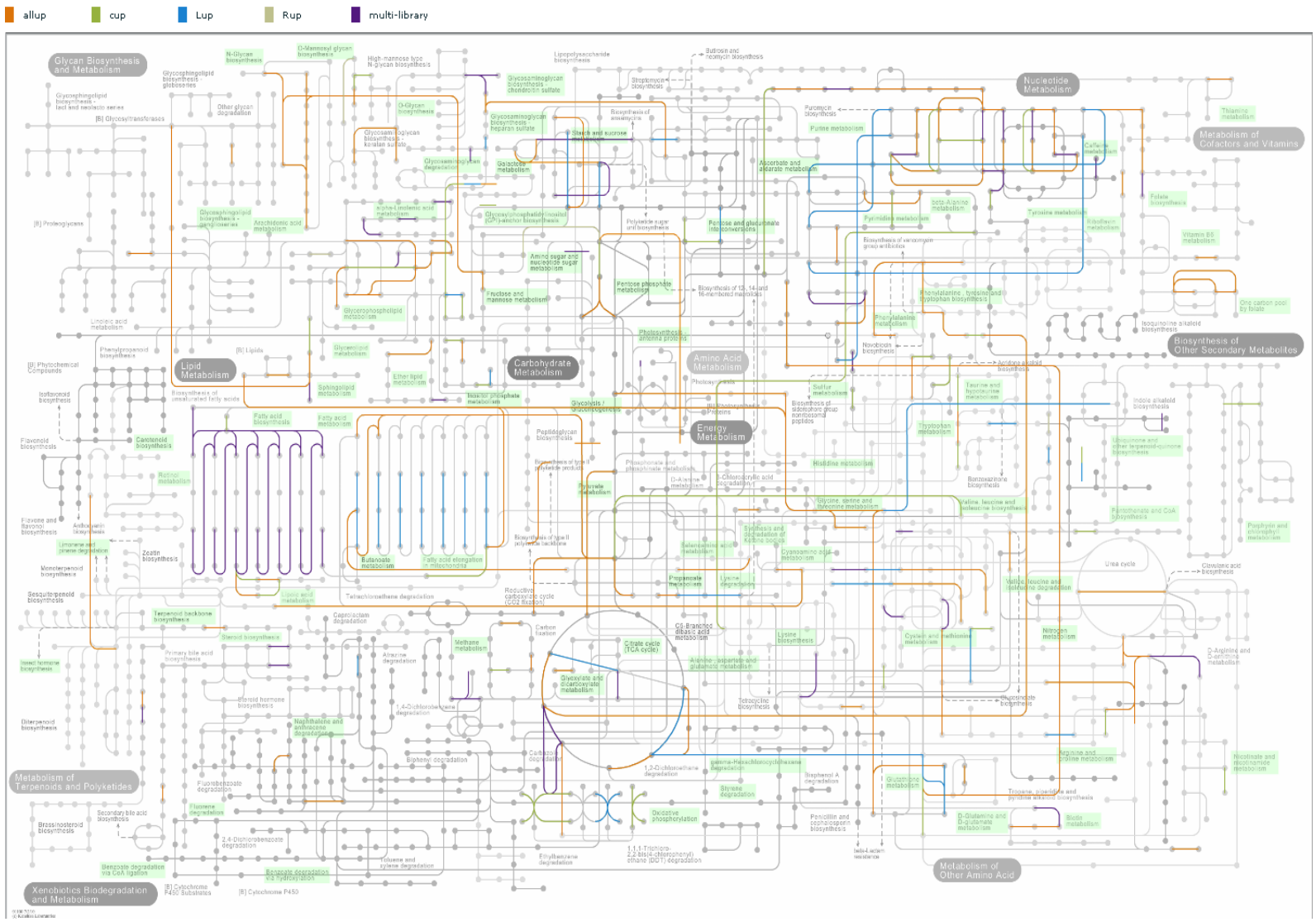

Figure S8 Metabolic pathways corresponding to transcripts increased in accumulation 5hPBM in all three strains (allup), in LVP (Lup), in CTM (cup) and in Rex-D (Rup) are visualized by LinkinPath (Ingsriswang et al., 2011). Multi-library corresponds to pathways elicited by transcripts increased in accumulation in more than one tested condition.