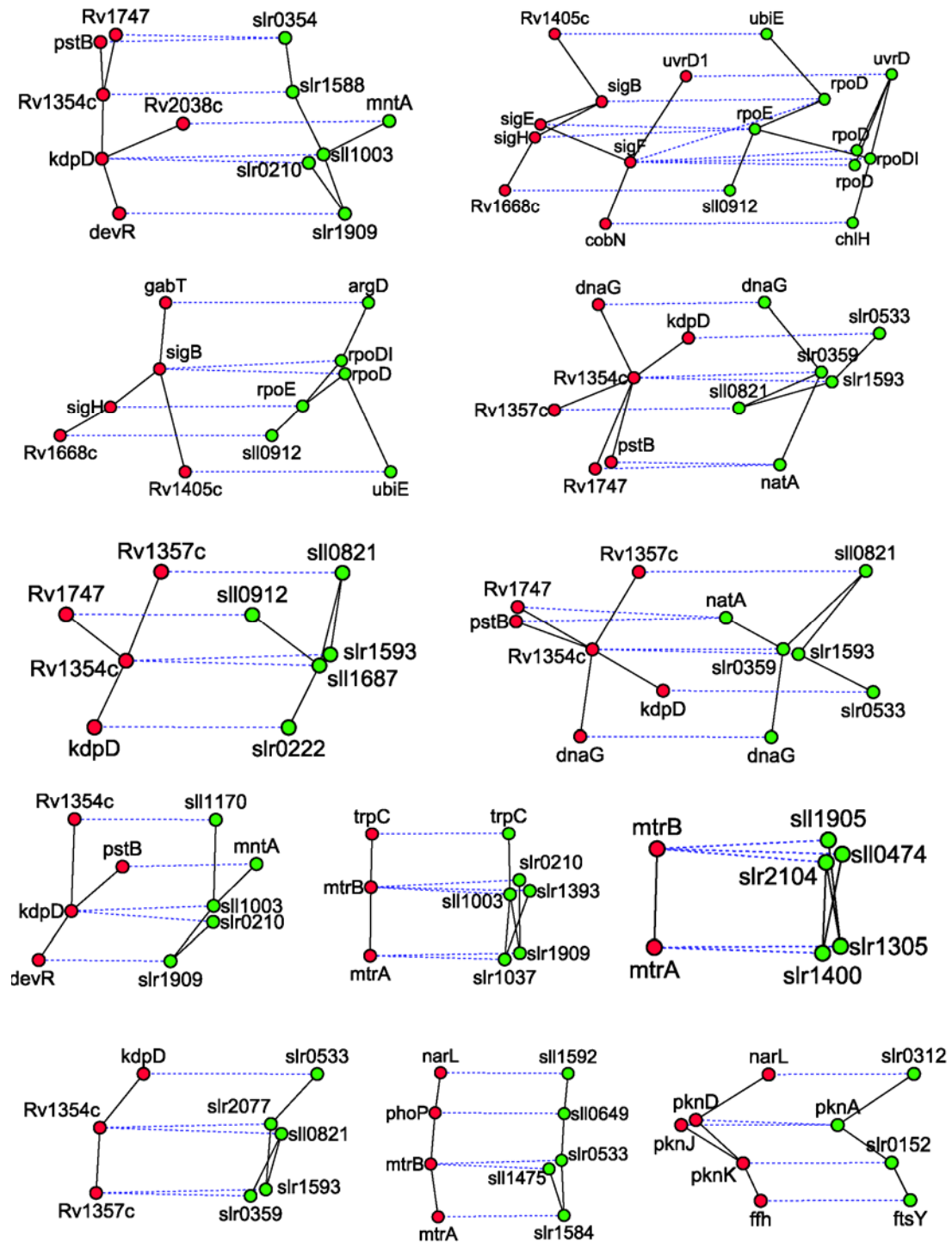


Fig. S6 Conserved subnetworks between *M. tuberculosis* and *Synechocystis* sp.



12 conserved subnetworks were identified between *M. tuberculosis* and *Synechocystis* sp using NetworkBlast. Red nodes represent the proteins from *M. tuberculosis* and green from *Synechocystis* sp. Blue dash line indicate cross-species sequence similar between proteins.