Hierarchical clustering of features

Samples

rig \ samples class

3_utr_parsedFeats.csv.gz_5_UTR_start_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_GC_freq_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_utr_seq_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_C_freq_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_end_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_end_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_mode_ratio_freq_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_T_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_start_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_A_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_C_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_A_freq_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_T_freq_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_CDS_Length_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_G_freq_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_Gene_start_bp_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_3_len_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_start_3_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_end_3_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3utr_counts_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_end_2_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_start_2_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_2_len_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_end_3_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_start_3_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_3_len_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_end_1_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_1_len_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_start_1_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_tot_5utr_len_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_start_1_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_end_1_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_UTR_1_len_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_3_utr_len_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_G_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_tot_3utr_len_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_Transcript_length_including_UTRs_and_CDS_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5utr_counts_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_start_2_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_end_2_by_Gene_name, Gene names (primary)
3_utr_parsedFeats.csv.gz_5_UTR_2_len_by_Gene_name, Gene names (primary)
Features
Involvement in disease
Keywords
Gene ontology (GO)
Post−translational modification
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Post−translational modification
Tissue specificity
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Protein families
Protein names
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Tissue specificity
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Keywords
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Features
Compositional bias
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Compositional bias
Mass, Length
Length, Mass
Mass
Length
Sequence
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Protein names
DNA binding
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Involvement in disease
Non−terminal residue
Developmental stage
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Developmental stage
Repeat
Zinc finger
Pathway
Nucleotide binding
human_gene_mirtar_targets.csv.gz_text_index_of_Protein families_with_attributes, Gene ontology (GO)
Protein families