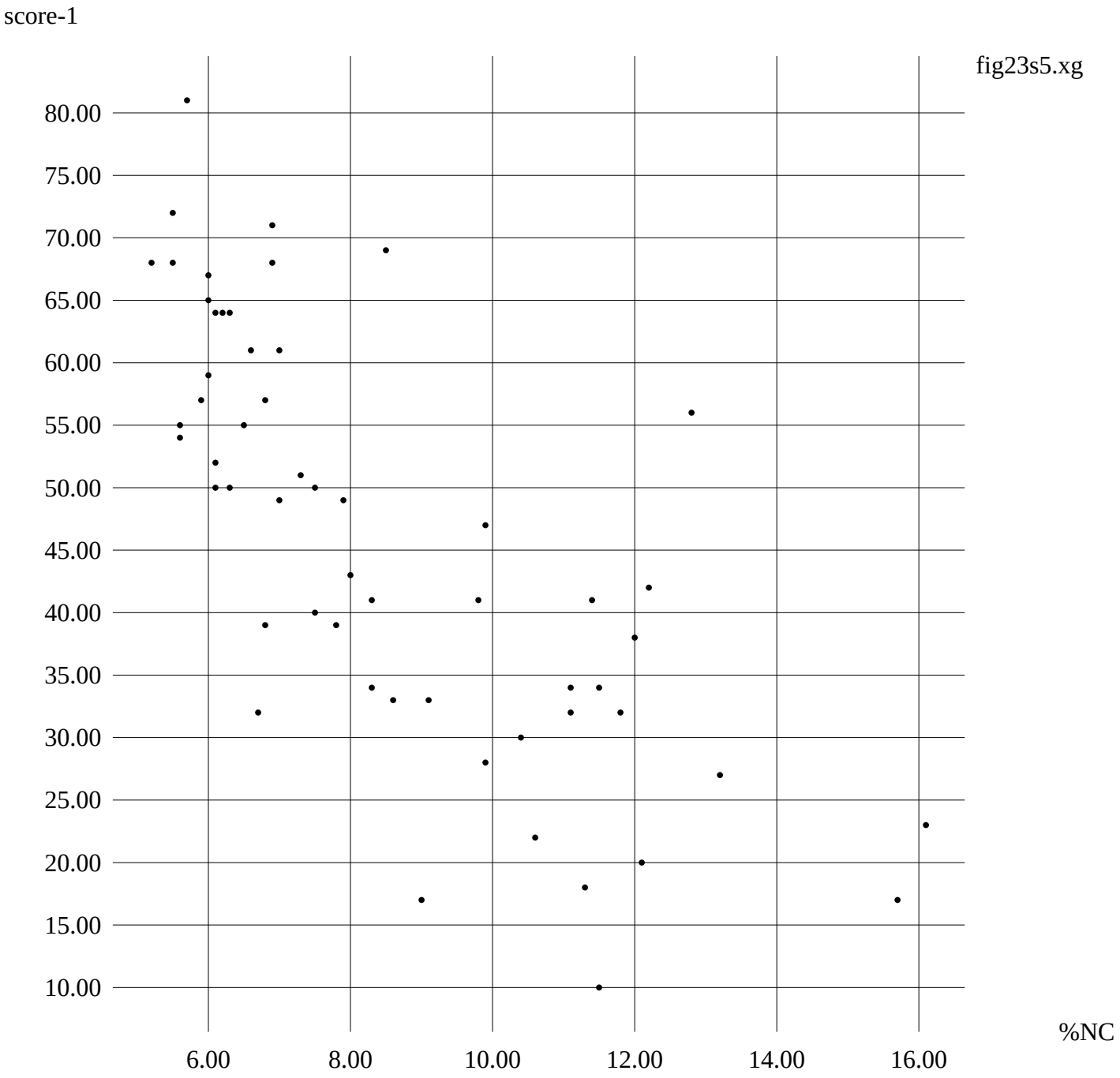Figure 5, 16S rRNA case: score-1 vs % non-canonical base pairs