

Figure S5. Isovolume of GAB1 PH domain generated by GRID. The area favoring H-bond acceptor (-6.0 kcal/mol) was displayed in magenta surface, whereas area favoring hydrophobic (-2.0 kcal/mol) probes was displayed in green surface. The protein backbone is illustrated with cyan lines, and the ligand (IP4) is in blue sticks. The crucial residues are labeled around isovolume surface.

