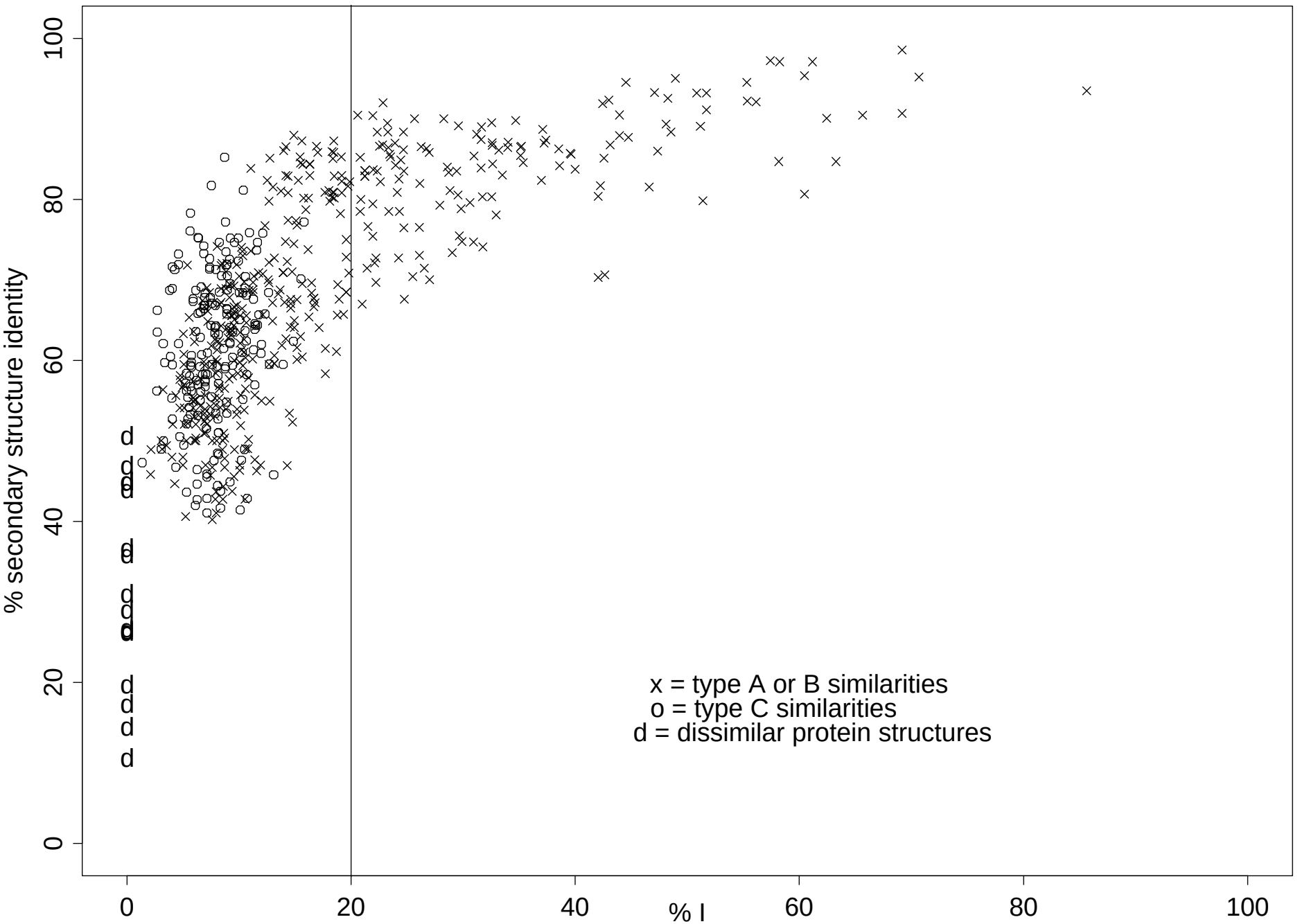
% secondary structure identity
0
20
40
60
80
100
% I
x = type A or B similarities
o = type C similarities
d = dissimilar protein structures