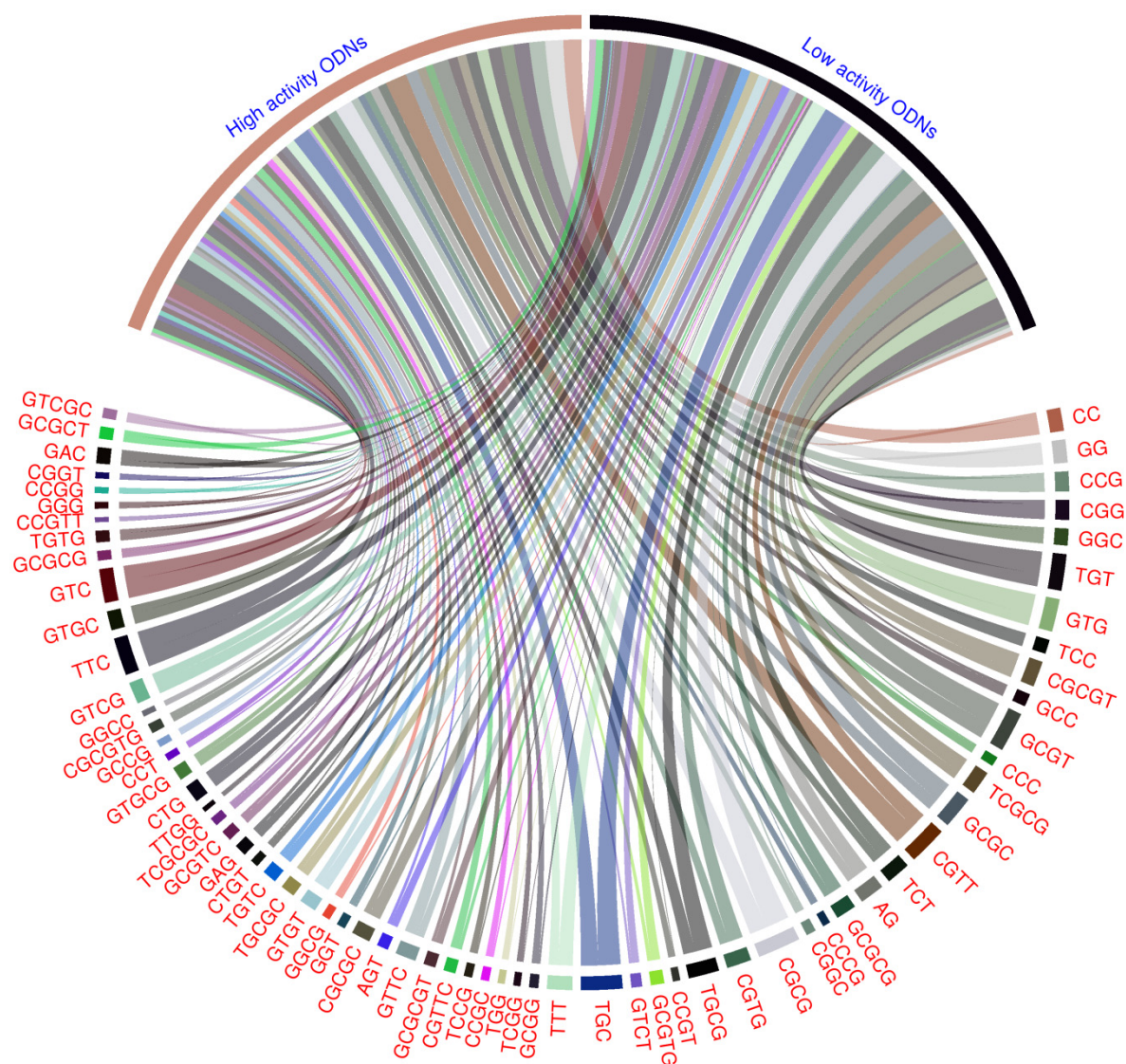


# Prediction of novel mouse TLR9 agonists using a random forest approach



**Additional File 1: Sequence motifs in mTLR9 active ODNs having an absolute difference in the occurrence above 10% in high and low activity groups of ODNs, arranged in a clockwise manner.** The width of the ribbon shows the average percentage composition of the motifs in each group.