



Figure S3. Metabolite–gene Pearson correlation network between DAMs and DEGs involved in phenylpropanoid biosynthesis (ko00940) in P3CR vs. P3R (A) and P5CR vs. P5R (B). Metabolite–gene pairs were connected by edges; red and green lines represent positive and negative correlation, respectively; yellow and blue nodes represent DAMs and DEGs, respectively. ↓, downregulation; ↑, upregulation.