Figure 2

(a) Sensitivity (TP/P) vs. L_{cut} for different prediction methods:
- PIP (de novo prediction)
- Essentiality
- Expression correlation
- MIPS function
- GO biological processes

(b) Comparison of various protein interaction databases (PIE, Gavin, Ho, Uetz, Ito) against different L_{cut} values.

(c) Sensitivity curve showing the performance of PIP and PIE methods, along with Voting results.