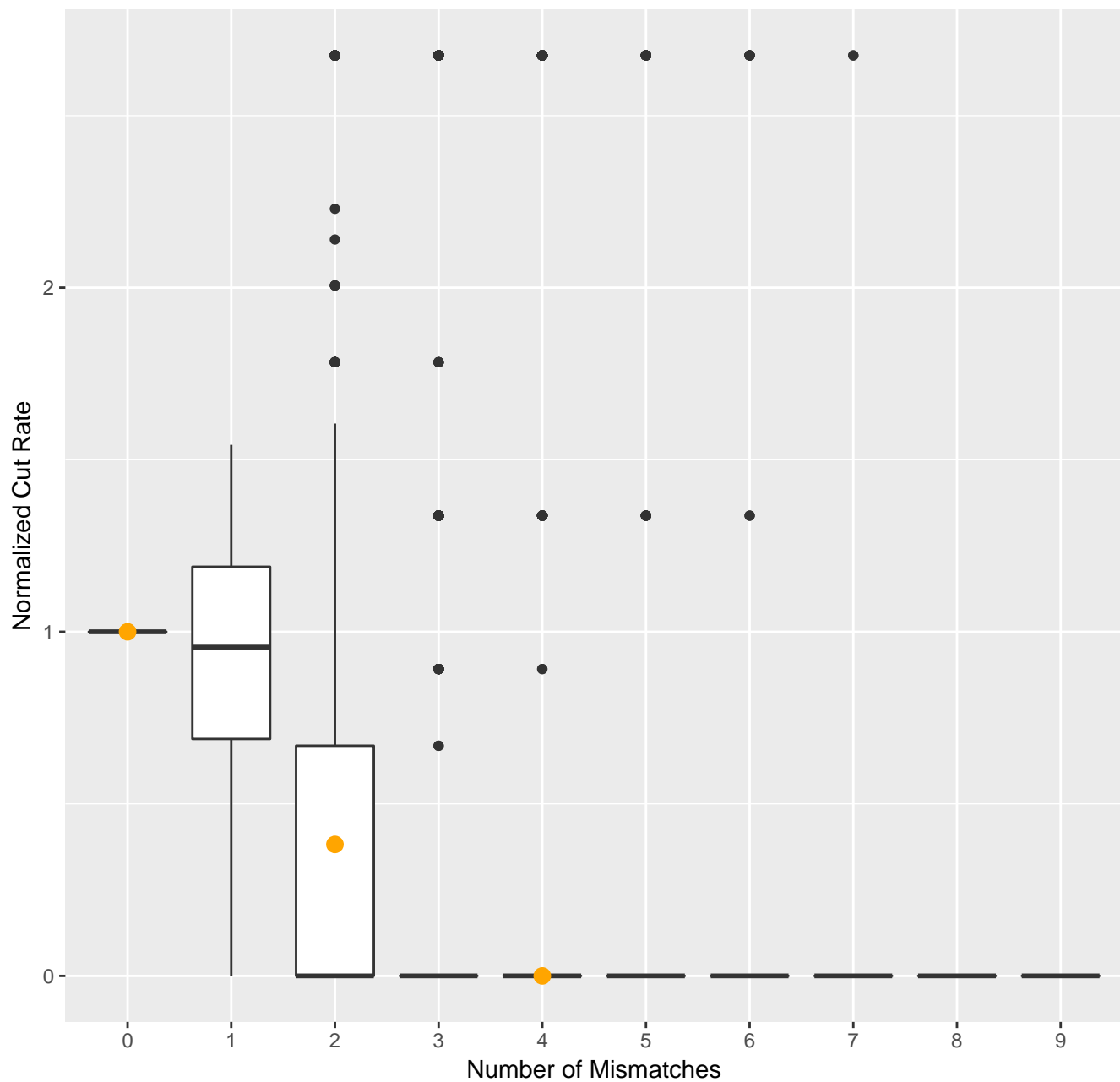Distribution of Normalized Cut Rate vs. Mismatches for 16hr+gDNA2

Cut Rate by Mean Mismatch Position in 16hr+gDNA2