

Resources for Translational Bioinformaticians

Jake Chen

In this project researchers developed software that can help extract results from PubMed literature to a comprehensive connectivity map, developing information on the relationships among drugs, proteins, and diseases. The relationships mined from literature can be thoroughly curated with the tool's web-based online annotation graphical user interface. These comprehensive connectivity maps cover disease-specific information and will become a valuable resource for translational bioinformaticians.