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About the Cover
Reactive stroma from prostate cancer and normal prostatic stroma was laser captured, RNAs were amplified and gene expression analyzed using expression microarrays. A total of 1141 genes were differentially expressed in reactive compared with normal stroma. The Human Protein Reference Database catalogue of over 35,000 known physical protein-protein interactions was queried to determine how the reactive stroma-associated genes relate to each other in terms of interactions between their associated proteins. A total of 292 interactions were visualized as a graphical network. A view of the central node is shown. For further details, please see Dakhova and colleagues on page 3979 in this issue.