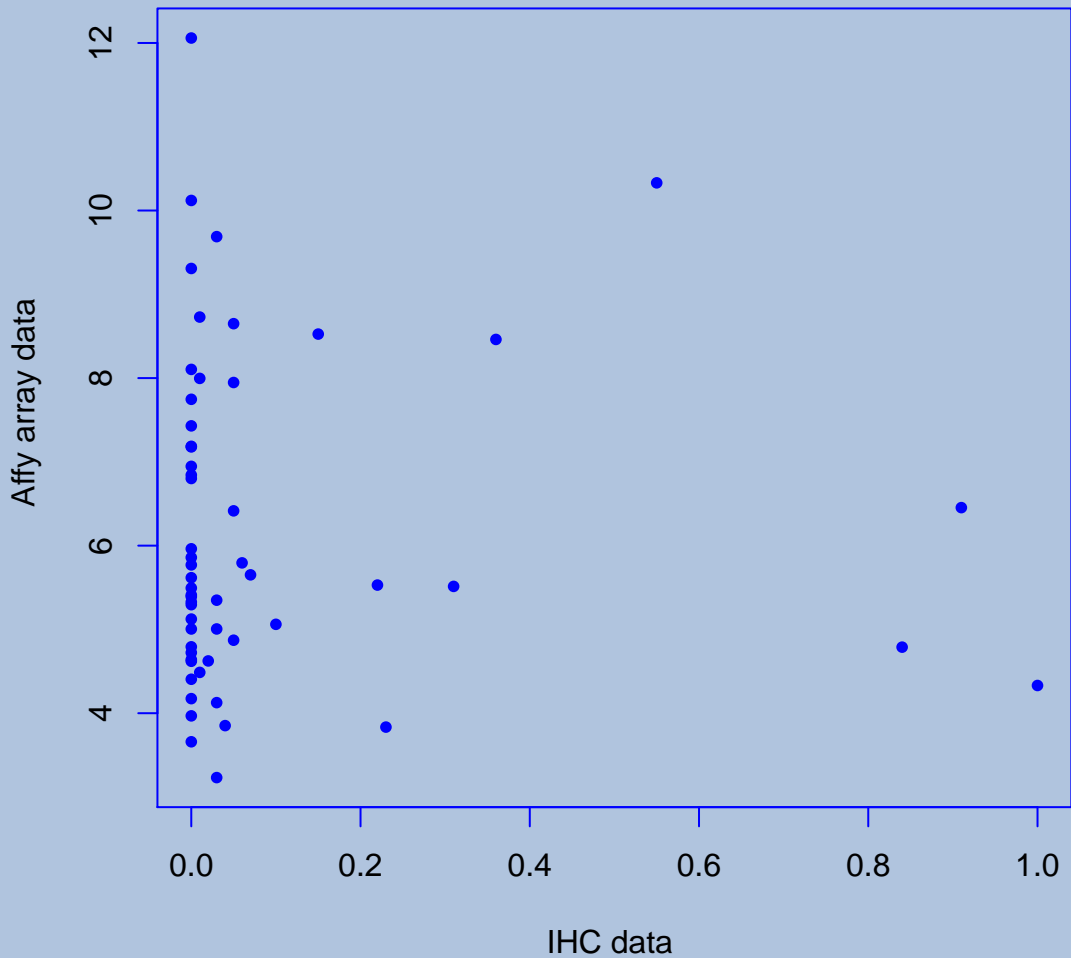
Luminal_CD26 , digitized
spearman = 0.24 , pearson = 0.24



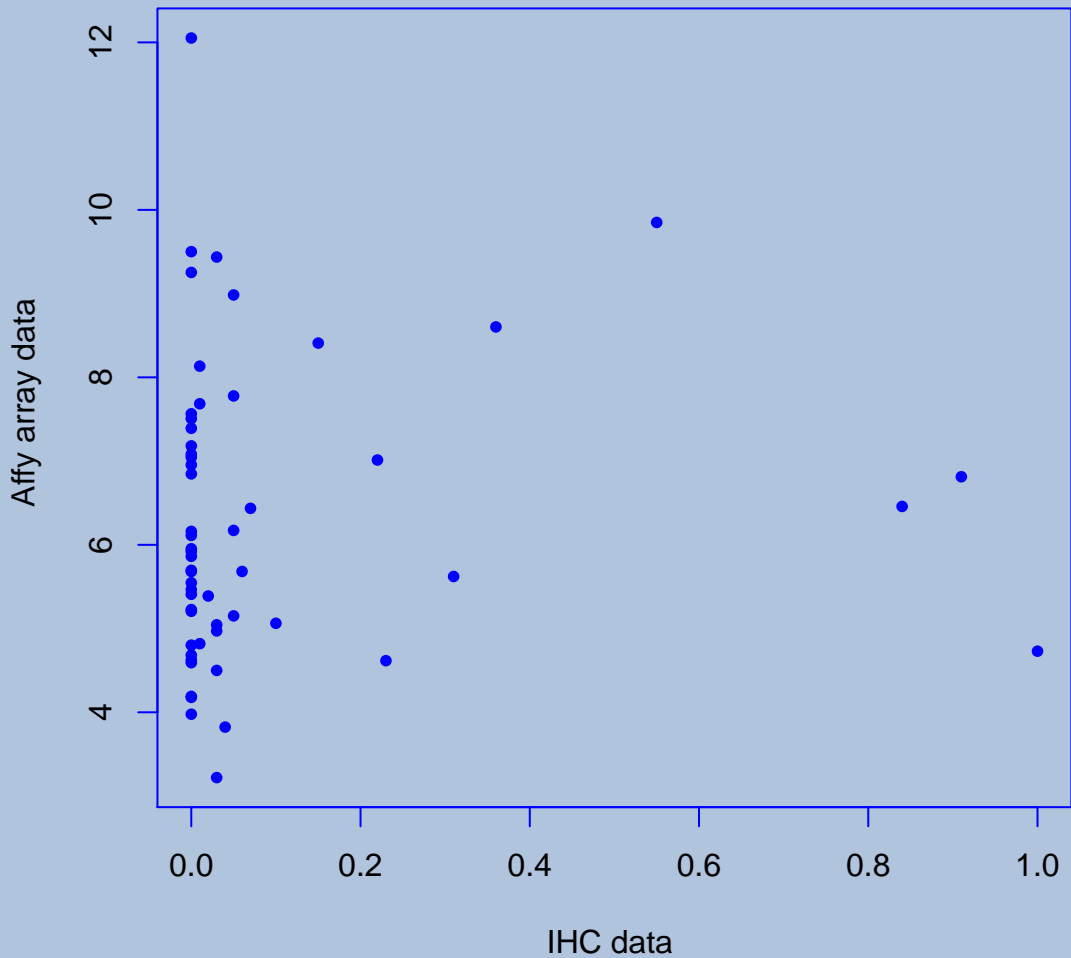
Luminal_CD26



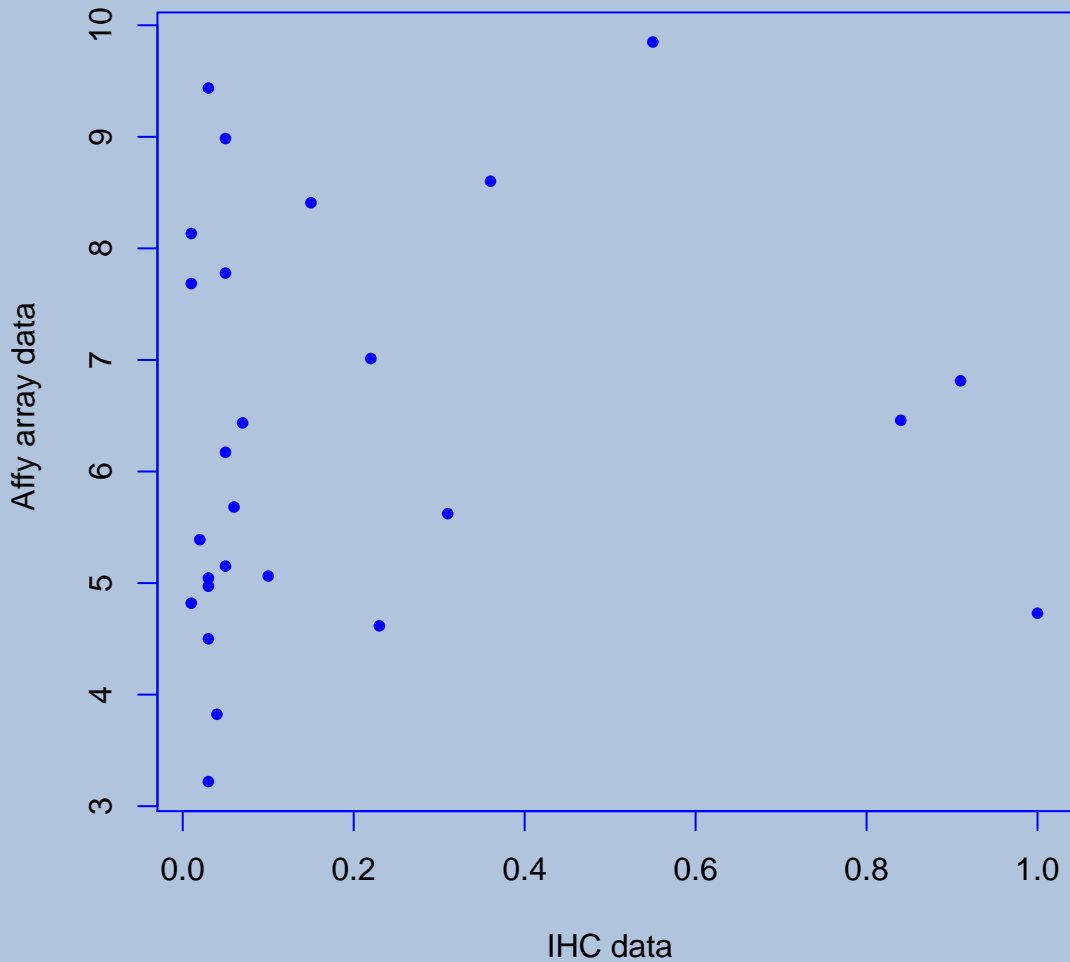
Stromal_CD49a , median
spearman = -0.00 , pearson = -0.01



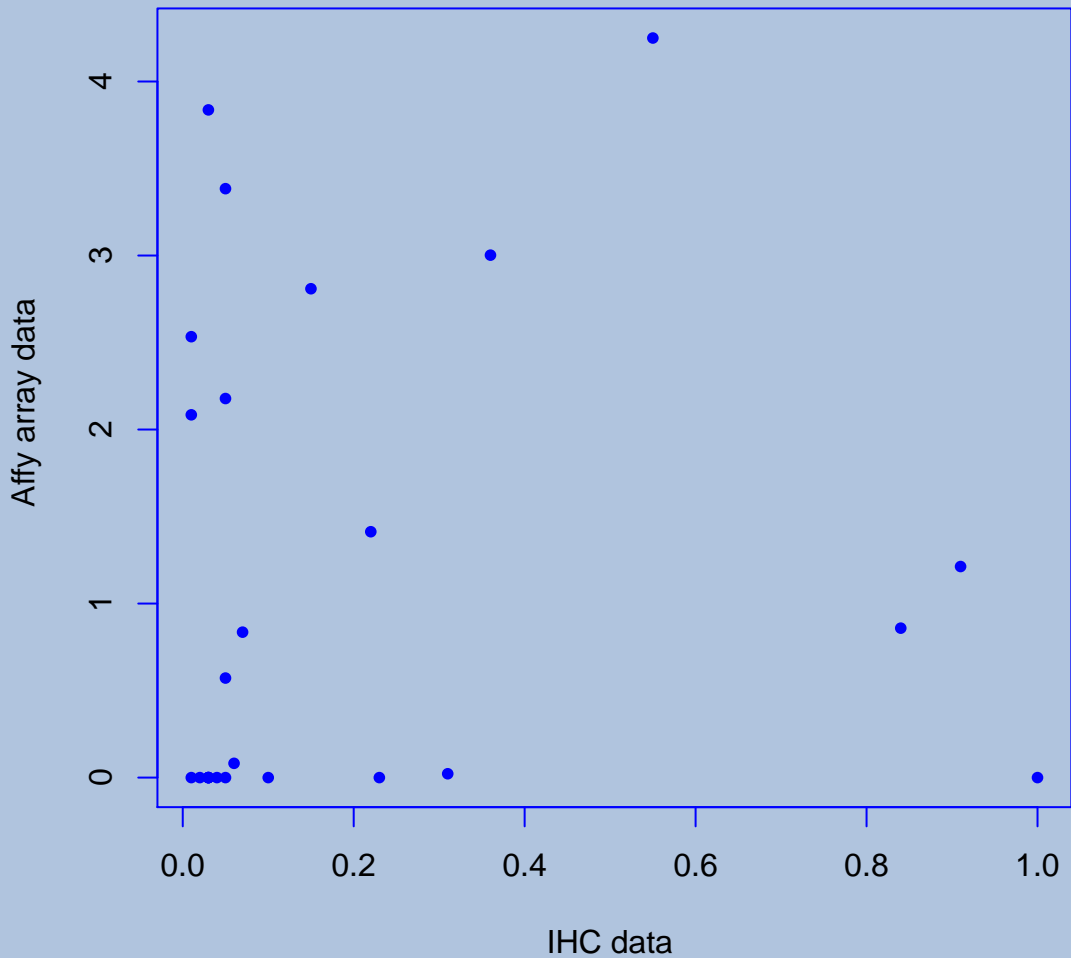
Stromal_CD49a , mean
spearman = 0.06 , pearson = 0.07



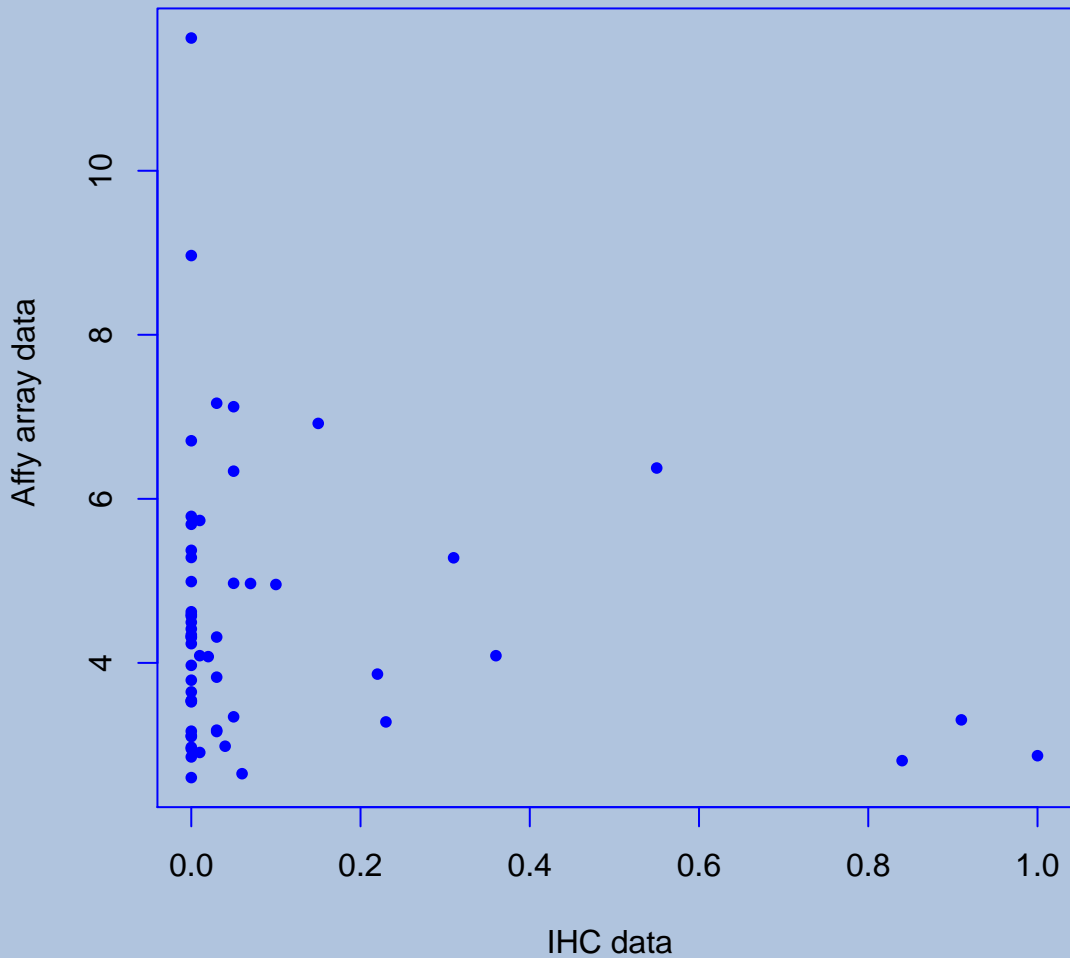
Stromal_CD49a , no_zeros
spearman = 0.19 , pearson = 0.10



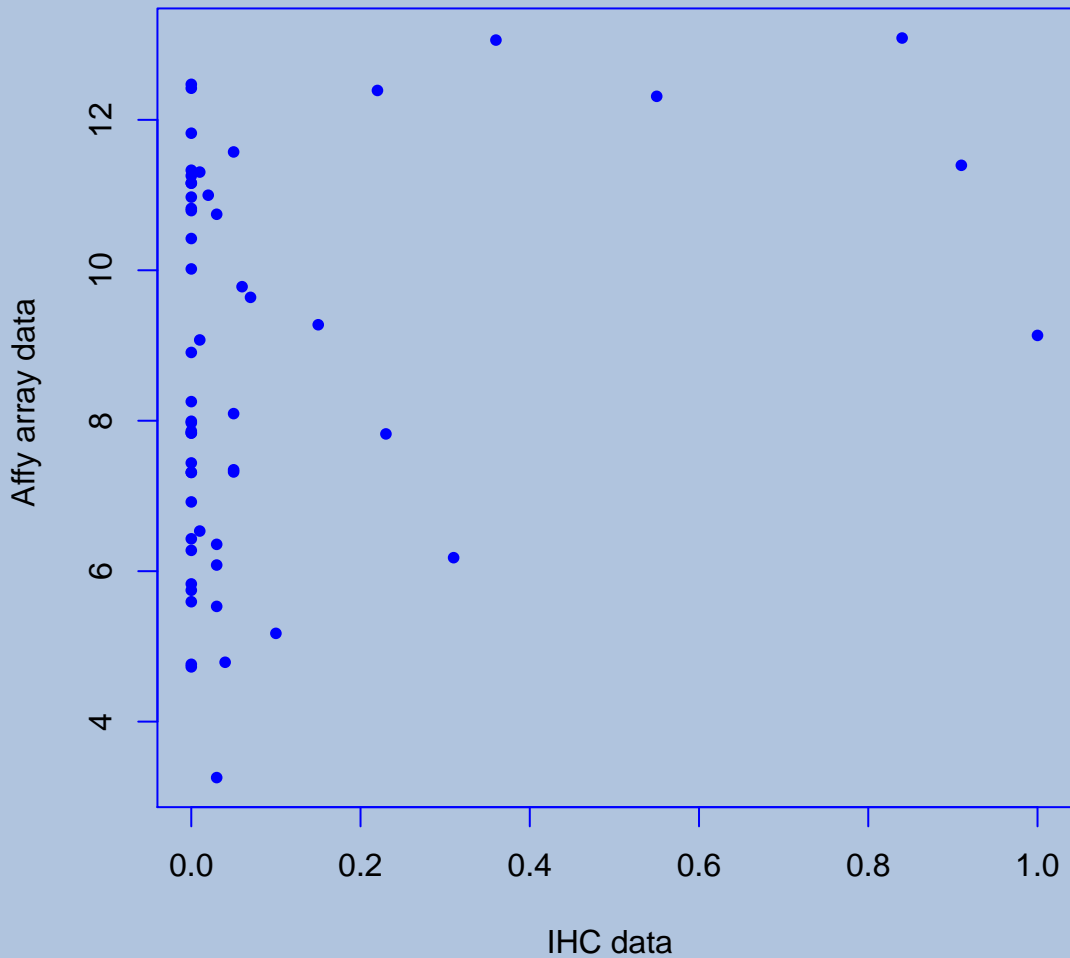
Stromal_CD49a , xform
spearman = 0.21 , pearson = 0.06



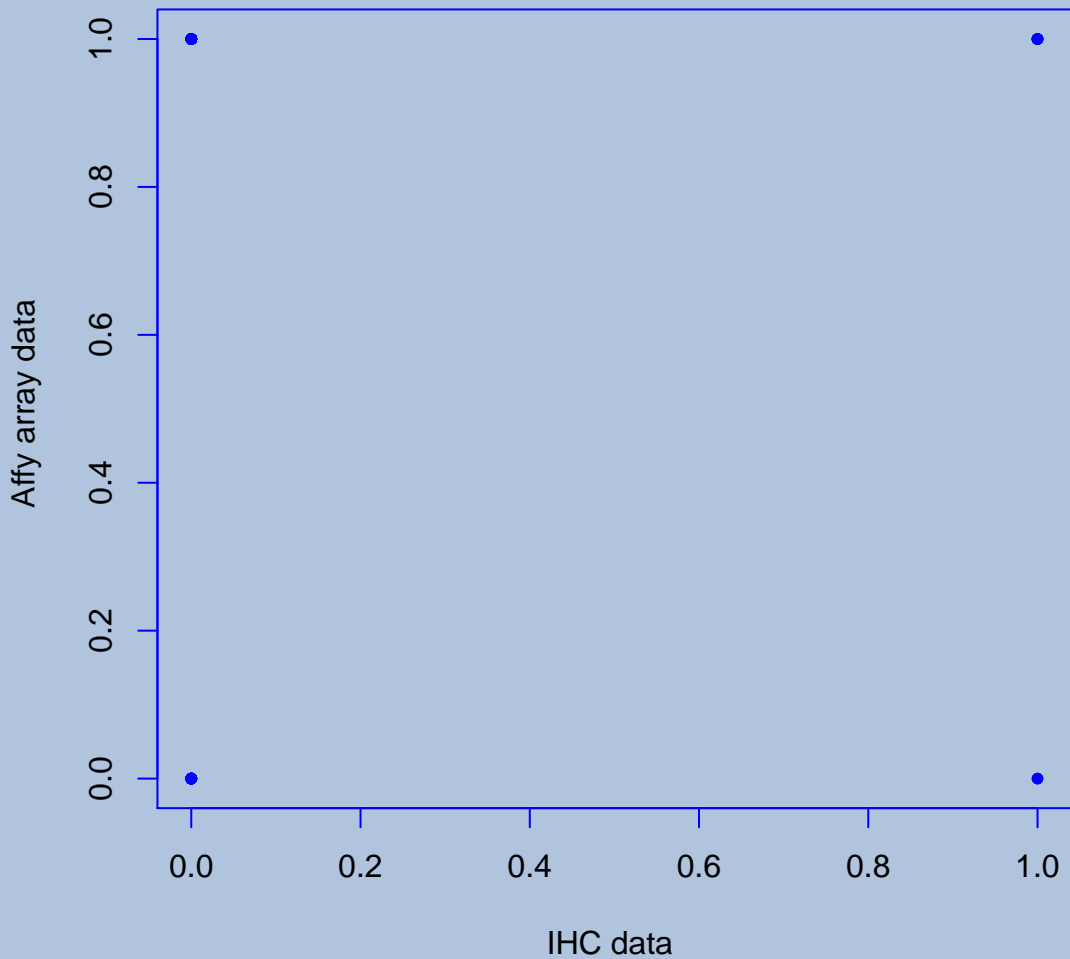
Stromal_CD49a , min
spearman = -0.05 , pearson = -0.15



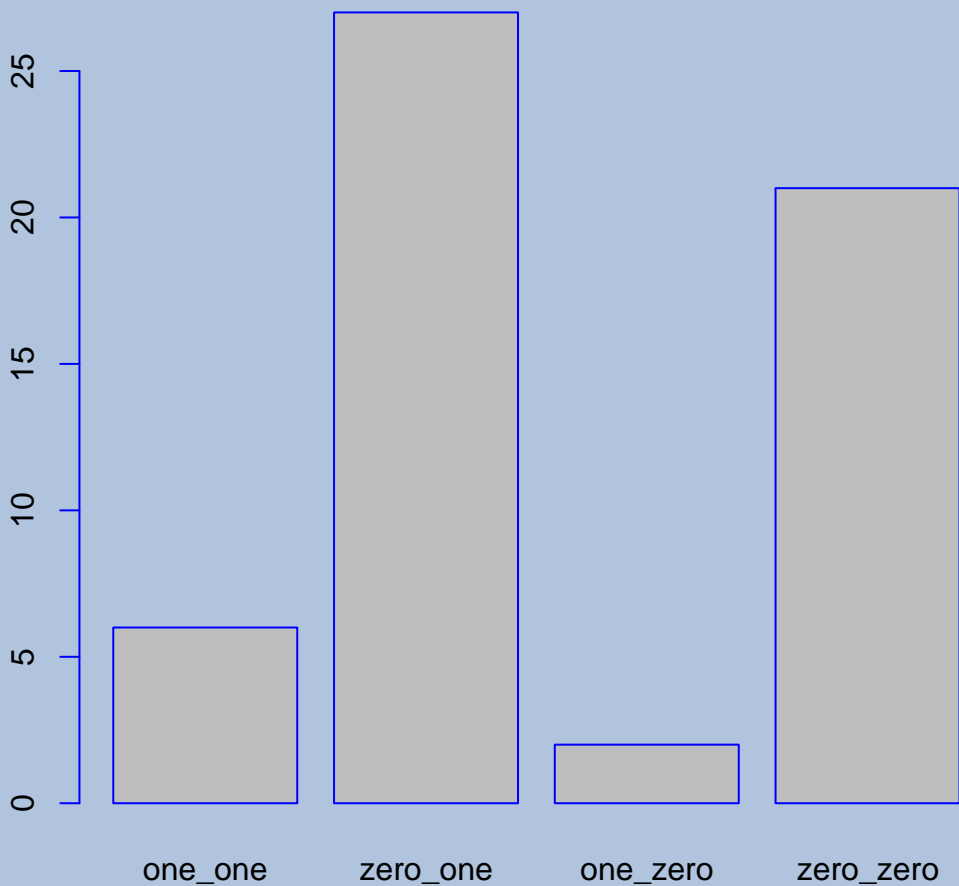
Stromal_CD49a , max
spearman = 0.13 , pearson = 0.30



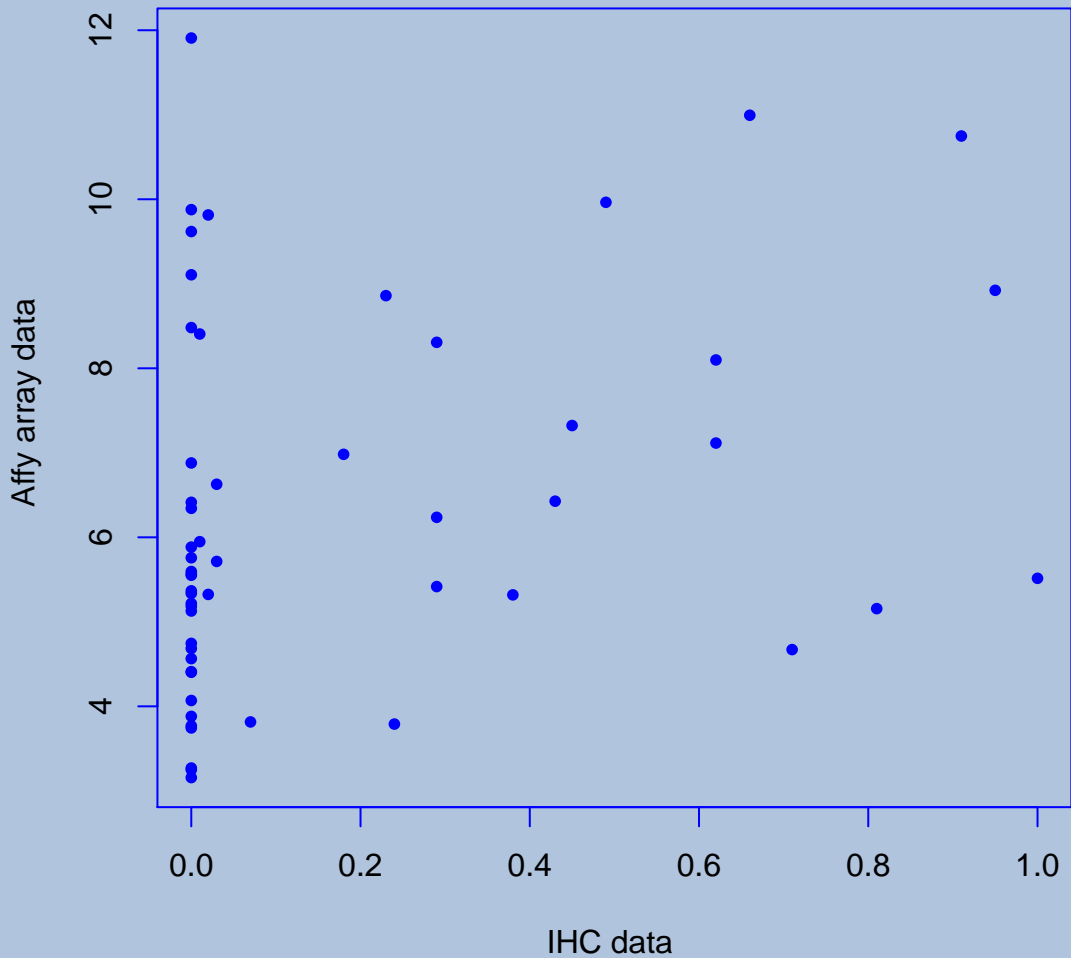
Stromal_CD49a , digitized
spearman = 0.13 , pearson = 0.13



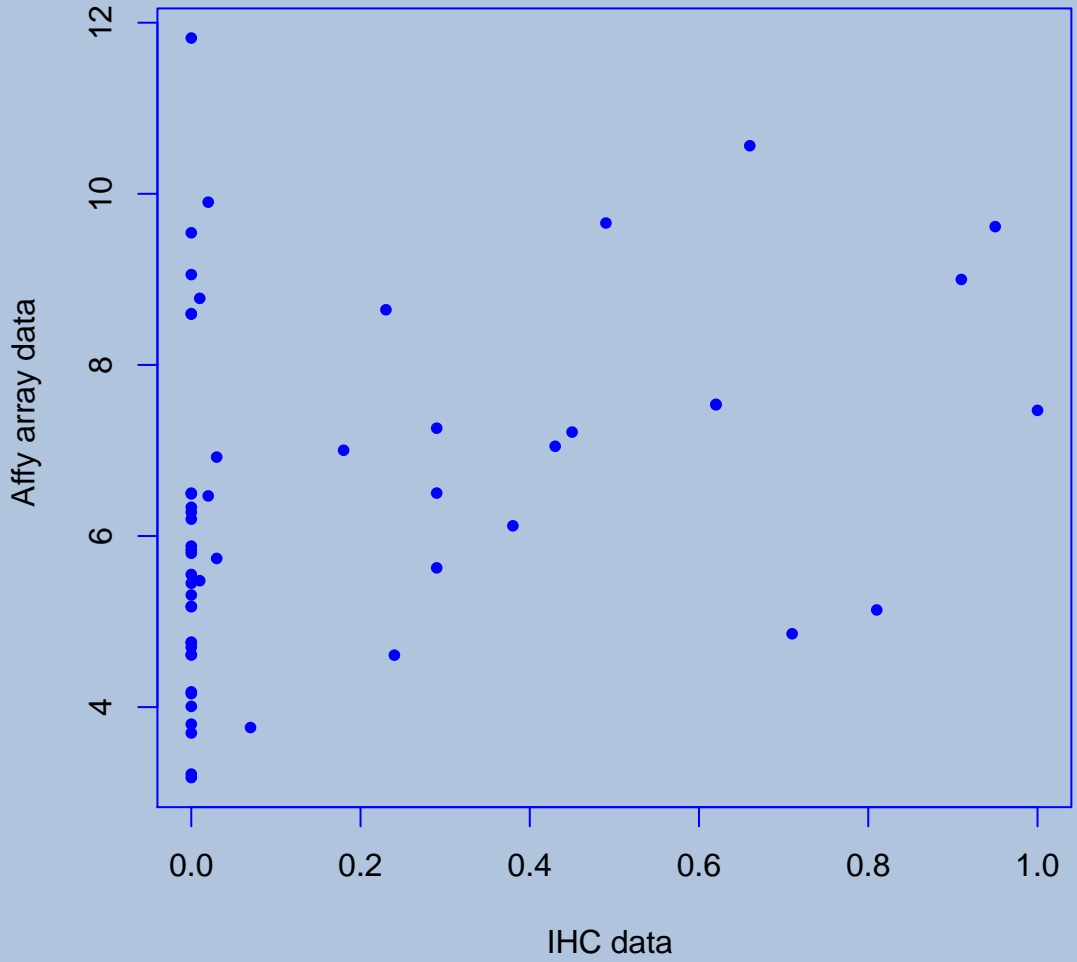
Stromal_CD49a



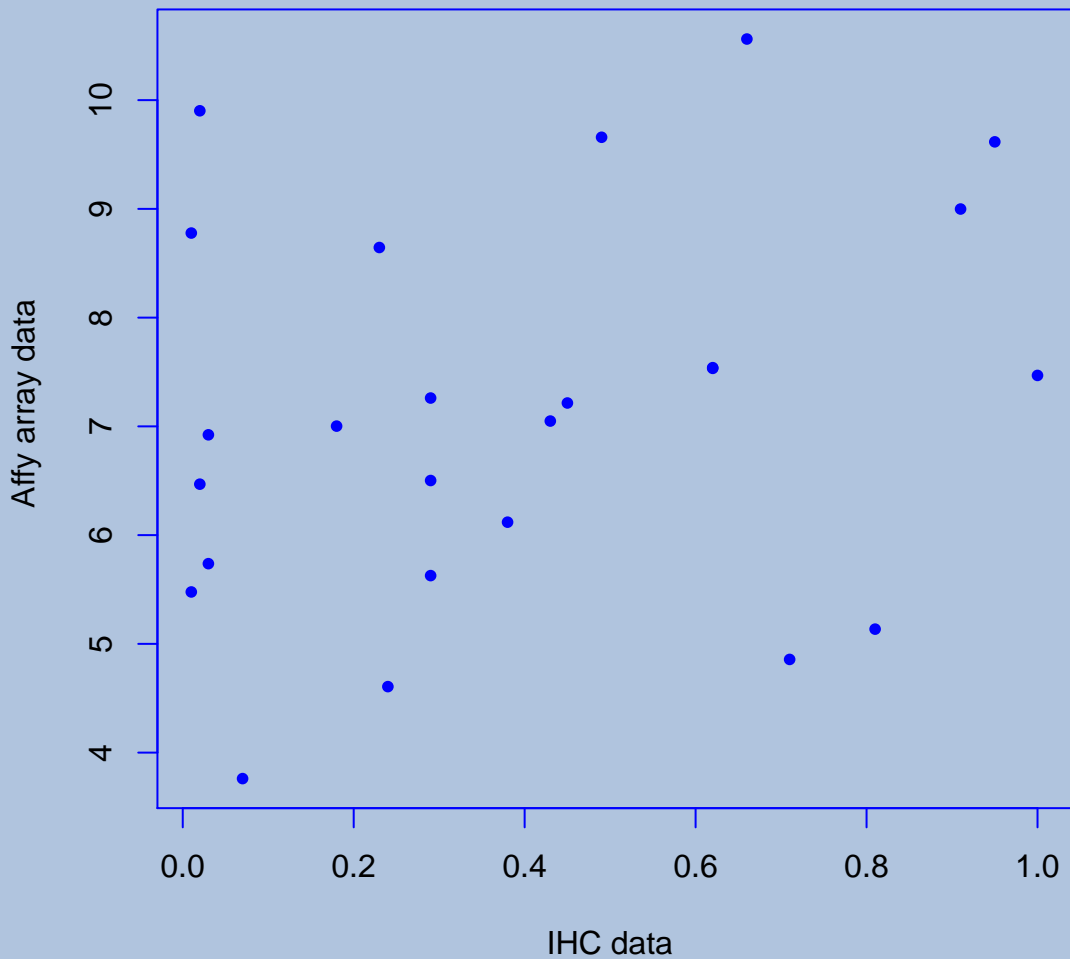
Basal_Epithelial_CD104 , median
spearman = 0.35 , pearson = 0.31



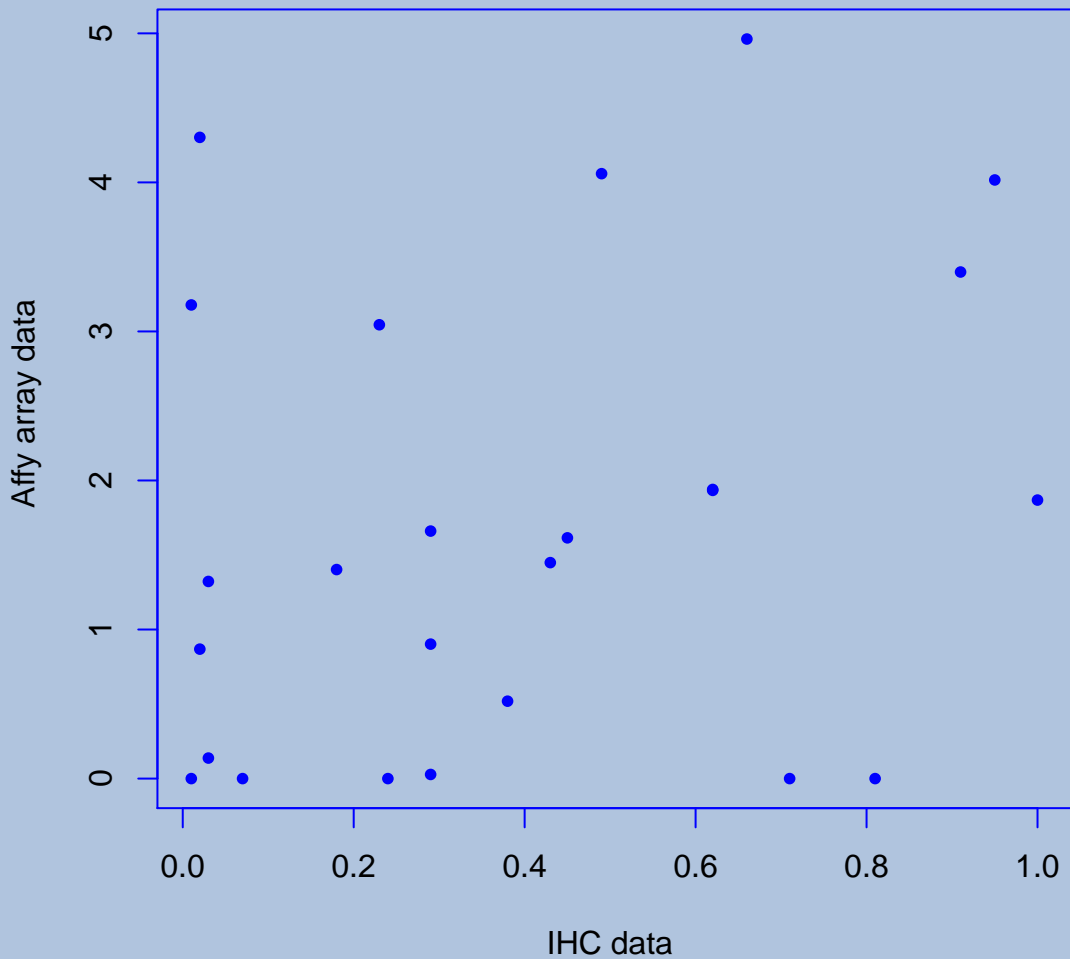
**Basal_Epithelial_CD104 , mean
spearman = 0.42 , pearson = 0.34**



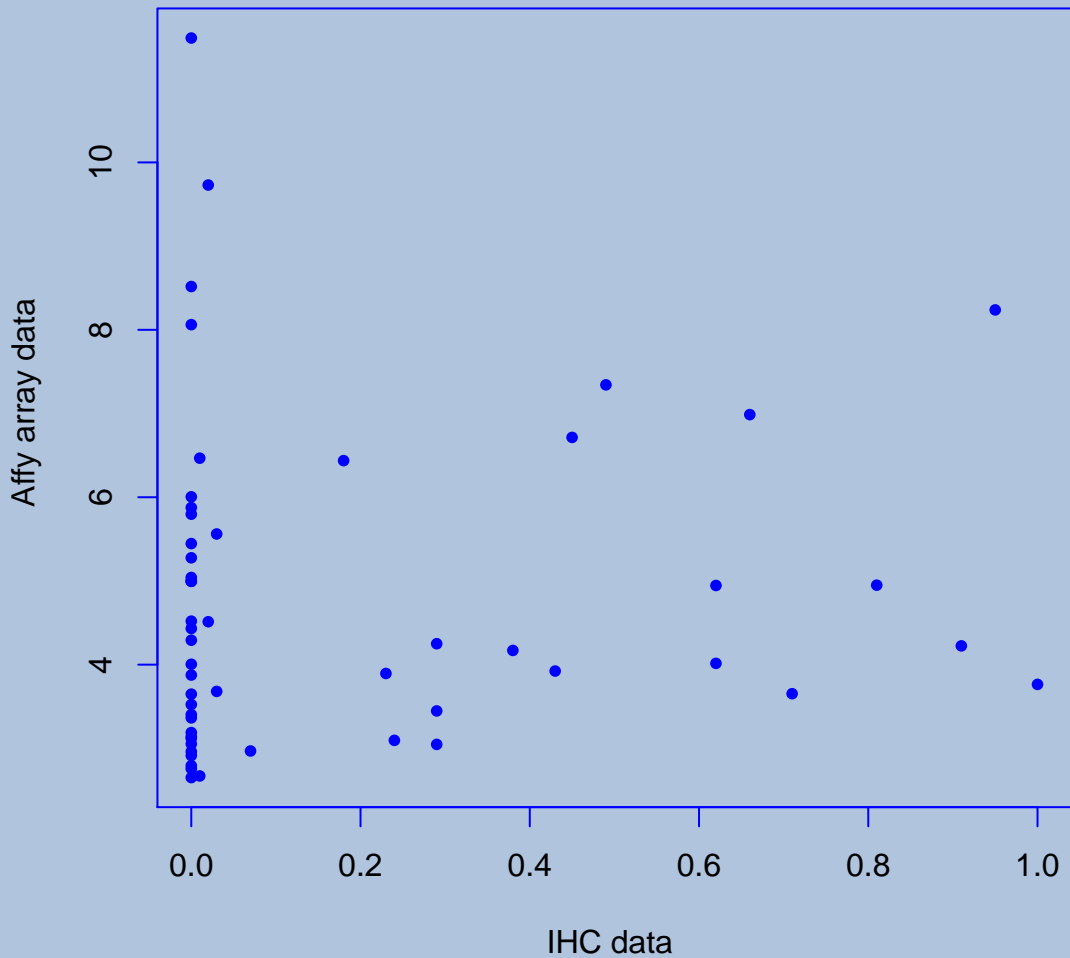
Basal_Epithelial_CD104 , no_zeros
spearman = 0.25 , pearson = 0.26



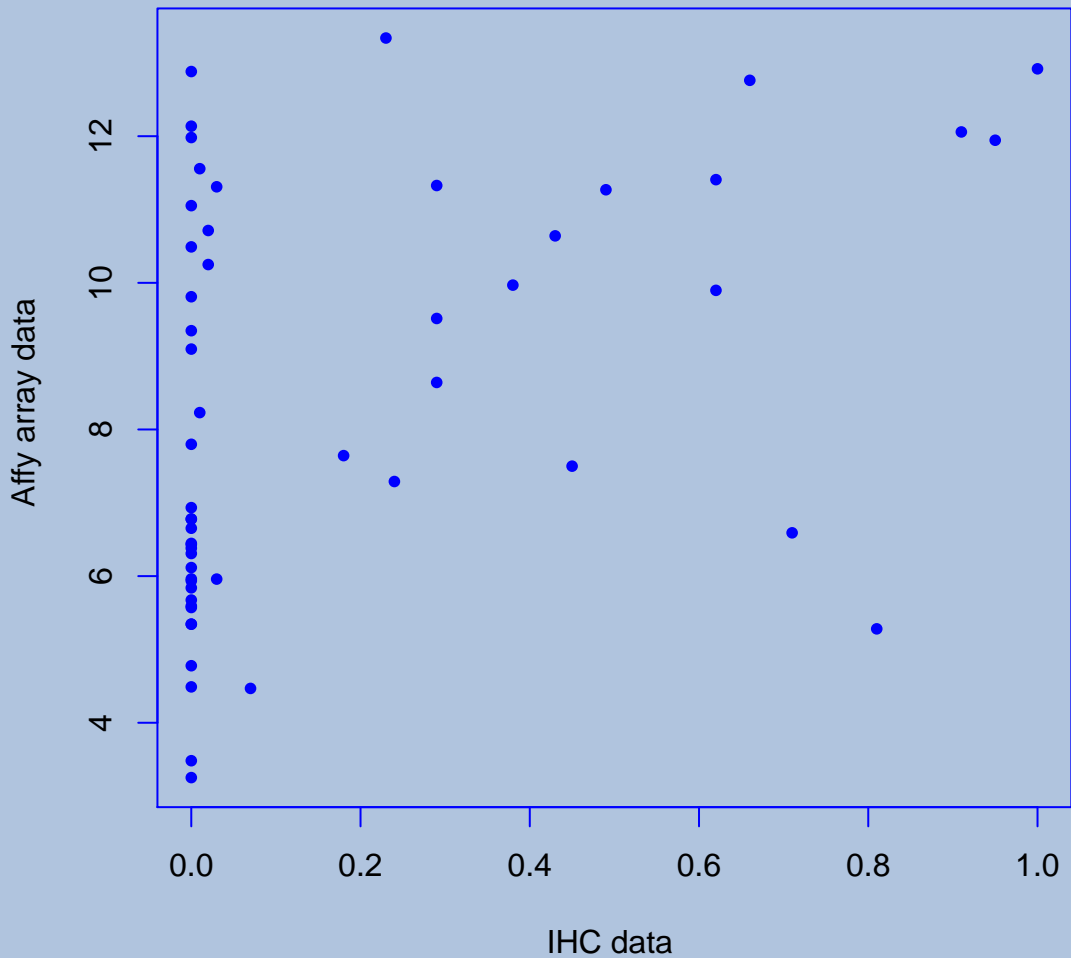
Basal_Epithelial_CD104 , xform
spearman = 0.25 , pearson = 0.27



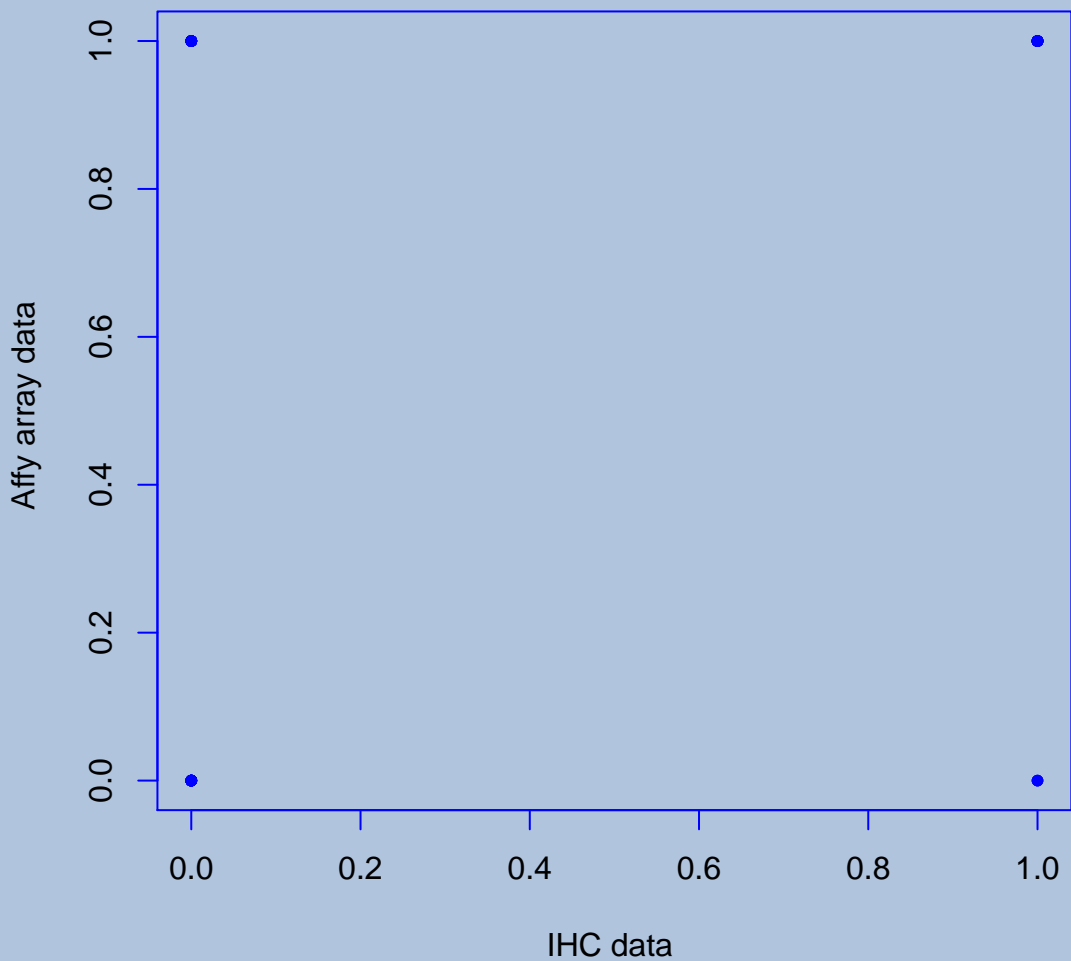
Basal_Epithelial_CD104 , min
spearman = 0.14 , pearson = 0.09



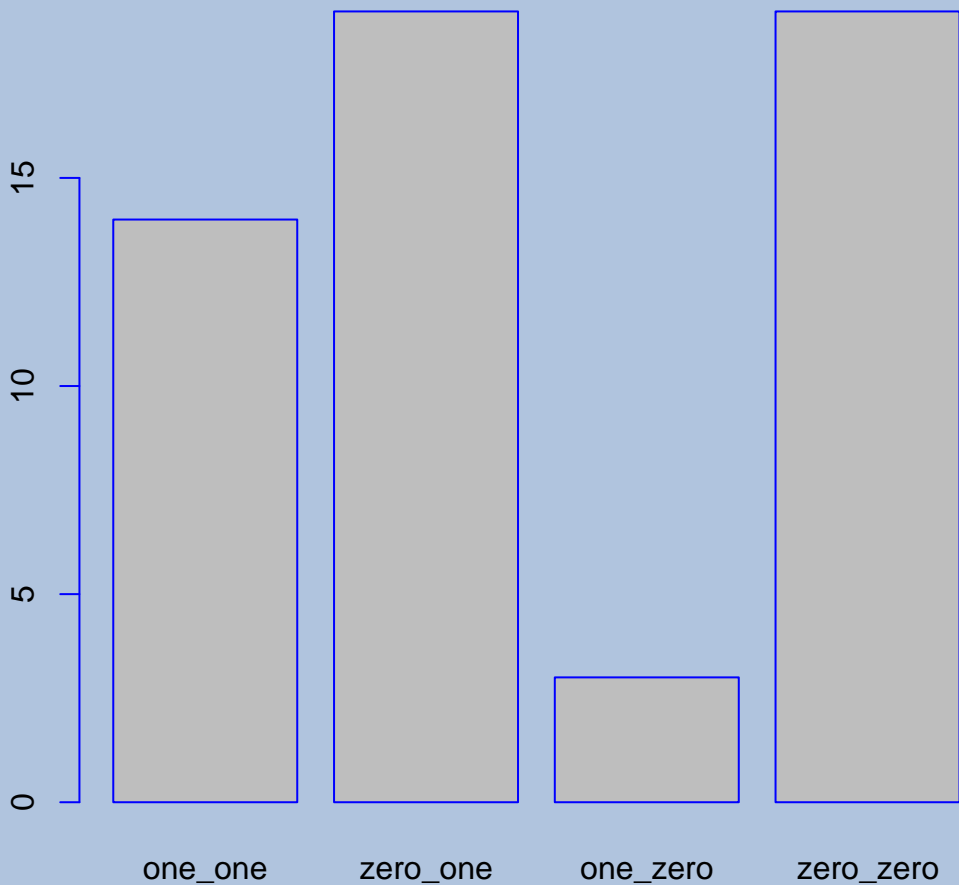
**Basal_Epithelial_CD104 , max
spearman = 0.46 , pearson = 0.42**



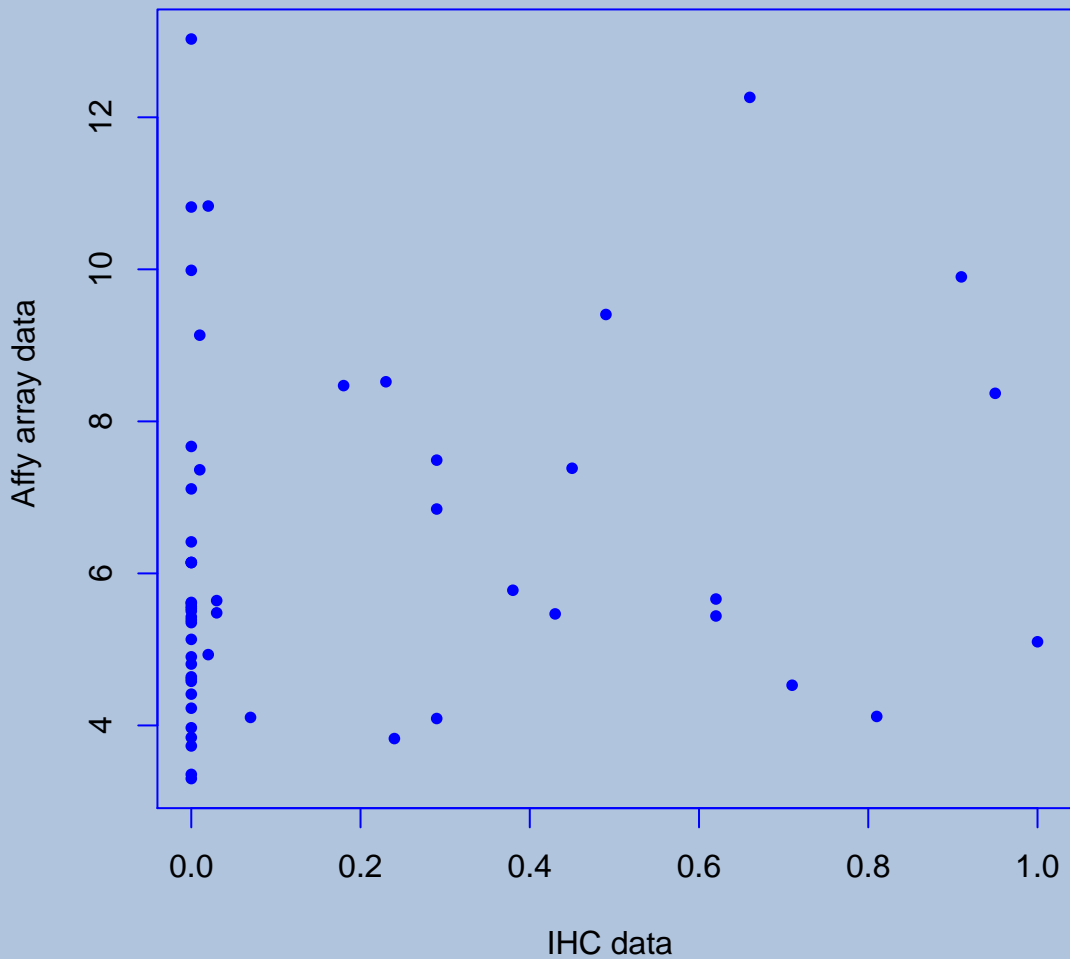
Basal_Epithelial_CD104 , digitized
spearman = 0.31 , pearson = 0.31



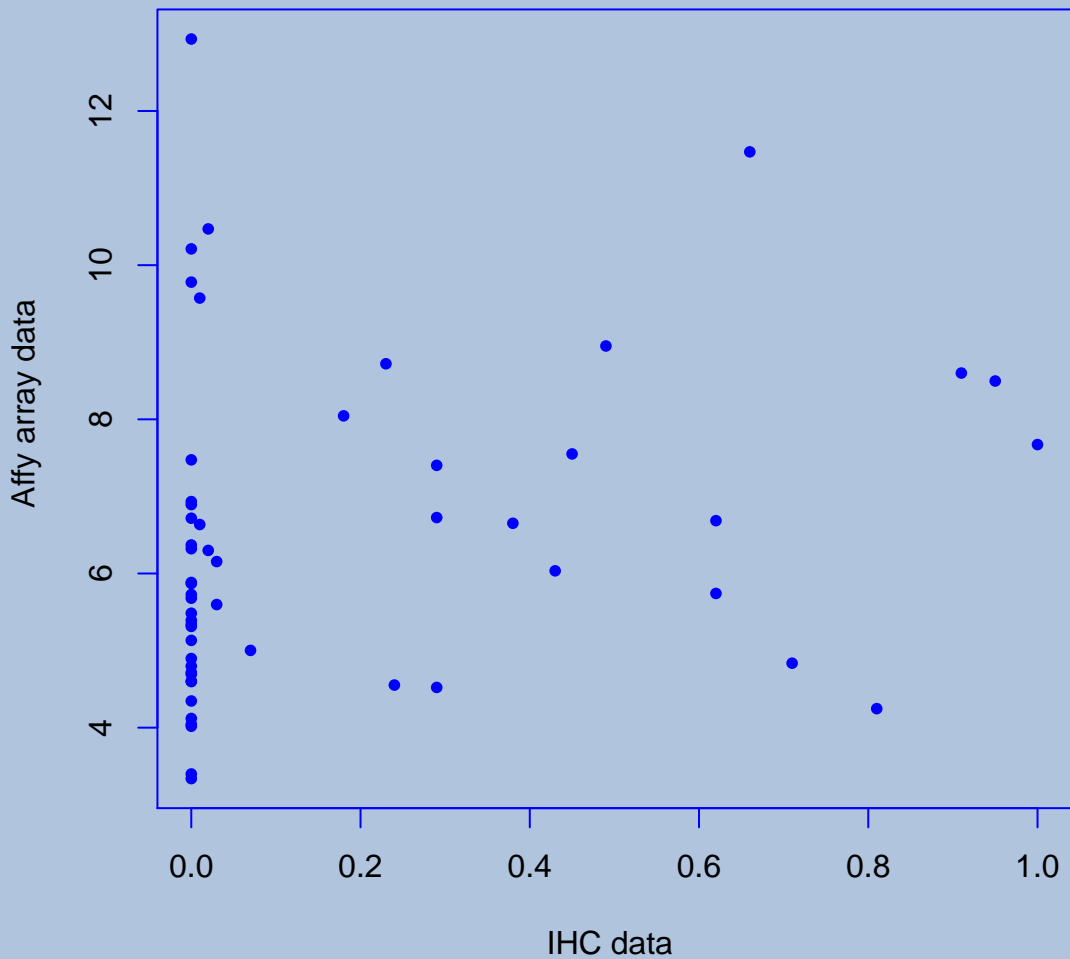
Basal_Epithelial_CD104



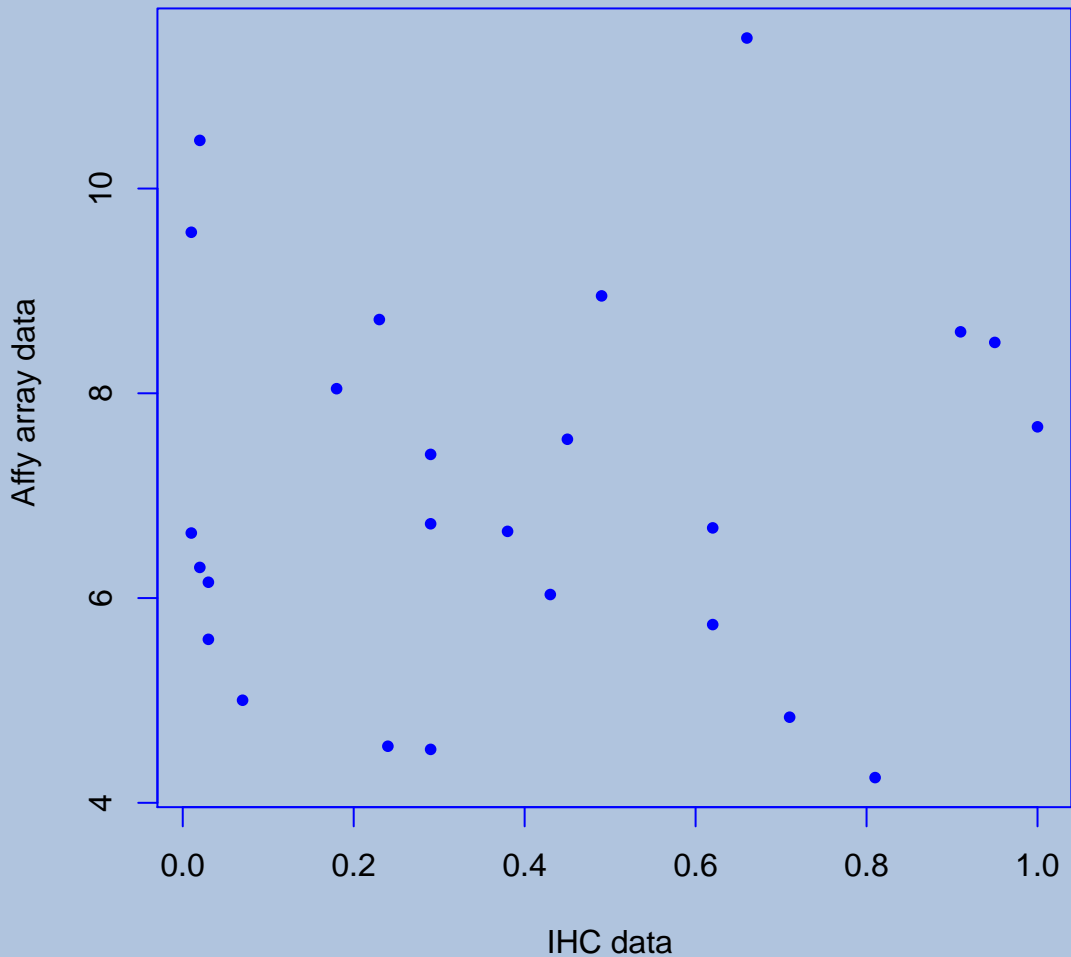
Progenitor_ABCG2 , median
spearman = 0.22 , pearson = 0.17



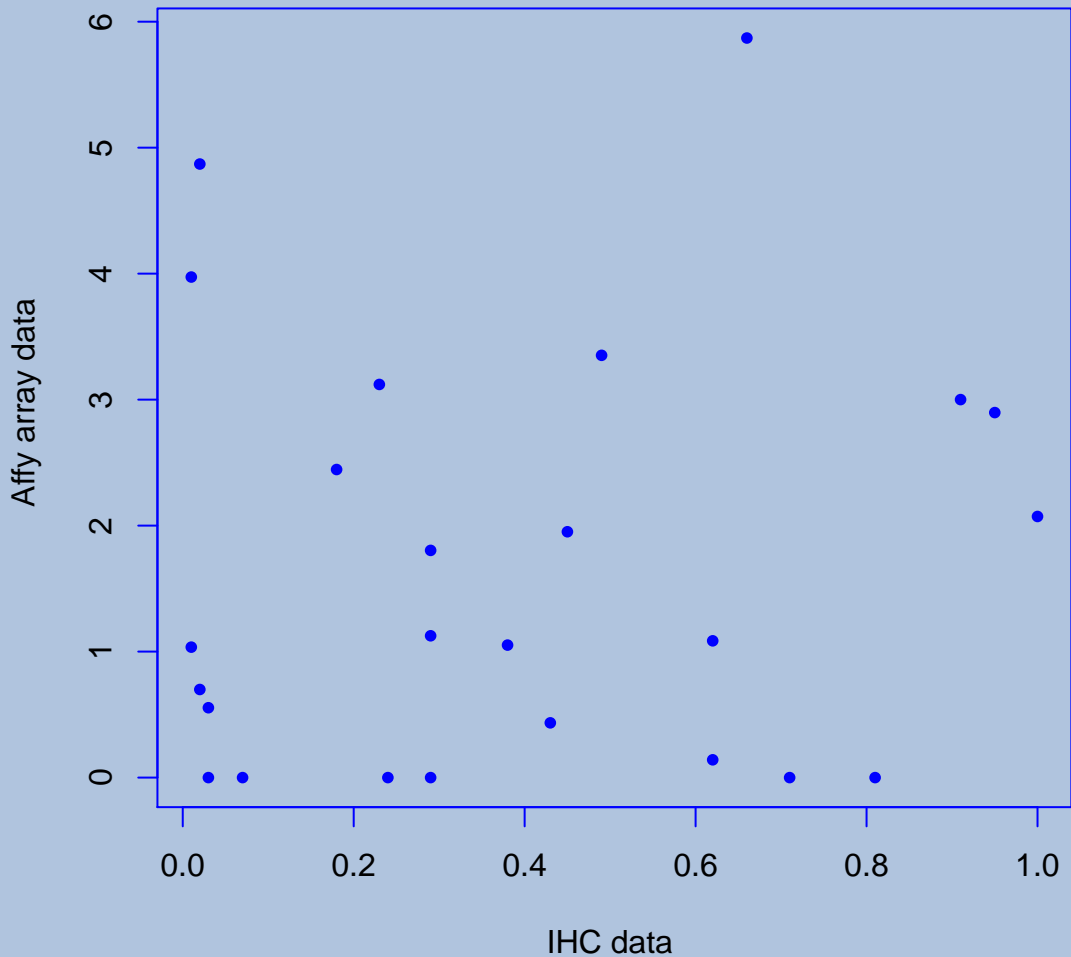
Progenitor_ABCG2 , mean
spearman = 0.34 , pearson = 0.23



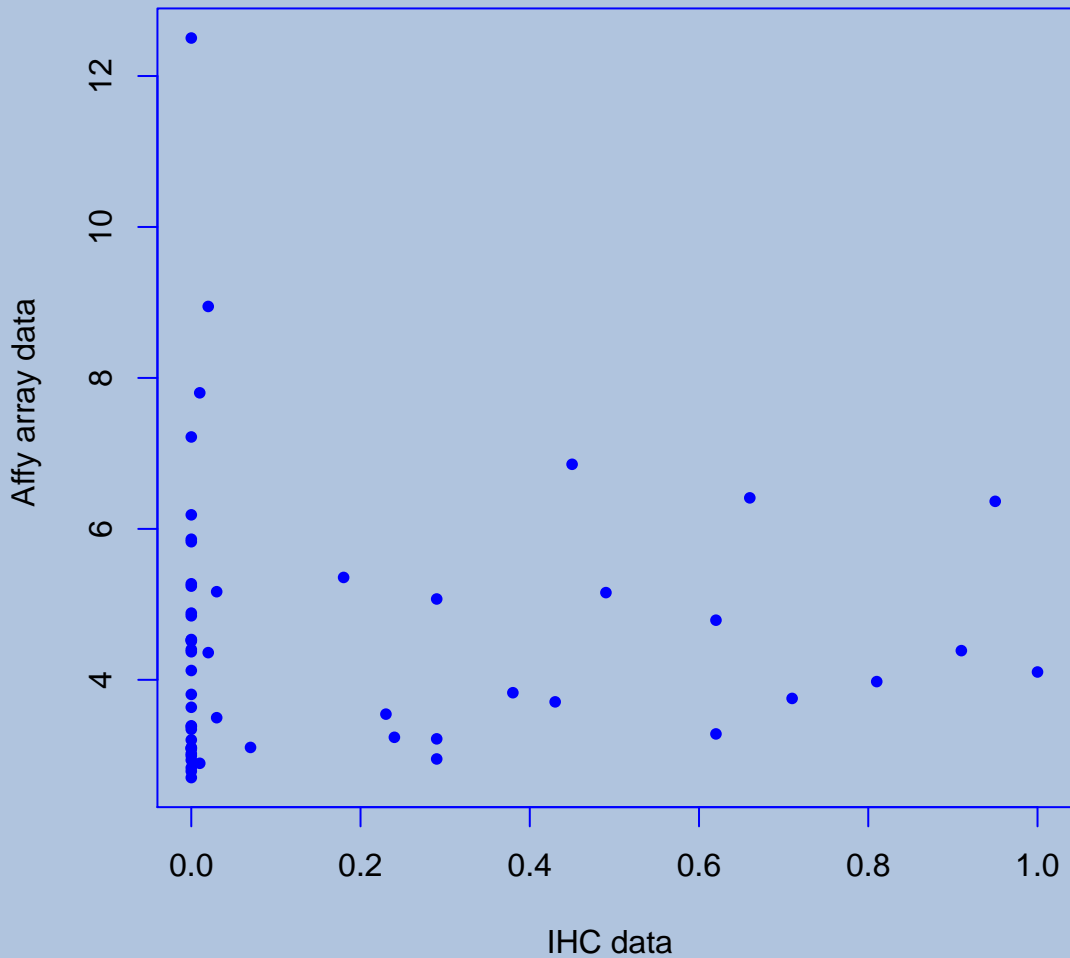
Progenitor_ABCG2 , no_zeros
spearman = 0.03 , pearson = 0.07



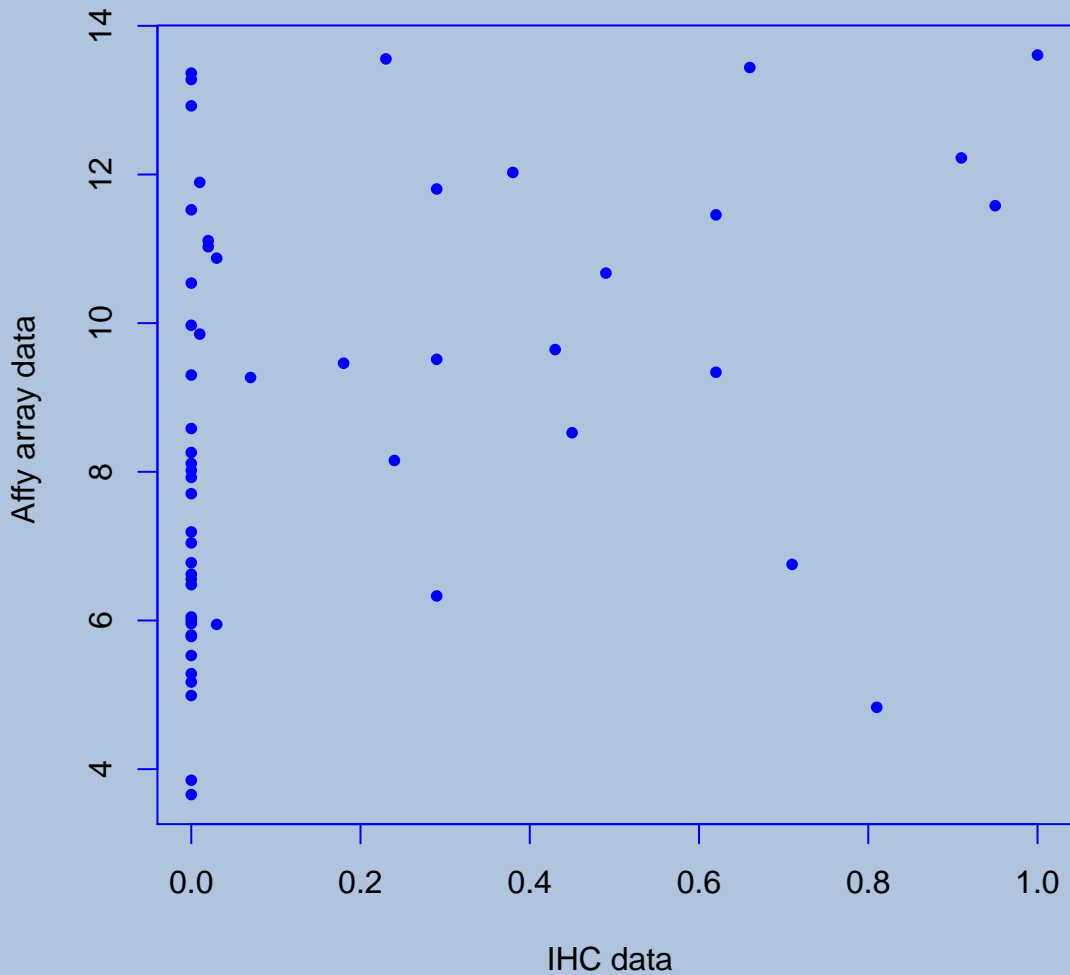
Progenitor_ABCG2 , xform
spearman = 0.06 , pearson = 0.10



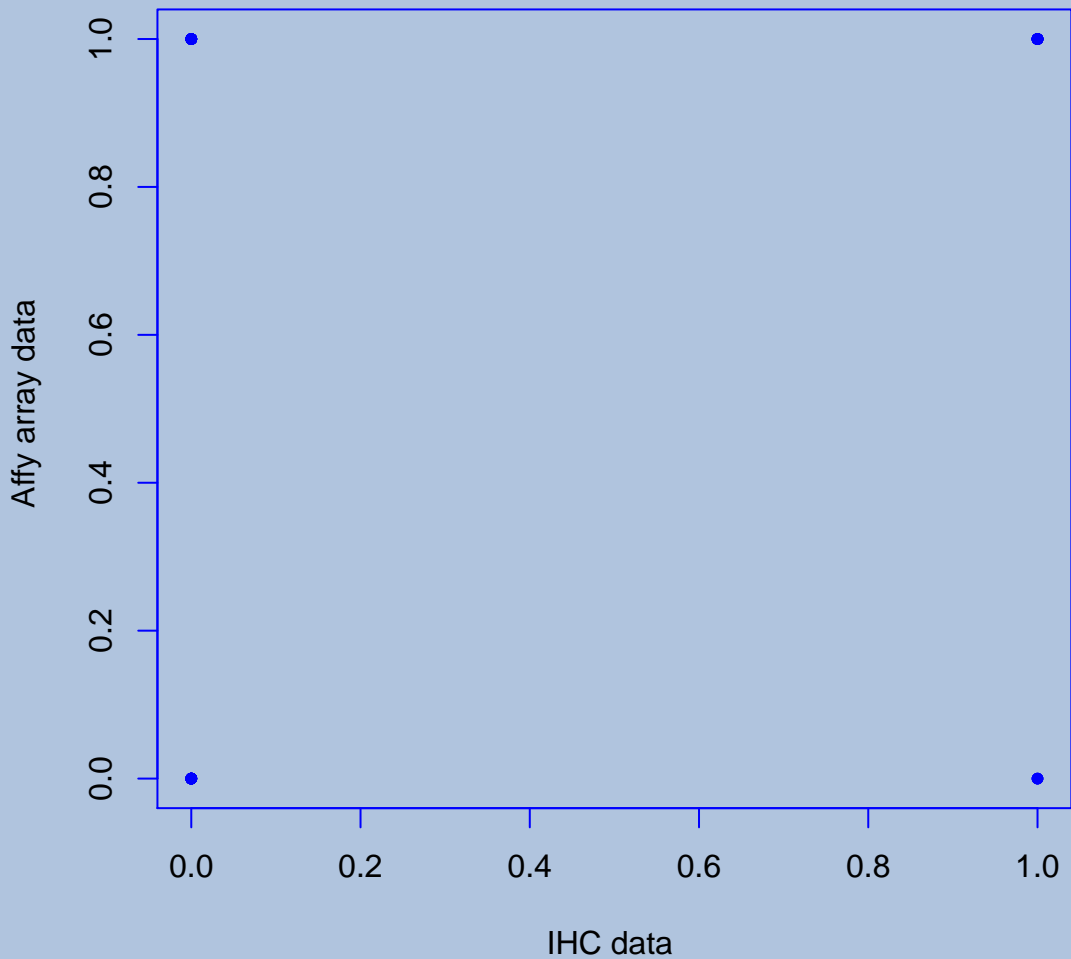
Progenitor_ABCG2 , min
spearman = 0.15 , pearson = 0.03



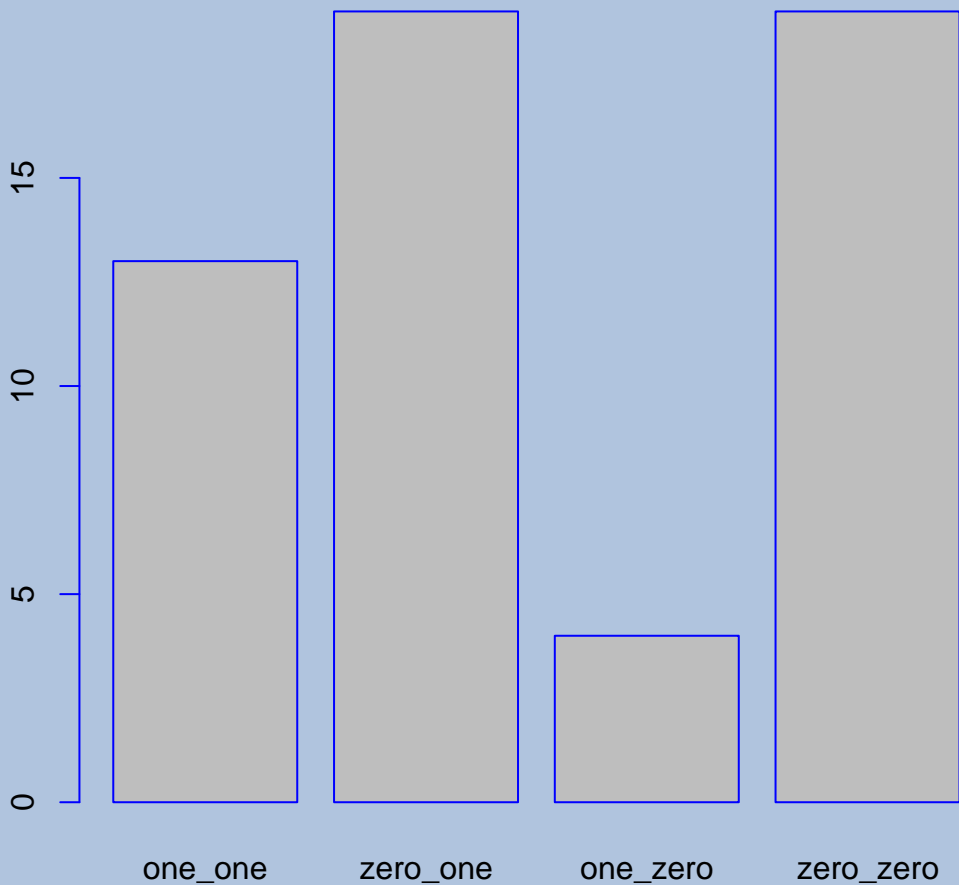
Progenitor_ABCG2 , max
spearman = 0.44 , pearson = 0.35



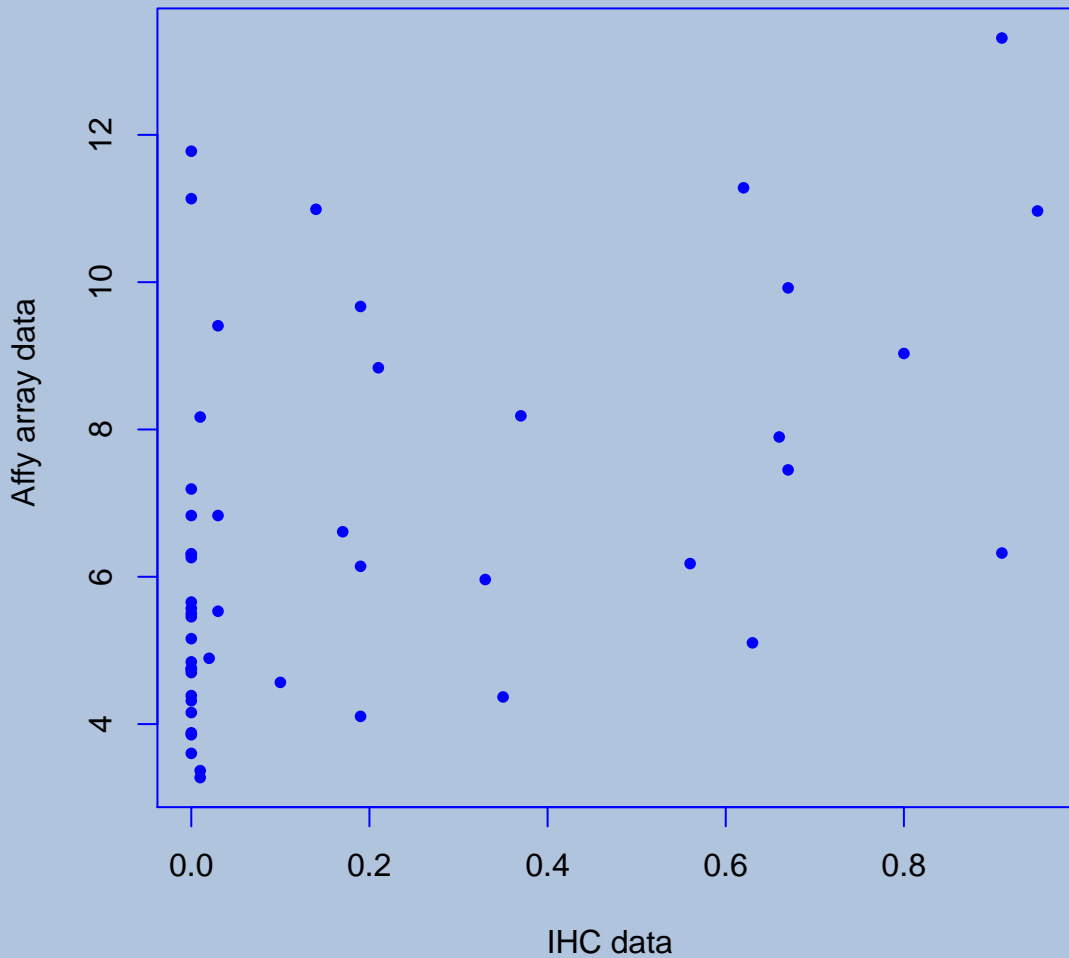
Progenitor_ABCG2 , digitized
spearman = 0.25 , pearson = 0.25



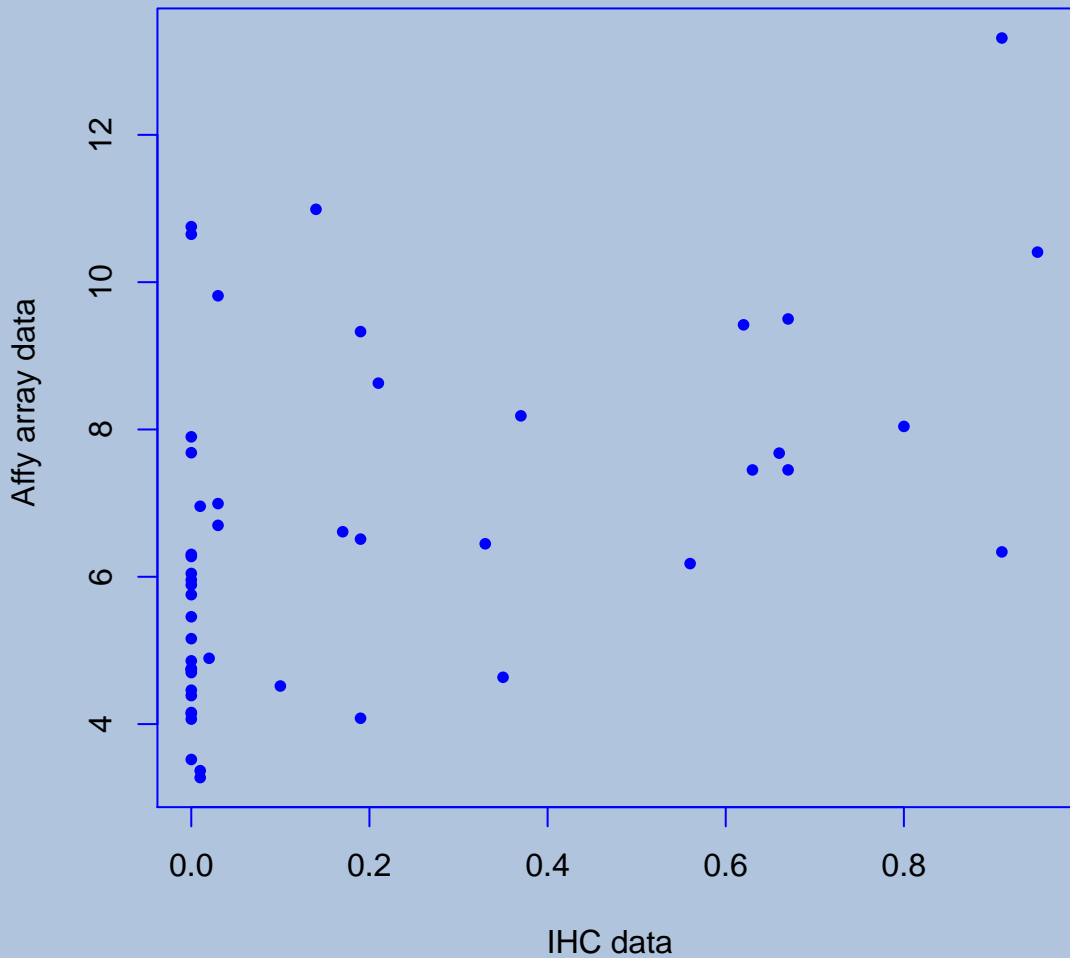
Progenitor_ABCG2



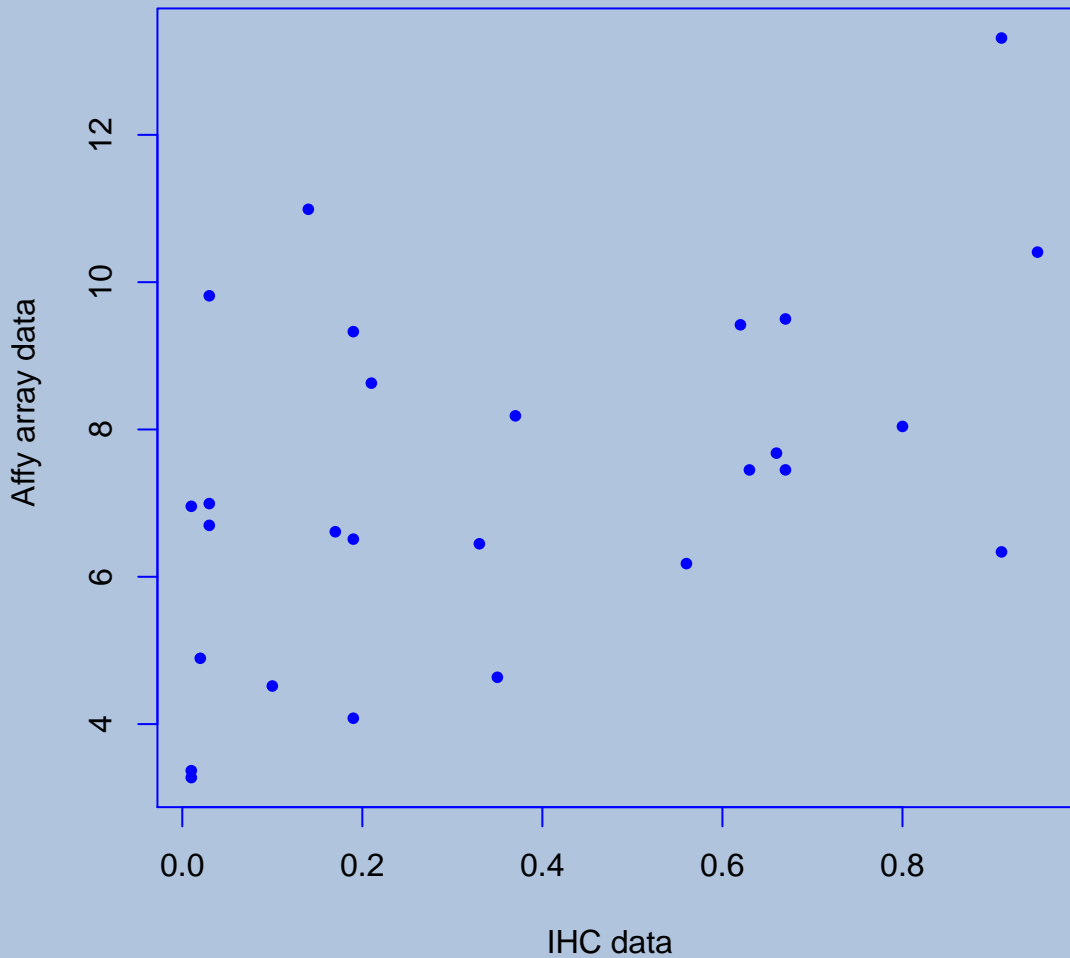
Cancer_CD26 , median
spearman = 0.45 , pearson = 0.49



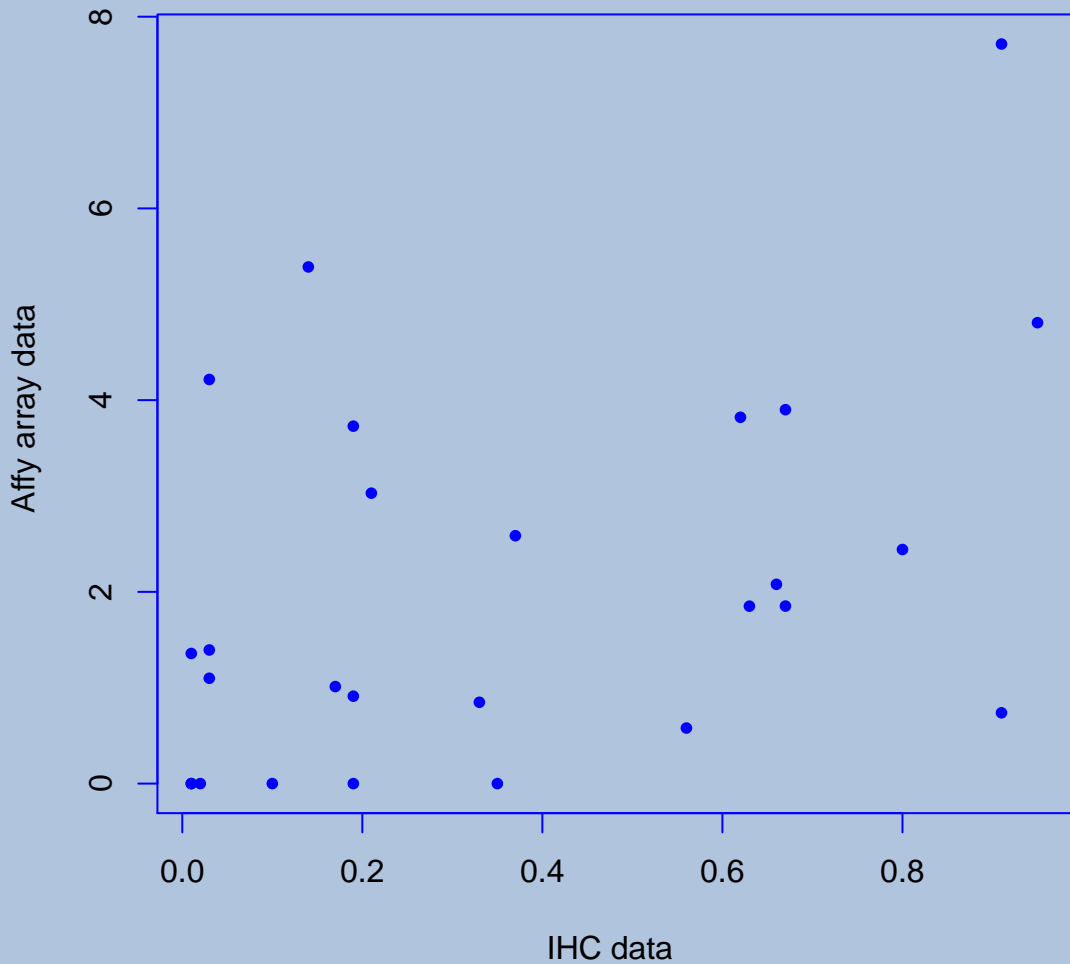
Cancer_CD26 , mean
spearman = 0.49 , pearson = 0.50



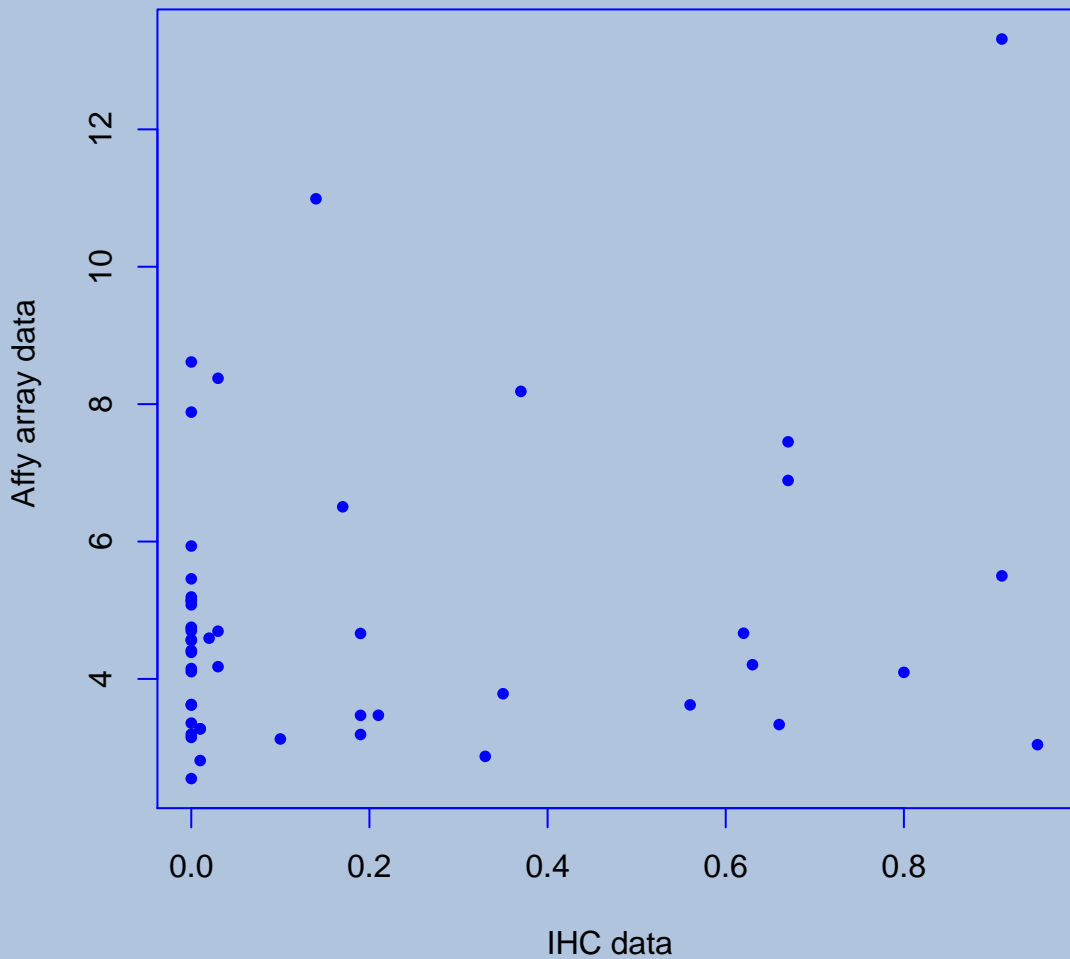
Cancer_CD26 , no_zeros
spearman = 0.46 , pearson = 0.48



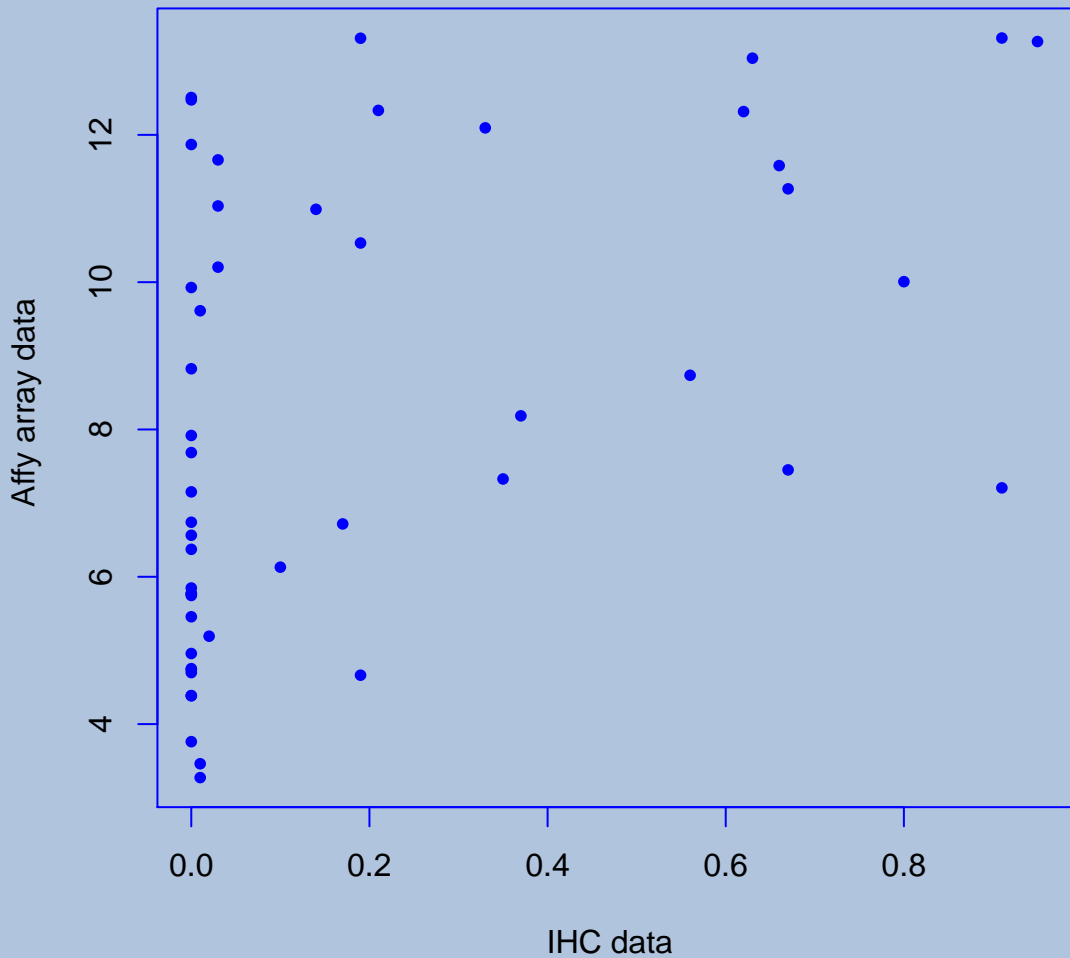
Cancer_CD26 , xform
spearman = 0.44 , pearson = 0.44



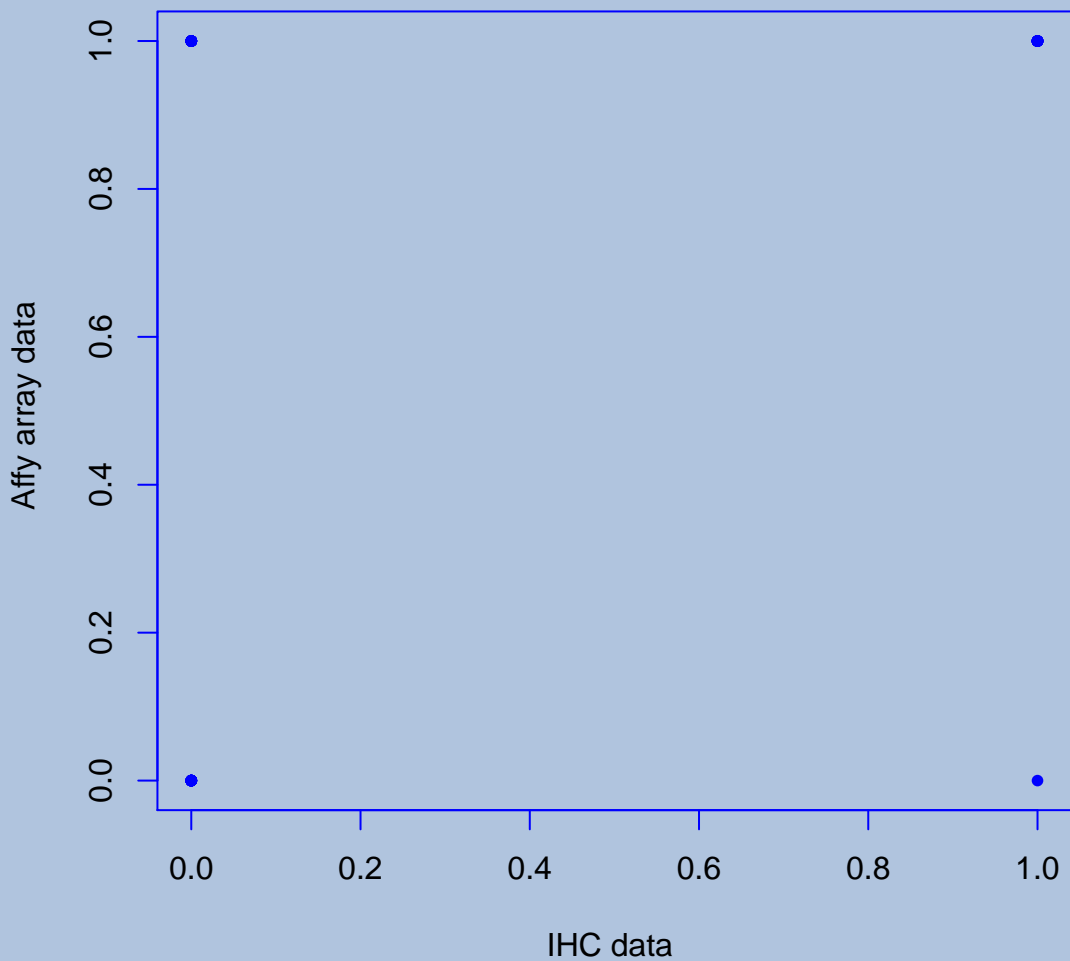
Cancer_CD26 , min
spearman = -0.01 , pearson = 0.21



Cancer_CD26 , max
spearman = 0.51 , pearson = 0.48



Cancer_CD26 , digitized
spearman = 0.40 , pearson = 0.40



Cancer_CD26

