

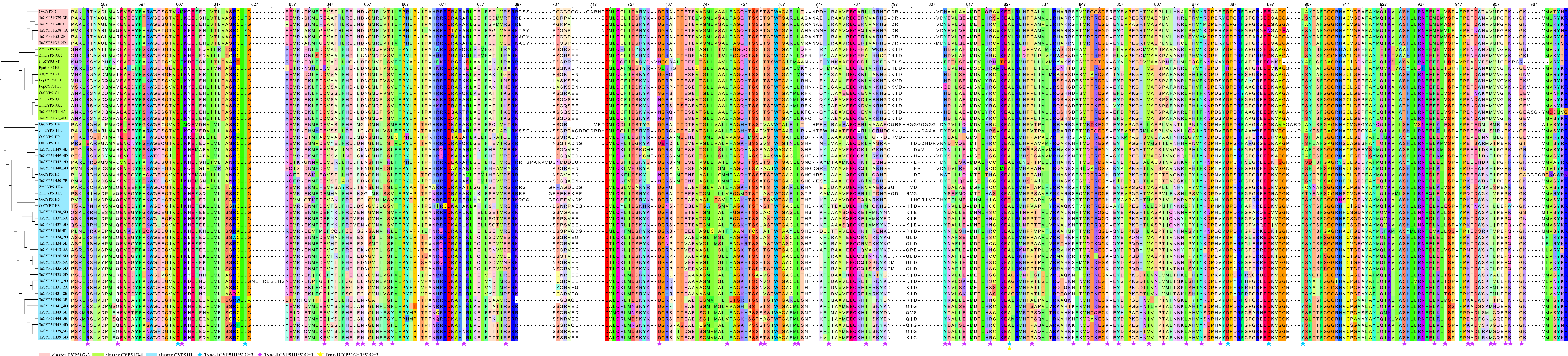


Figure S10. The multiple sequence alignment of CYP51 members. Sites whose posterior probability is larger than cutoff value (0.67) are marked in star or triangle.