## GXB\_clinical\_vars\_step2

Load GXB\_Clinical\_Vars.csv

Find unique values in column "Suggested.GXB.Clinical.Variable.Name"

GXBvars <- variabletable\$Suggested.GXB.Clinical.Variable.Name
unique(GXBvars)</pre>

```
[1] sample_barcode
##
## [2] array_sample_id
## [3] sample_title
## [4] survival_time
## [5] treatment_adjuvant
## [6] ?endocrine status
## [7] pharmaceutical treatment
## [8] disease_free_survival_event
## [9] disease_free_survival
## [10] distant met free survival event
## [11] distant_met_free_survival
## [12] vital_status
## [13] ER status
## [14] HER2_status
## [15] histologic_grade
## [16] histologic diagnosis
## [17] lymph_node_status
## [18] age_at_initial_pathologic_diagnosis
## [19] PR_status
## [20] tumor size
## [21] type_treatment
## [22] IMS
## [23] DMFS 10Y TIME
## [24] DMFS_10Y_EVENT
## [25] IBS
## [26] IDS
## [27] ICR
## [28] bone_metastasis_event
## [29] cohort_specific_ID
## [30] series
## [31] lung_metastasis_event
## [32] ?lung_met_first_event
## [33] ?lung met free survival
## [34] metastasis_event
## [35] disease_free_survival_event; disease_free_survival
## [36] <NA>
## [37] disease free survival event; distant met free survival event
## [38] disease free survival; distant met free survival
## [39] sample_storage
## [40] breast_cancer_cause_of_death
## [41] lymph_node_examined_count
## [42] RFS_5Y_EVENT
## [43] bone metastasis free survival
## [44] lung_metastasis_free_survival
## [45] number_of_lymphnodes_positive
## [46] tissue type
## [47] van_t_veer_signature
## [48] GGI_indicator
## [49] OS_10Y_TIME
## [50] angio_invasion_indicator
## [51] ?NPI
## [52] risk_AOL_indicator
## [53] risk NPI indicator
```

```
## [54] risk_SG
## [55] risk_veridex_indicator
## [56] genetic_grade_signature_status_sws_classifier
## [57] probability1_by_sws_classifier
## [58] probability3_by_sws_classifier
## [59] p53_mutation_status
## 58 Levels: ?endocrine_status ... vital_status
```