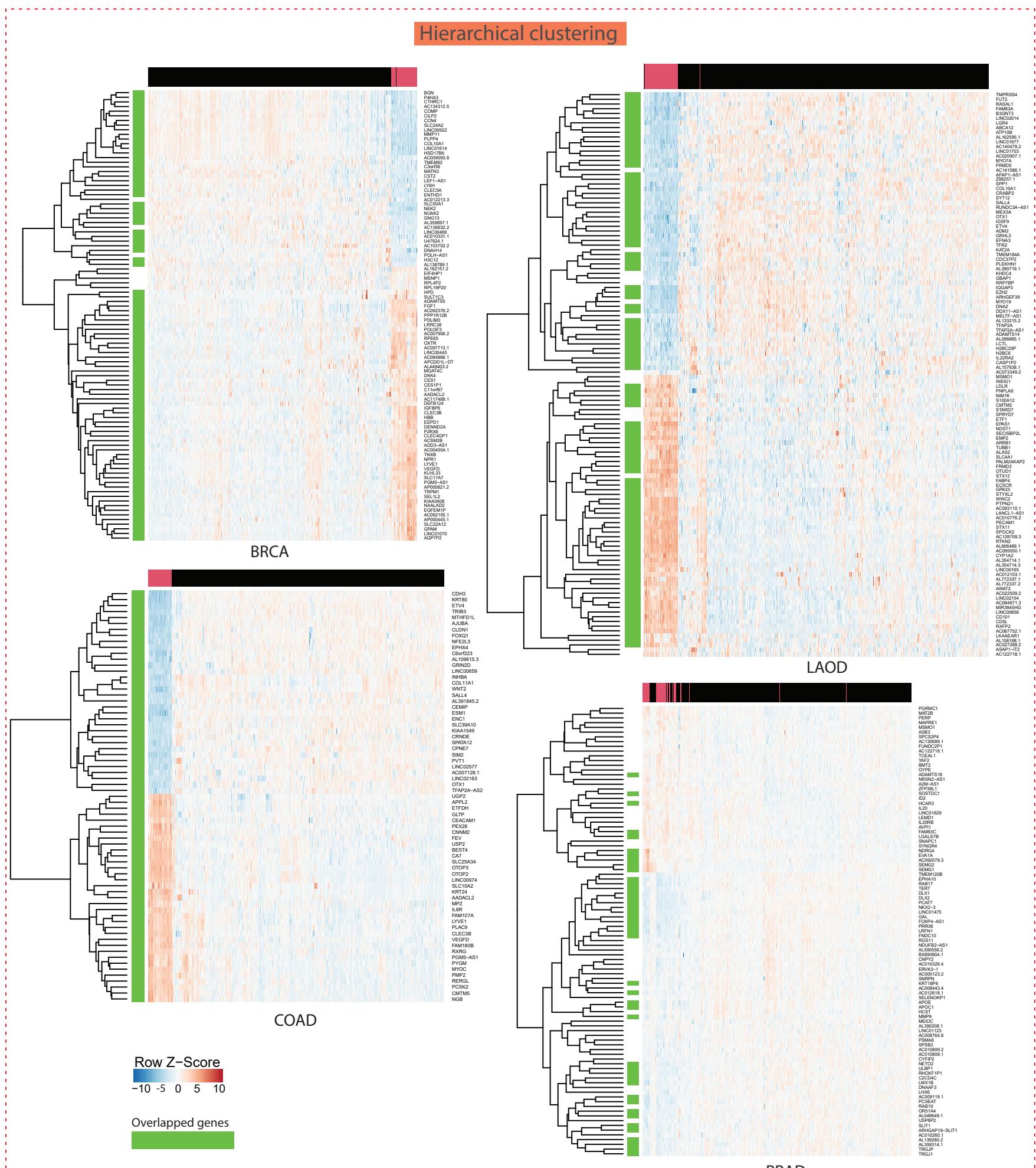


A Systems Biology Approach to Unveil Shared Therapeutic Targets and Pathological Pathways Across Major Human Cancers

Aftab Alam¹, Mohd Faizan Siddiqui², Rifat Hamoudi^{3,4,5}, Uday Kishore^{6,7}, Maria Fernandez Cabezudo^{7,8}
and Basel K. Al-Ramadji^{1,7,9*}



*The genes highlighted in green are the ones that limma had also selected

* The samples highlighted in red are **Solid Tissue Normal**, the samples highlighted in black are **Primary solid Tumor**.