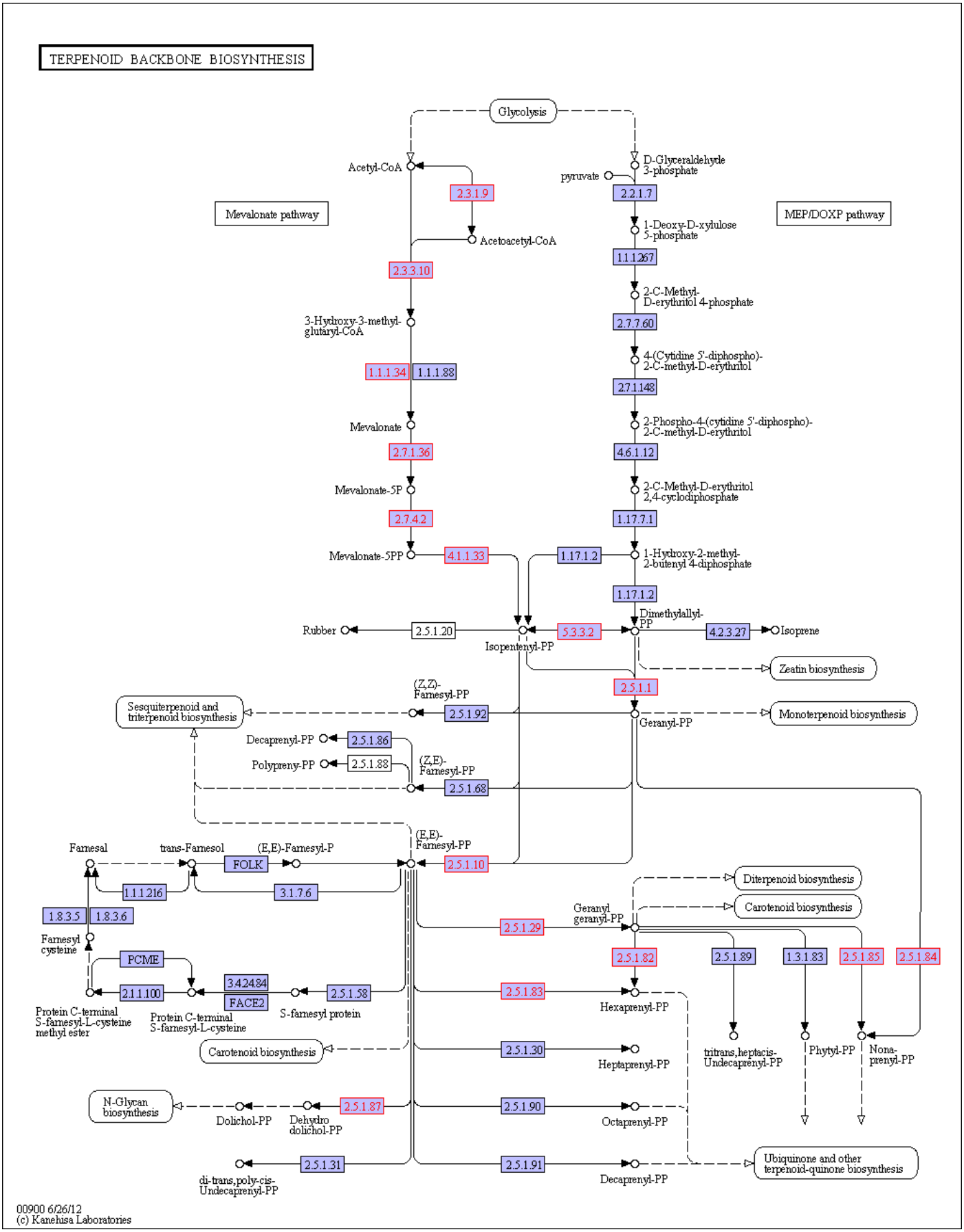

The terpenoid backbone biosynthesis pathway of *L. rhinocerotis* by KEGG mapper (http://www.genome.jp/kegg/mapper.html). Blue box marked in red indicates existing homologous genes of the enzyme.