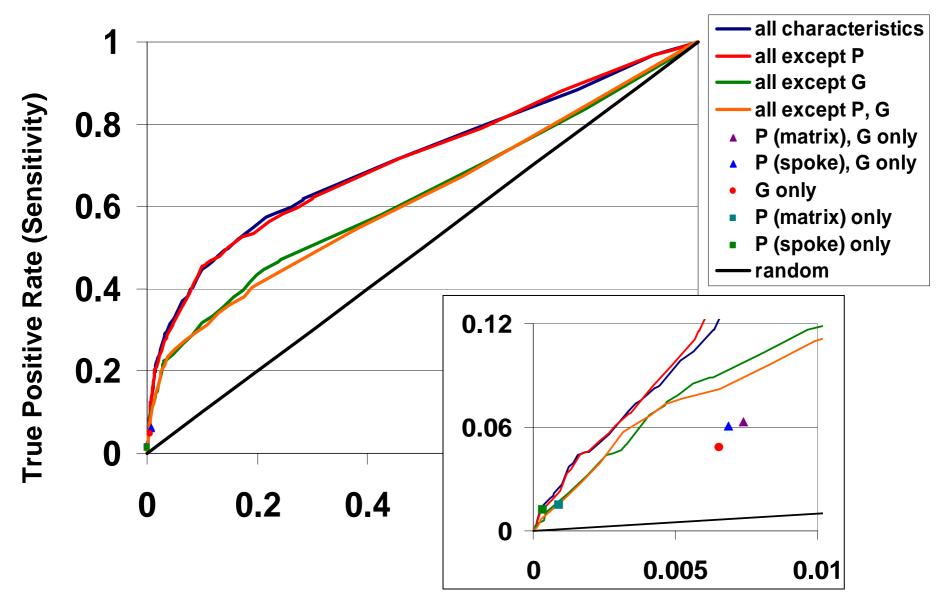
Supplementary Figure 1a



False Positive Rate (1 – Specificity)

Supplementary Figure 1b

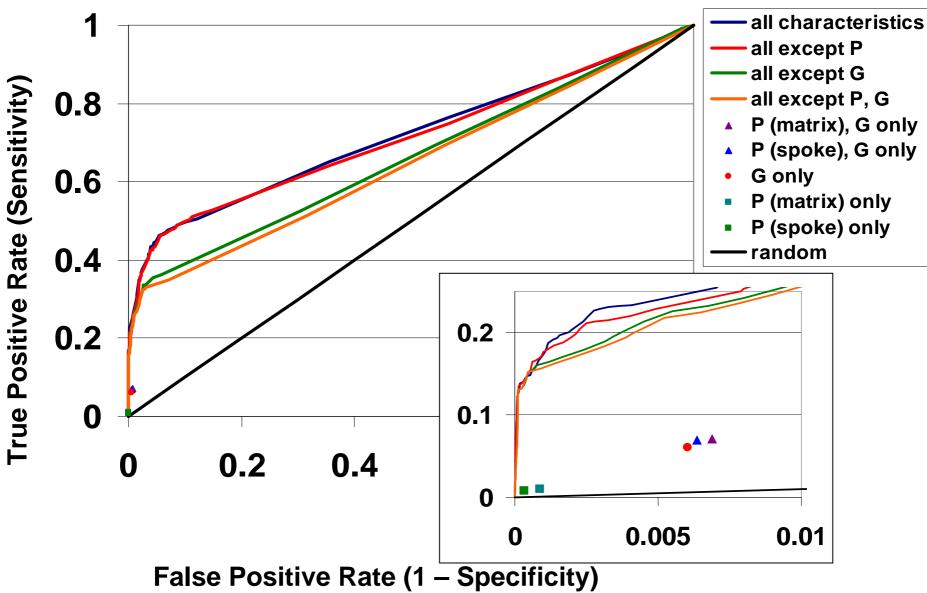
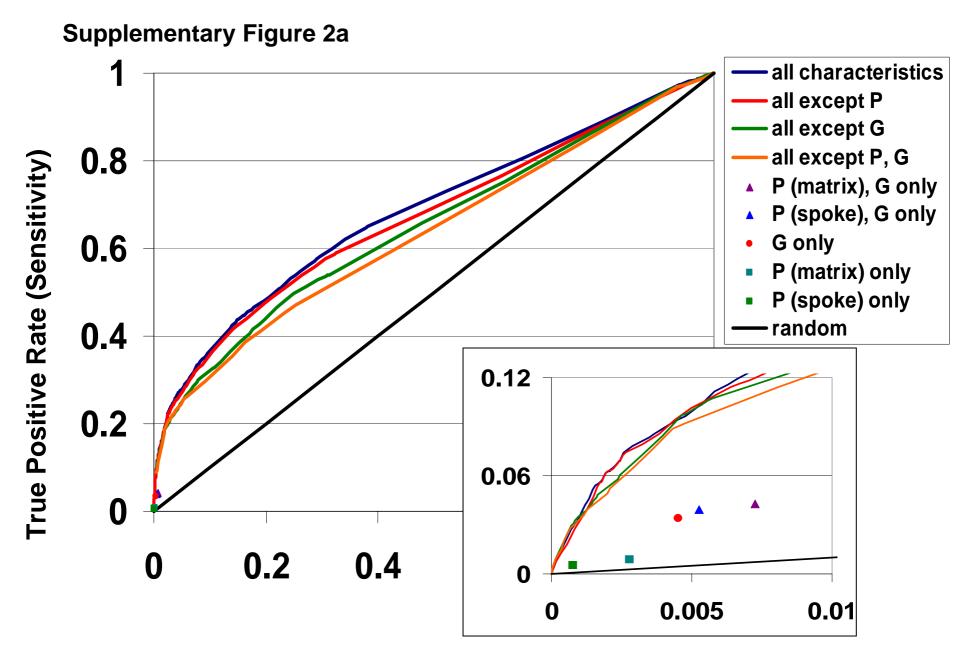
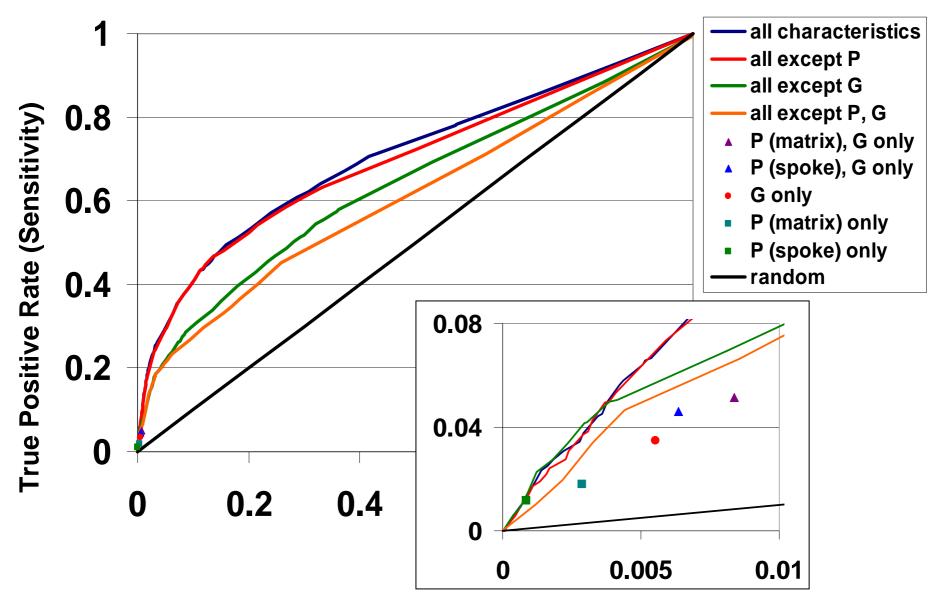


Figure 1. Performance when predicting 'shared component' (a) and 'shared function' (b) with and without genetic (G) [20] and/or Gavin et al. physical (P) [11] interaction information.



False Positive Rate (1 – Specificity)

Supplementary Figure 2b



False Positive Rate (1 – Specificity)

Supplementary Figure 2c

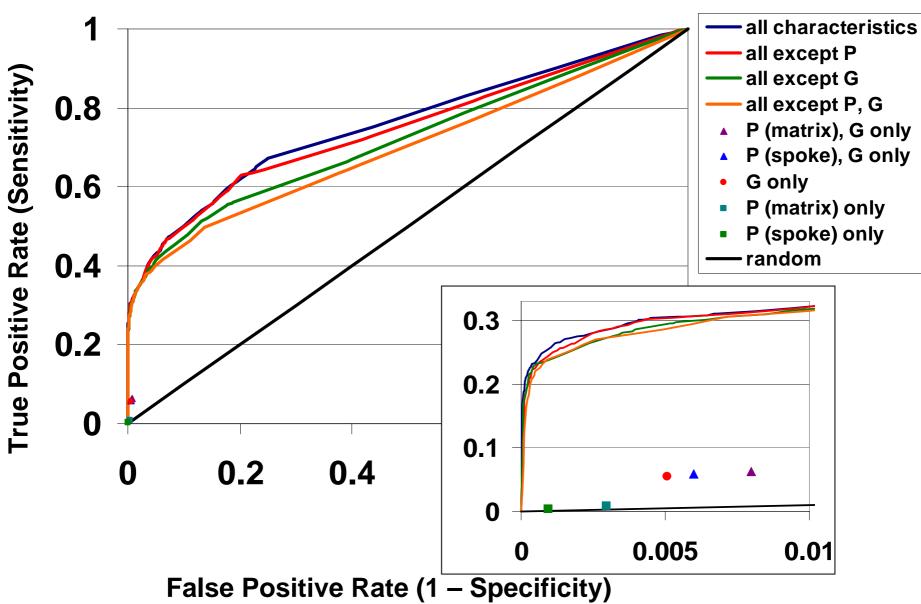
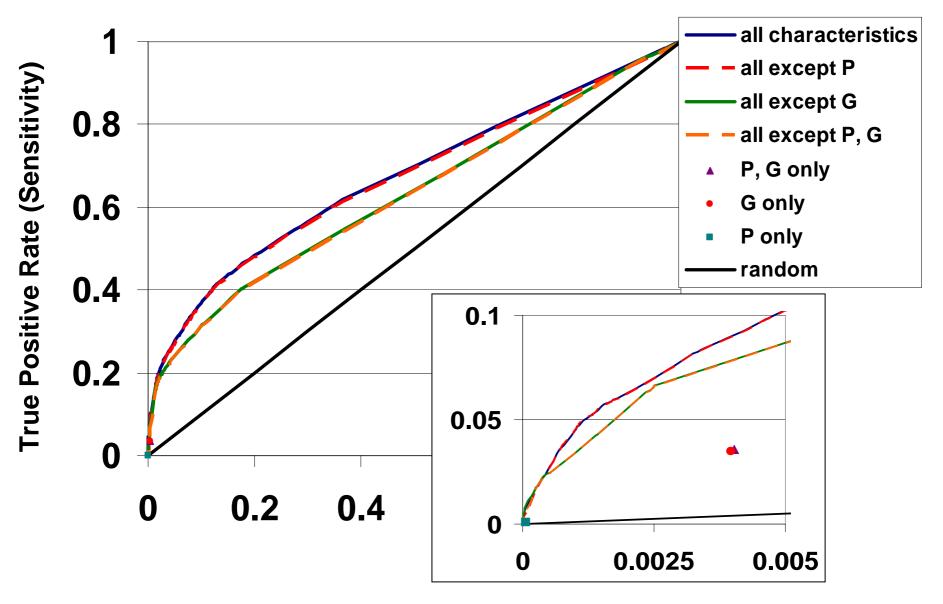


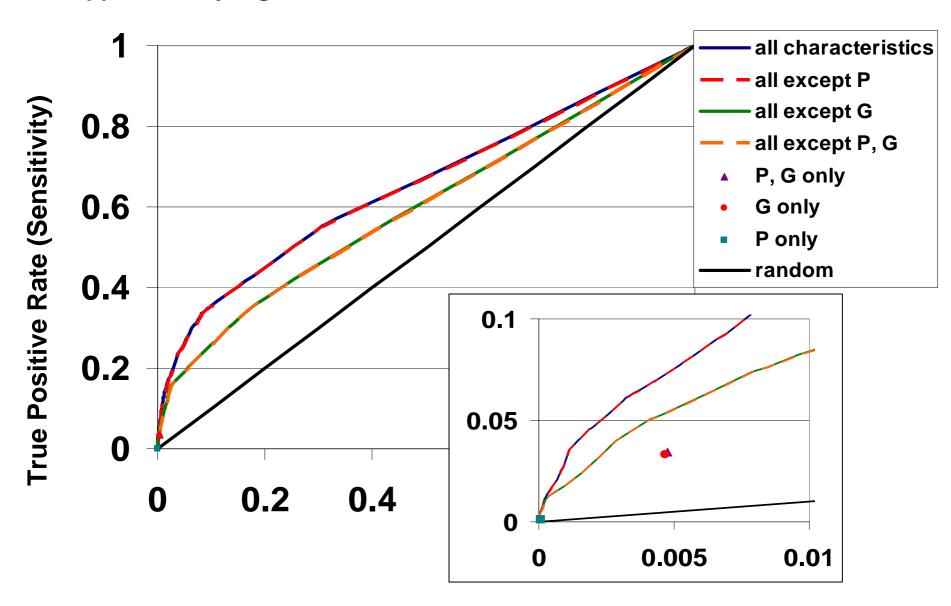
Figure 2. Performance when predicting 'shared process' (a), 'shared component' (b), and 'shared function' (c), with and without genetic (G) [20] and Ho et al. physical (P) [9] interaction information.

Supplementary Figure 3a



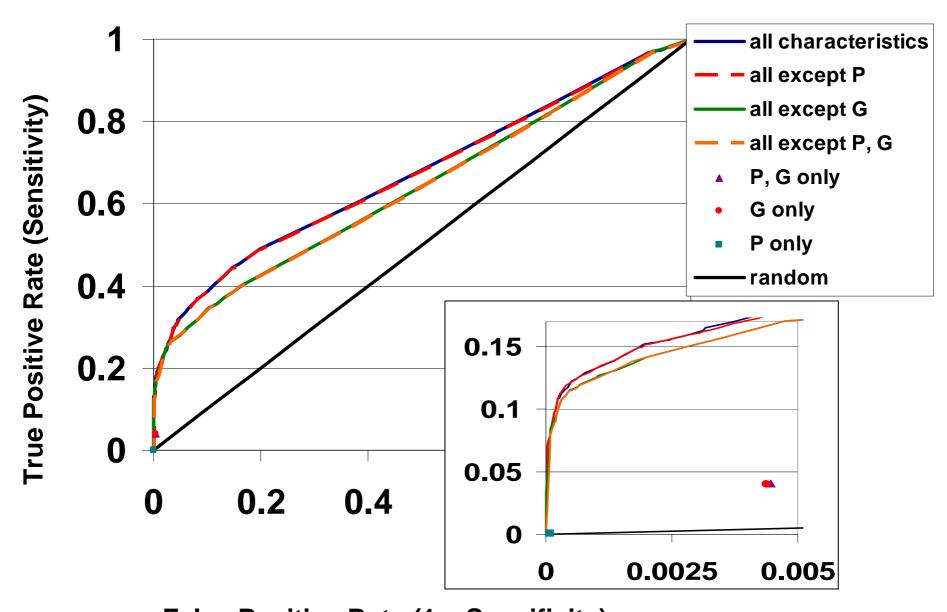
False Positive Rate (1 – Specificity)

Supplementary Figure 3b



False Positive Rate (1 – Specificity)

Supplementary Figure 3c



False Positive Rate (1 – Specificity)

Figure 3. Performance when predicting 'shared process' (a), 'shared component' (b), and 'shared function' (c), with and without genetic (G) [20] and/or yeast two hybrid physical (P) [12,13] interaction information.