

Figure 1: Disconnectivity Graph for the BLN-46 model protein, $E_{threshold} = -48.0\epsilon$, $N_{sp} = 1341$.

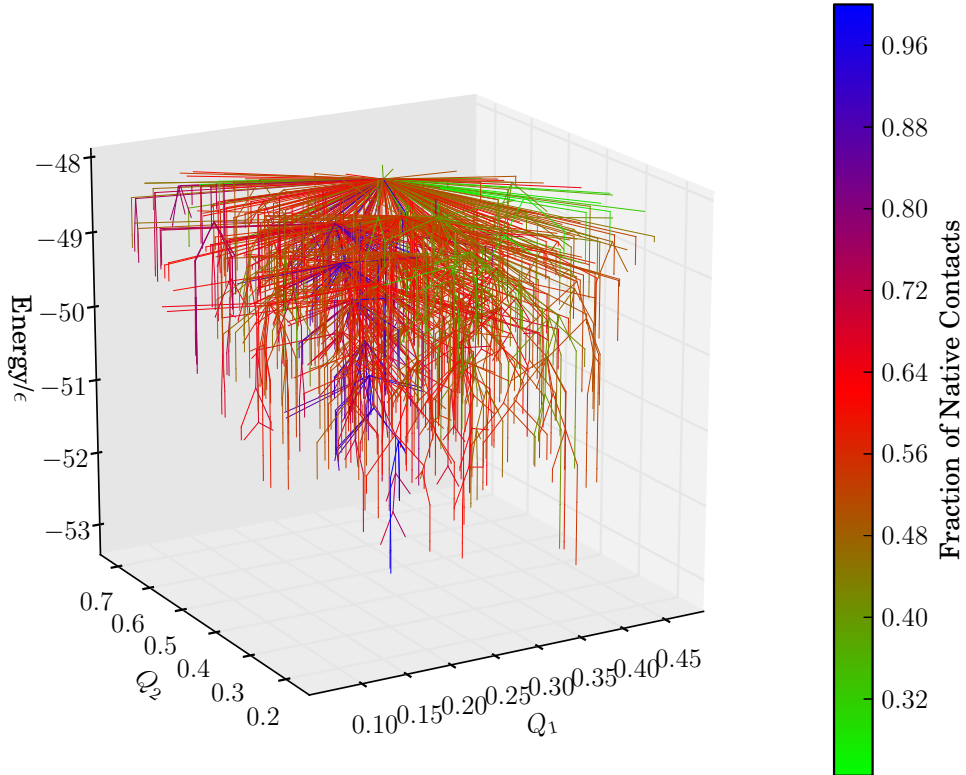


Figure 2: 3D Metric Disconnectivity Graph for the BLN-46 model protein, $E_{threshold} = -48.0\epsilon$, $N_{sp} = 1341$. Ordered according to first two principal components Q_1 and Q_2 , which capture 24% and 14% of variance respectively.