



Supplemental Figure 1. The Xenbase and Echinobase software stacks. The majority of code is shared between the two resources and features developed for one site benefit the other (light green). When features are resource specific, conditional code is utilized as indicated by dark green. Database content is mostly unique to each resource, except for shared content such as core ontologies (e.g. GO), disease relationships, human protein-protein interaction data etc. Double ended arrows indicate database queries.