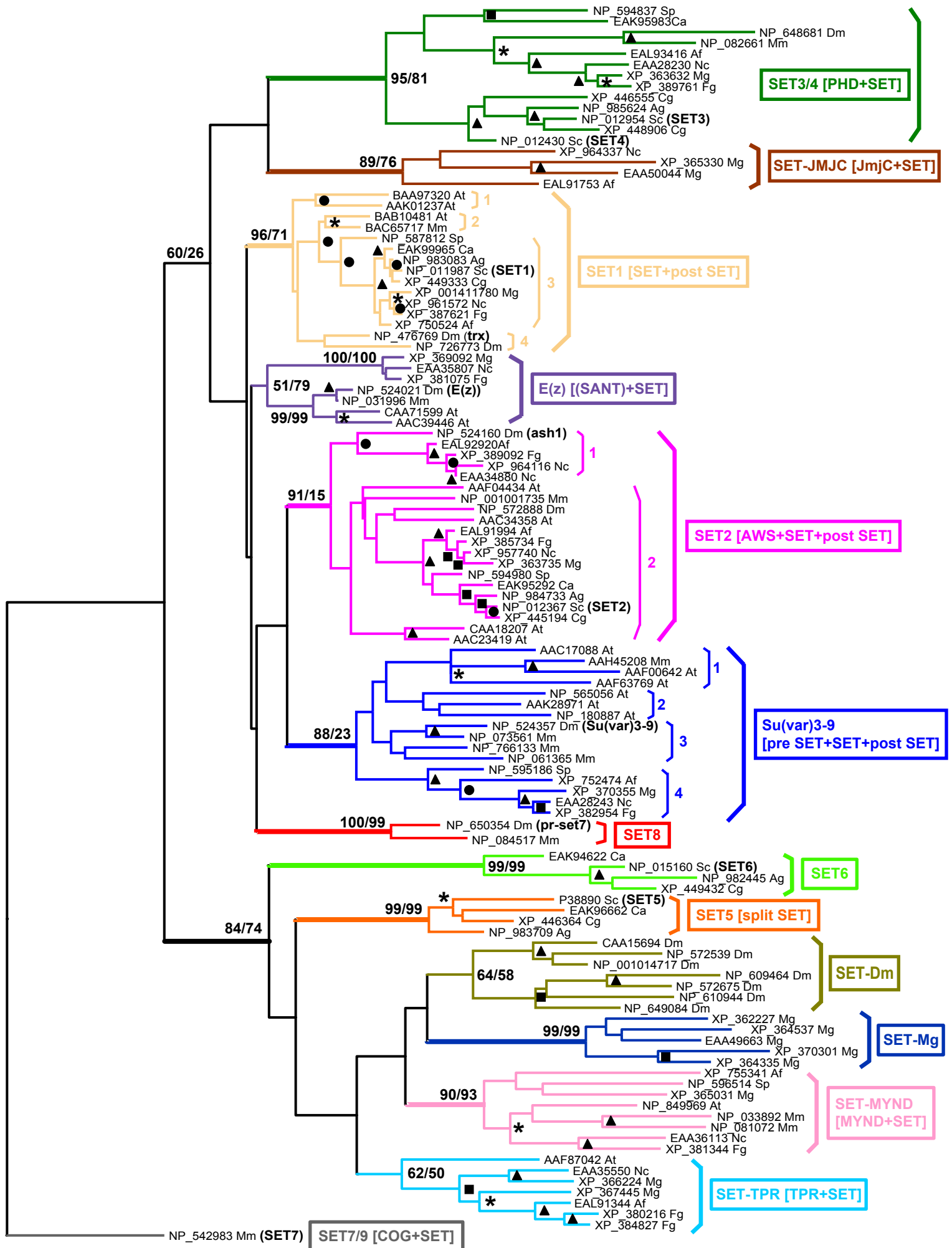




0.2 amino acid substitutions per site