(A)

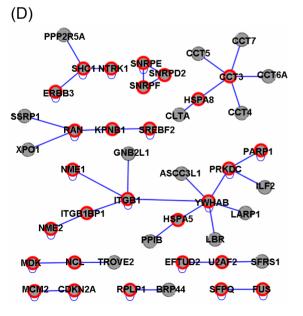
| ` ' | |
|-------------------------|------------|
| GO (Molecular function) | Percentage |
| Protein Binding | 24% |
| Ion Binding | 14% |
| Nucleic Acid Binding | 13% |
| Hydrolase Activity | 10% |
| Nucleotide Binding | 10% |
| | |

(B)

| GO (Biological process) | Percentage |
|-------------------------------------|------------|
| Cellular Physiological Process | 27% |
| Metabolism | 21% |
| Regulation of Cellular Process | 9% |
| Regulation of Physiological Process | 9% |
| Cell Communication | 8% |

(C)

| | GIS | Lee_NIH | Kim_NIH | SMD1648 | SAGE | EHCO conserved |
|---|-------|---------|---------|---------|-------|----------------|
| | up | up | up | up | up | up |
| query | 161 | 161 | 46 | 463 | 112 | 139 |
| no PPI information | 22 | 12 | 14 | 102 | 44 | 19 |
| # of query forms PPI | 45 | 52 | 4 | 128 | 7 | 47 |
| query also hub | 28 | 29 | 2 | 81 | 2 | 18 |
| query query PPIs | 55 | 62 | 4 | 180 | 5 | 48 |
| ratio of query forms PPI out of total query | 32.4% | 34.9% | 12.5% | 35.5% | 10.3% | 39.2% |



(E)

