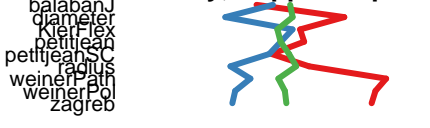
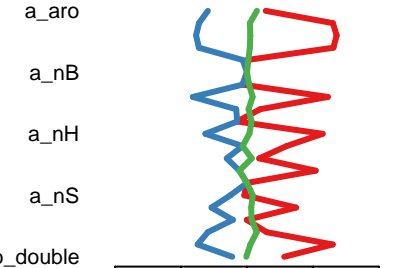


(EII,3,6015) Adjacency, distance matrix, connectivity, Kappa shape in



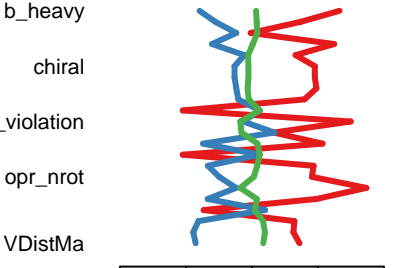
-3.0 -1.5 0.0 1.5 3.0

(EII,3,6015) Atom and bond counts



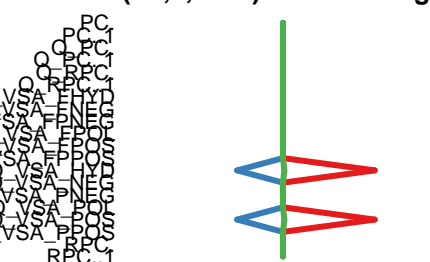
-3.0 -1.5 0.0 1.5 3.0

(EII,3,6015) Atom and bond counts



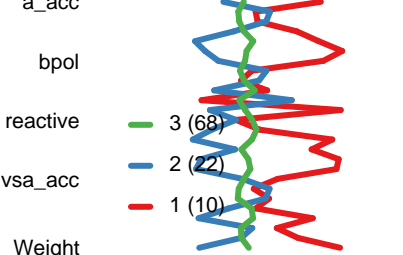
-3.0 -1.5 0.0 1.5 3.0

(EII,3,6015) Partial charge

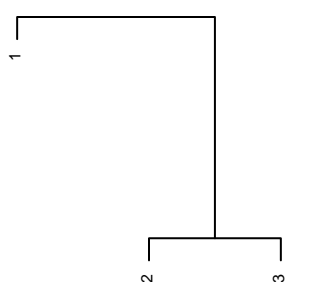
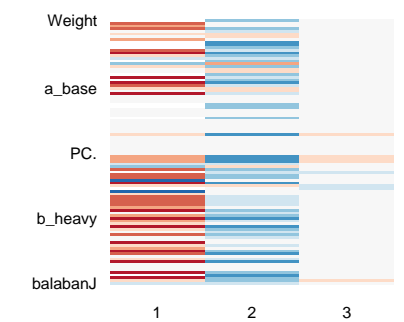


-3.0 -1.5 0.0 1.5 3.0

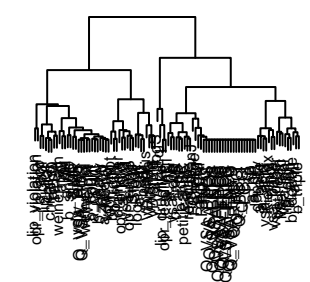
(EII,3,6015) Physical proper and pharmacophore feature descriptors



-3.0 -1.5 0.0 1.5 3.0



dist(avg_p)
hclust (*, "complete")



dist(t(avg_p))
hclust (*, "complete")