

Title: Quantitative Proteomic Analysis Reveals Novel Insights into Intracellular Silicate Stress-Responsive Mechanisms in the Diatom *Skeletonema dohrnii*

Authors: Thangaraj Satheeswaran, Shang Xiaomei, Sun Jun*, Liu Haijiao

Figure S1. An iPath metabolic biochemistry map of protein expressed and identified as down-regulated in the Si-limited and Si-added cells of *S. dohrnii*



