Cluster evaluation - Spec2Vec - dilation

Legend:
- Spec2Vec, cutoff = 0.9
- Spec2Vec, cutoff = 0.9 + dilation (cutoff=0.85)
- Spec2Vec, cutoff = 0.9 + 2x dilation (cutoff=0.85 + 0.8)
- Spec2Vec, cutoff = 0.9 + 3x dilation (cutoff=0.85 + 2x 0.8)
- Spec2Vec, cutoff = 0.9 + 3x dilation (cutoff=0.85 + 3x 0.8)
- Spec2Vec, cutoff = 0.9 + 3x dilation (cutoff=0.85 + 3x 0.8 +

x-axis: cluster size (<2, 2-5, 6-10, 11-50, 51-100)
y-axis: mean molecular similarity of nodes in cluster