Network Topology Analysis and Hub Target Identification Panel A: Degree Distribution Panel B: Centrality Analysis EGFR Mean: 9.3 100 -Betweenness Centrality 3. PTGS2 Count TOP1 CYP3A4 CYP2C9 TRPA1 PTPN1 ALDH1A1 HIF1A CYP1A2 15 20 10 20 25 5 10 15 Degree Centrality Degree **Panel C: Centrality Correlation Panel D: Hub Protein Connectivity EGFR** 24 0.96 0.49 -0.34eigenvector PTGS2 22 -0.31 -0.15 closeness -0.34CYP2C9 **Hub Proteins** CYP3A4 18 0.69 -0.15 0.49 betweenness ALDH1A1 16 -0.31 0.96 degree 0.69 HIF1A CYP1A2 20 Degree Centrality