## SUPPLEMENTAL FIGURE AND TABLE INVENTORY

## Supplemental Figures

- Figure S1. Number of Kinases Bound Plotted Against the Number of Targets Bound (Included on Page 1 of Supplemental figures and legends)
- Figure S2. Protein array vs Breitkreutz AC/MS complex comparison (Included on Page 2 of Supplemental figures and legends)
- 3. **Figure S3. Rio1 Modules** (Included on Page 4 of Supplemental figures and legends)
- 4. Figure S4. GO analysis enrichment diagram for CAMK Targets

  (Included on Page 5 of Supplemental figures and legends)

## **Supplemental Tables**

- Table S1. Protein Array Kinase interactions (File uploaded as PDF Fasolo\_Table S1)
- 2. Table S2: Targets that failed to vaidate by co-IP/Immunoblot (File uploaded as PDF Fasolo\_Table S2)
- 3. Table S3: Markov Analysis of phophomotif enrichment (File uploaded as PDF Fasolo\_Table S3)
- Table S4. Over-enriched interaction modules (File uploaded as PDF Fasolo\_Table S4)

- Table S5. GO Analysis for all kinase targets (File uploaded as PDF Fasolo\_Table S5)
- 6. **Table S6. GO Analysis for CAMK kinase targets** (File uploaded as PDF Fasolo\_Table S6)
- 7. **Table S7. Yeast Strains used in this study** (File uploaded as PDF Fasolo\_Table S7)