

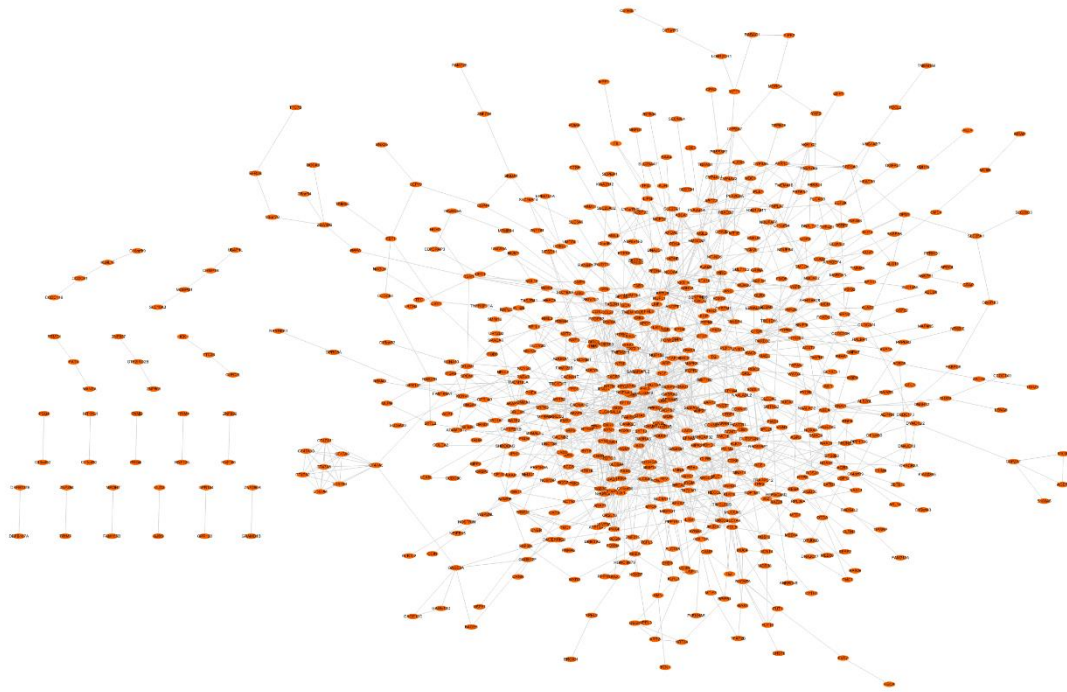
**Identification of hub genes and potential molecular mechanisms related to
radiotherapy sensitivity in rectal cancer based on multiple datasets**

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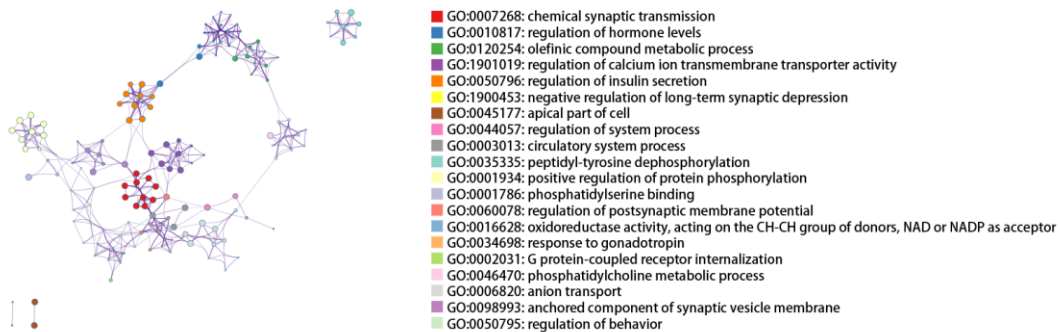
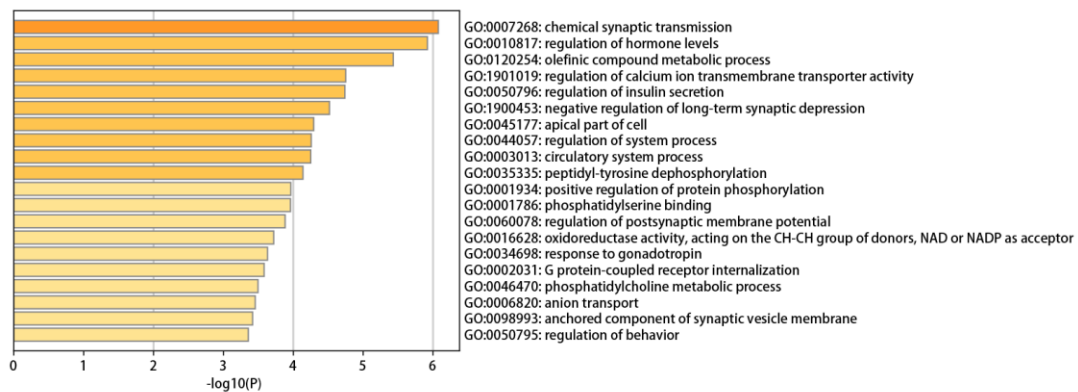
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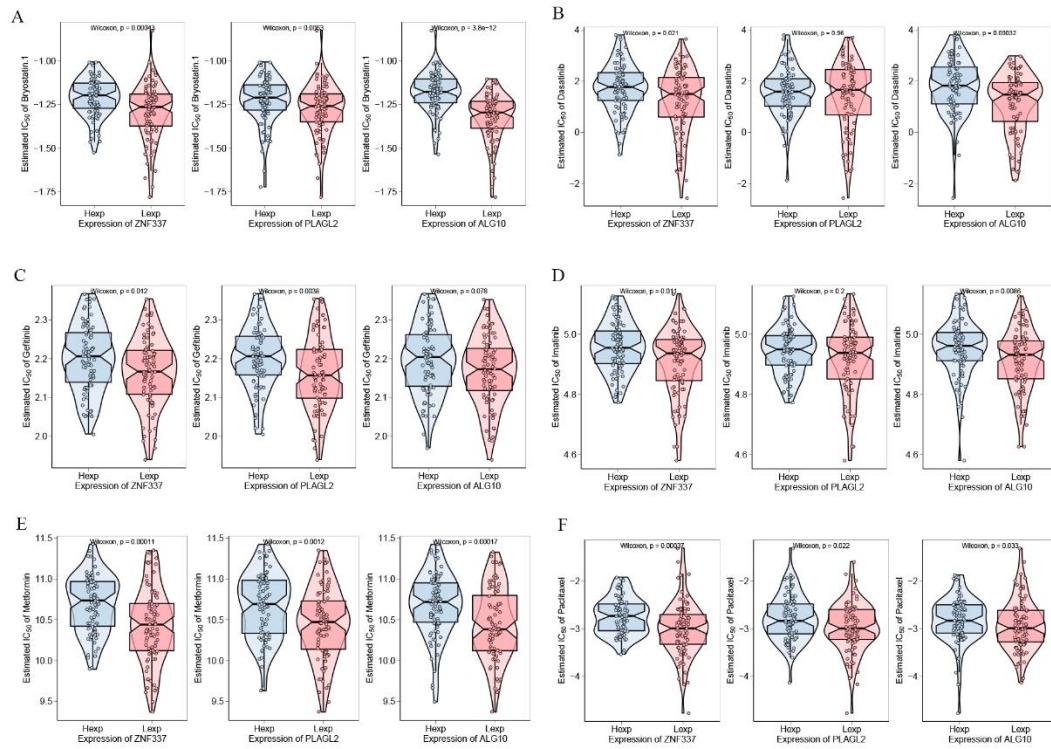
Supplementary Figures and Legends



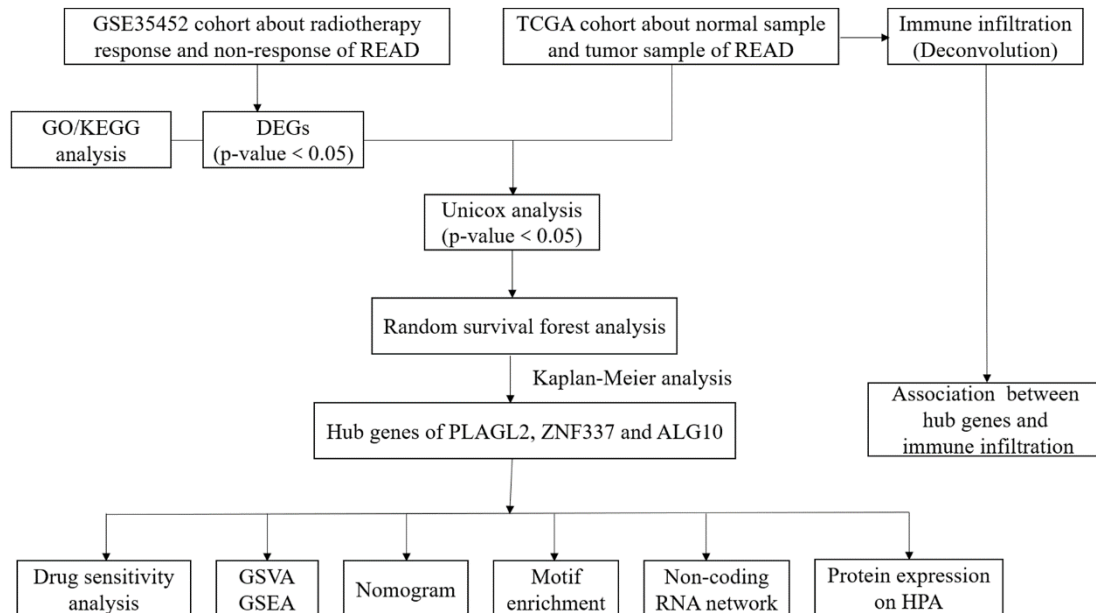
Supplementary Figure 1. Functional enrichment of DEGs by metascape enrichment analysis.



Supplementary Figure 2. Protein interaction network analysis of DEGs by Cytoscape



Supplementary Figure 4. The sensitivity analysis of chemotherapeutic drugs between different expression levels of key genes based on GDSC database. A: Bryostatin.1; B: Dasatinib; C: Gefitinib; D: Imatinib; E: Metformin; F: Paclitaxel.



Supplementary Figure 5. The flowchart of the current study.

Supplemental Table

Supplementary Table 1. Information of the DEGs identified from database. A total of 1119 DEGs were identified from GSE35452 dataset, including 544 up-regulated genes and 575 down-regulated genes in READ cohort.

DEGs	Gene name
Upregulated (544)	LOC100131756, LRRIQ3, LOC101927151, BRCA2, RP11-495P10.9, RP3-522P13.1, UMODL1, C10orf62, MED14OS, LHX8, LOC101928401, SPACA3, DRD3, ENKUR, LOC101928288, CSMD2-AS1, RECQL, TMC1, TRA@, MTNR1B, SEMA3A, GABRA6, SPATC1, SNX25, STX16, RP11-171N4.1, UNC93A, LIG4, LOC284865, TRHR, MTCPI, B3GALTL, LOC100506538, LOC101926912, RP11-356B19.11, SLC6A4, LOC149373, C5orf64, PTPN11, LOC284939, LINC00424, C18orf15, RP11-585P4.5, LOC283075, USP12, GTF2IRD2B, NAA11, LINC01206, SYN1, C7orf25, DIAPH3, SCPEP1, RNF166, FAM230B, UQCC2, RP11-66N11.8, PALM, LINC00116, VAPB, GPR132, TRIM65, SLC26A4-AS1, KIR2DL2, FOXK1, LOC101929025, TPP2, LOC284798, CXCR3, OR51A10P, CMC4, RP11-1012E15.1, MANEA, RNF6, ANGPT1, KCNV1, CYP2A7, PFDN4, LINC00593, MTIF3, LOC729461, SPATC1L, MRGPRX4, LINC01348, PIPOX, CCND2, ZFP1, KLHL18, LINC00052, LINC01424, LINC00703, LOC101927988, MMP27, LOC101928545, YAF2, NRTN, RBM41, POLB, DOCK9, EDARADD, LOC101929289, CTB-78F1.1, LOC100271832, GPR107, GCNT4, GABRB1, PLB1, LINC01420, LOC101929709, HM13, LRRC37BP1, LOC101927650, TUSC8, AC064852.4, PPIAL4B, PPIAL4C, PPIAL4G, C14orf39, SNORD3A, SNORD3B-1, SNORD3B-2, SNORD3C, SNORD3D, LINC00323, DTD1, GRINA, FAM200A, UBD, LINC00461, MIR9-2, NAALAD2, SLC10A3, GS1-164F24.1, CXCL11, DEFB107A, DEFB107B, C10orf53, RP11-498C9.17, RFPL2, ADCY5, PHF2P1, LOC101929248, ZNF92, MED14, POU5F1, BC037833, RP11-492E3.2, LOC100652824, MIR3934, TCP11L1, FAM72A, FAM72B, FAM72C, FAM72D, SKIL, OR11A1, DDAH1, TMPRSS11B, EFR3B, PDGFRA, LOC101928435, LOC101927003, BC031255, RP11-260O18.1, NKAP, LOC101928446, ZNF280B, CYP4A11, CEP44, PARVB, RP11-360K13.1, CIDCEP, AGER, MED30, BMP15, TGIF2LY, CXXC1P1, ABCB8, LOC285766, HRG, GPR143, MORC4, TCFL5, CASC2, POM121, CCDC116, RP11-402G3.5, TTC7B, LOC101927308, PTPRO, CEP19, AK054708, RP11-547D24.1, SGOL1, LOC646482, CTA-331P3.1, OGFRL1, LOC653786,

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