

READ - Rectum adenocarcinoma

Subtype (CMS)	Biology & Expression	Genomic Alterations	Clinical Features
CMS1: MSI Immune	<ul style="list-style-type: none"> • Strong immune and inflammatory signatures (high PD-L1, cytotoxic T-cell markers) • CpG island methylator phenotype (CIMP-high) 	<ul style="list-style-type: none"> • BRAF^{V600E} mutations • MLH1 promoter methylation leading to MSI-H • Low SCNA burden 	<ul style="list-style-type: none"> • Often right-sided tumors • Better overall survival but poor after relapse • Candidate for immunotherapy
CMS2: Canonical	<ul style="list-style-type: none"> • Epithelial differentiation with upregulated WNT and MYC targets • High epithelial marker expression 	<ul style="list-style-type: none"> • APC and TP53 mutations • Chromosomal instability (high SCNA) 	<ul style="list-style-type: none"> • Generally left-sided tumors • Best overall prognosis • Sensitive to standard chemo-targeted regimens
CMS3: Metabolic	<ul style="list-style-type: none"> • Metabolic dysregulation (lipid, glucose, glutamine pathways) • Mixed epithelial signatures 	<ul style="list-style-type: none"> • KRAS mutations enriched • Intermediate SCNA burden 	<ul style="list-style-type: none"> • Variable location • Intermediate prognosis • Potential for metabolic-targeted therapies
CMS4: Mesenchymal	<ul style="list-style-type: none"> • Strong TGF-β activation and EMT programs • High stromal and angiogenesis signatures 	<ul style="list-style-type: none"> • Frequent TP53 and SMAD4 alterations • High SCNA and stromal infiltration 	<ul style="list-style-type: none"> • Worst relapse-free and overall survival • Resistant to standard therapies • May benefit from stroma-targeted approaches