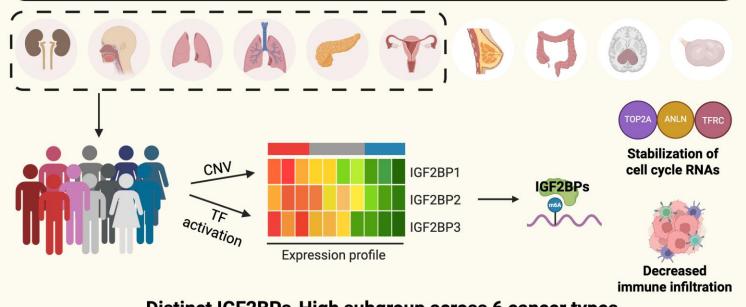
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Pan-cancer multi-omics analysis of m6A associated subgroups



Distinct IGF2BPs-High subgroup across 6 cancer types

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Decreased immune infiltration



IGF2BP: Expression profile

Cover feature: Proteogenomic Analysis on RNA m6A Modification-Associated Genes Identifies a Distinct Subgroup with High IGF2BPs Expression Across Cancer Types Yebin Ryu, Eunhyong Chang, Hayoon Park, Sung-Yup Cho, Joon-Yong An





