

Article

Using Geometric Approaches to the Common Metabolomics in Acute Lymphoblastic Leukemia and Rhabdomyosarcoma: Expanding and Integrating Pathway Simulations

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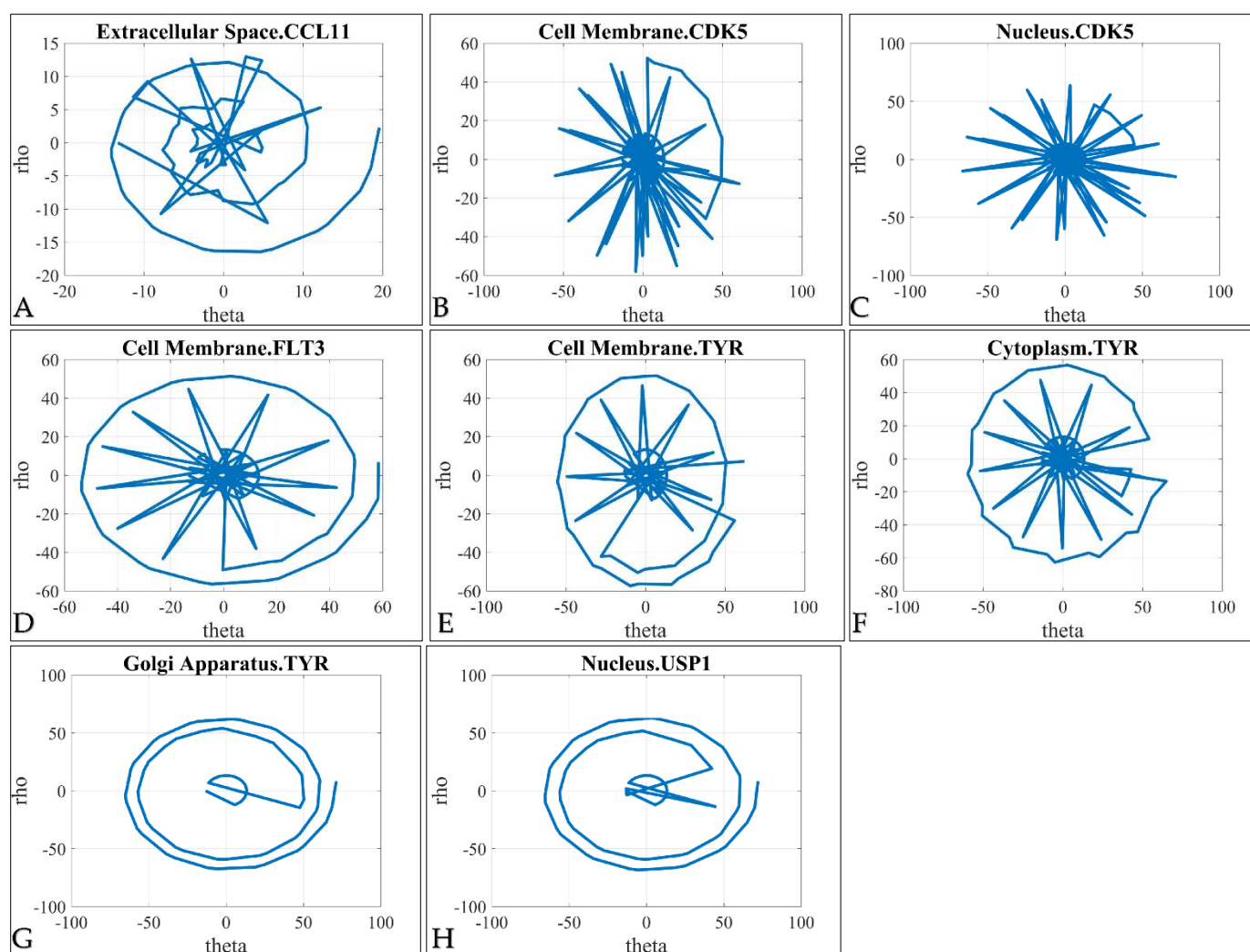


Figure S2. Polar transformation of time-series data. All species, namely extracellular CCL11 (a), membrane CDK5 (b), membrane FLT3 (c), membrane TYR (d), cytoplasmatic TYR (e), nucleolar CDK5 (f), nucleolar USO1 (g) and Golgi TYR (h) manifested quasi-turbulent dynamics.