



Fig. S22. The relationship between gene average gravitation (aveG) scores with four features: average Pearson Correlation Coefficient (avePCC), mutation rate, gene cDNA length, and gene connectivity (degree) in 9 cancer types. We separated all genes into three different categories based on their aveG scores: low aveG group (down 2,000 genes having the lowest aveG scores), middle aveG group (middle 2000 genes having the the aveG scores ranked between 4,500 to 6,500), high aveG group (top 2,000 genes have the highest aveG scores). P: probability.