

Combining Plasma Extracellular Vesicle Let-7b-5p, miR-184 and Circulating miR-22-3p Levels for NSCLC Diagnosis and Drug Resistance Prediction

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Supplementary information 1. Bioinformatics pipeline for survival analysis

```
#code adapted from: #https://www.jianshu.com/p/5e4285289137 #https://bioc.ism.ac.jp/packages/3.2/bioc/vignettes/TCGAbiolinks/inst/doc/tcgaBiolinks.html #https://www.rdocumentation.org/packages/survminer/versions/0.1.1/topics/ggsurvplot

#hsa-mir-184

samplesTP <- TCGAquery_SampleTypes(barcode = colnames(miR_matrix), typesample = c("TP"))
mir184_exp <- miR_matrix[c("hsa-mir-184"), samplesTP]
names(mir184_exp) <- sapply(strsplit(names(mir184_exp), '-'), function(x) paste(x[1:3], collapse="-"))
clinical$GENE <- mir184_exp[clinical$submitter_id]

df1<-subset(clinical,select=c(submitter_id,vital_status,days_to_death,days_to_last_follow_up,GENE))
df1$years_death<- df1$days_to_death/365#convert days to years
df1$years_to_last_follow_up <- df1$days_to_last_follow_up/365
df1$os<-ifelse(df1$vital_status=='Alive',df1$years_to_last_follow_up,df1$years_death)#alive sample uses
df1 <- df1[!is.na(df1$GENE),]#Remove samples with 0 expression
df1[df1$vital_status=='Dead',]$vital_status <- 2
df1[df1$vital_status=='Alive',]$vital_status <- 1
df1$vital_status <- as.numeric(df1$vital_status)

#Determine the optimal cutpoint of variables
df1.cut <- surv_cutpoint(df1,
  time = "os",
  event = "vital_status",
  variables = c("GENE"))

summary(df1.cut)

##      cutpoint statistic
## GENE 13.91485  2.567961

#Categorize variables
df1.cat<-surv_categorize(df1.cut,variables = NULL, labels = c("low", "high"))
head(df1.cat)

##            os vital_status GENE
## 1 2.5945205          1   low
## 2 2.9616438          2   low
## 3 0.7342466          2   low
## 4 0.8410959          2   low
## 5 0.0000000          2   low
## 6 1.4931507          1   low
```

```
df1.cat
```

```
##          os vital_status GENE
## 1    2.59452055      1  low
## 2    2.96164384      2  low
## 3    0.73424658      2  low
## 4    0.84109589      2  low
## 5    0.00000000      2  low
## 6    1.49315068      1  low
## 7    1.47123288      1  low
## 8    0.06027397      2  low
## 9    8.14520548      1  low
## 10   NA               2  low
## 11   0.63013699      1  low
## 12   3.41369863      1  low
## 13   5.77808219      1  low
## 14   1.40547945      1  low
## 15   2.56986301      1  low
## 16   2.00000000      1  low
## 17   5.10684932      1 high
## 18   0.38082192      1  low
## 19   0.78630137      1  low
## 20   4.19452055      2  low
## 21   1.66301370      2 high
## 22   1.63835616      2  low
## 23   2.21369863      2  low
## 24   0.61643836      1  low
## 25   3.14520548      1  low
## 26   5.92054795      1  low
## 27   0.76986301      2  low
## 28   4.44109589      1  low
## 29   1.62465753      2  low
## 30   2.16712329      1  low
## 31   1.24657534      1  low
## 32   1.79726027      2  low
## 33   7.33150685      1  low
## 34   0.96712329      1  low
## 35   0.38630137      1  low
## 36   1.96986301      2  low
## 37   4.79452055      1  low
## 38   1.66301370      1  low
## 39   3.00547945      1  low
## 40   1.33424658      1  low
## 41   0.36712329      1  low
## 42   2.43287671      1  low
## 43   0.01095890      2  low
## 44   7.09589041      1  low
## 45   4.92602740      2  low
## 46   0.13150685      1  low
## 47   1.19178082      1 high
## 48   2.38082192      2  low
## 49   4.87123288      2  low
## 50   3.95068493      1  low
## 51   3.39452055      1  low
```

## 52	1.80000000	1 low
## 53	10.79452055	1 low
## 54	2.45479452	1 low
## 55	5.18630137	1 low
## 56	1.12054795	1 low
## 57	0.70410959	2 low
## 58	1.22739726	1 low
## 59	0.96164384	1 low
## 60	2.08219178	2 low
## 61	0.84383562	2 low
## 62	0.16986301	1 low
## 63	7.10958904	1 low
## 64	4.11232877	2 low
## 65	2.37260274	1 low
## 66	1.19178082	1 low
## 67	6.46575342	1 low
## 68	1.69041096	1 low
## 69	2.03013699	1 low
## 70	1.89041096	1 low
## 71	2.43561644	1 low
## 72	1.61917808	1 low
## 73	1.22739726	1 low
## 74	4.03835616	1 low
## 75	1.78356164	1 low
## 77	0.04109589	1 low
## 78	0.96986301	2 low
## 79	3.16986301	1 low
## 80	2.54794521	1 low
## 81	1.28219178	2 low
## 82	3.10958904	2 low
## 83	0.09041096	2 low
## 84	1.30958904	2 low
## 85	3.98356164	2 low
## 86	1.99452055	1 low
## 87	1.13424658	2 low
## 88	0.57534247	2 low
## 89	0.15890411	2 low
## 90	2.33424658	1 high
## 91	2.72054795	1 low
## 92	1.17534247	2 low
## 93	2.70684932	1 low
## 94	1.78356164	1 low
## 95	5.85479452	1 low
## 96	3.27945205	2 low
## 97	1.30684932	1 low
## 98	0.50410959	1 low
## 99	0.92054795	2 low
## 100	1.25205479	1 low
## 101	8.38082192	1 low
## 102	1.85479452	2 low
## 103	1.13424658	1 low
## 104	1.58356164	1 high
## 105	2.67397260	2 low
## 106	2.54520548	2 low

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## 107 0.44109589          2 low
## 108 1.28493151          2 low
## 109 1.03013699          2 low
## 110 2.02739726          1 low
## 111 1.83561644          1 low
## 112 2.50136986          1 high
## 113 0.45205479          1 low
## 114 1.37808219          2 low
## 115 1.56986301          1 high
## 116 0.51232877          2 low
## 117 2.72602740          2 low
## 118 1.58356164          1 low
## 119 2.73150685          1 low
## 120 2.26301370          2 low
## 121 1.47671233          1 high
## 122 1.13698630          1 high
## 123 2.36712329          2 low
## 124 1.98082192          1 low
## 125 0.92876712          2 low
## 126 2.20547945          1 high
## 127 0.49041096          2 low
## 128 1.49863014          1 low
## 129 1.16712329          1 low
## 130 4.15342466          2 high
## 131 0.23835616          2 low
## 132 1.01369863          2 low
## 133 0.31780822          2 low
## 134 3.91506849          1 low
## 135 2.95616438          1 high
## 136 2.93424658          1 low
## 137 1.25205479          2 low
## 138 1.65753425          1 low
## 139 1.90136986          2 low
## 140 1.55616438          1 low
## 141 2.26575342          1 low
## 142 0.05205479          2 low
## 143 0.00000000          1 low
## 144 7.75890411          1 low
## 145 2.34246575          2 low
## 146 2.72602740          2 low
## 147 1.55616438          1 low
## 148 0.50958904          1 low
## 149 2.36438356          1 low
## 150 3.18630137          1 low
## 151 1.78904110          1 low
## 152 1.92054795          1 high
## 153 3.57534247          1 low
## 154 3.54246575          2 low
## 155 NA                  1 low
## 156 2.47671233          1 low
## 157 3.08493151          1 low
## 158 0.49041096          1 low
## 159 0.66849315          2 low
## 160 1.96712329          1 low

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## 161 1.51232877    1 low
## 162 1.78630137    1 low
## 163 0.06575342    1 low
## 164 8.68219178    2 high
## 165 0.32602740    1 low
## 166 19.34794521   1 low
## 167 5.29315068    1 high
## 168 1.67123288    1 high
## 169 4.65753425    1 low
## 170 6.70958904    1 low
## 171 1.64383562    1 low
## 172 3.38356164    2 low
## 173 3.77808219    2 low
## 174 0.70136986    1 low
## 175 1.16712329    1 low
## 176 4.43013699    1 low
## 177 1.20547945    1 low
## 178 2.01917808    2 low
## 179 2.25753425    1 low
## 180 1.87123288    1 low
## 181 1.63287671    1 low
## 182 0.04931507    2 low
## 183 1.26027397    2 low
## 184 7.16986301    2 low
## 185 0.83013699    2 low
## 186 3.37808219    1 low
## 188 0.09863014    1 low
## 189 2.93698630    1 low
## 190 1.33698630    2 low
## 191 2.60273973    2 low
## 192 3.08219178    1 low
## 193 1.71506849    1 high
## 194 1.92876712    1 high
## 195 2.08493151    1 low
## 196 1.63013699    1 low
## 197 5.40821918    1 low
## 198 2.40821918    2 low
## 199 0.70684932    2 low
## 200 0.68493151    2 low
## 201 4.38356164    2 low
## 202 1.67123288    1 low
## 203 1.01917808    1 low
## 204 1.65205479    1 high
## 205 1.22191781    1 low
## 206 0.21643836    1 low
## 207 1.41095890    1 low
## 208 9.95890411    1 low
## 210 1.82465753    2 low
## 211 0.03013699    1 low
## 212 1.96986301    1 low
## 213 3.25753425    1 low
## 214 0.51780822    2 low
## 215 0.13698630    1 low
## 216 0.45753425    2 low

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## 217 2.31506849      1 low
## 218 7.73424658      1 low
## 220 2.58630137      1 low
## 221 5.65753425      1 low
## 222 4.10410959      2 low
## 223 0.41369863      1 low
## 224 1.16986301      1 low
## 225 2.70410959      2 low
## 226 1.47945205      1 low
## 227 0.42191781      2 low
## 228 1.80273973      1 low
## 229 0.33972603      2 low
## 230 0.84109589      1 low
## 231 0.66849315      2 high
## 232 0.87945205      2 low
## 233 2.60821918      2 low
## 234 1.55616438      1 low
## 235 2.29863014      1 high
## 236 3.75068493      1 low
## 237 7.34520548      2 low
## 238 1.15616438      1 high
## 239 1.36986301      2 low
## 240 6.81643836      1 low
## 241 18.44383562     1 low
## 242 4.44383562      2 low
## 243 0.26575342      2 low
## 244 1.44109589      1 low
## 245 3.22739726      1 low
## 246 NA                1 low
## 248 3.31232877      2 low
## 249 3.46575342      2 high
## 250 1.30958904      1 low
## 251 6.48767123      1 low
## 252 1.45479452      1 low
## 253 1.30410959      1 low
## 254 4.08767123      2 low
## 255 1.27671233      1 low
## 256 7.16712329      1 low
## 257 1.18904110      2 low
## 258 NA                2 low
## 259 3.33150685      1 low
## 260 3.83561644      1 high
## 261 1.64109589      1 low
## 262 6.55616438      2 low
## 263 1.92876712      1 high
## 264 8.93424658      1 low
## 265 7.38630137      1 low
## 266 9.05479452      1 high
## 267 1.90958904      2 low
## 268 2.84931507      1 low
## 269 0.33150685      2 low
## 270 2.81095890      2 low
## 271 3.09589041      1 low
## 272 1.21095890      2 high

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## 273 3.20821918      2 low
## 274 0.79726027      2 low
## 275 3.36712329      2 low
## 276 1.62191781      1 low
## 277 0.23013699      1 low
## 278 2.60000000      1 low
## 279 1.71506849      2 low
## 280 1.13972603      1 low
## 281 0.50958904      1 low
## 282 3.19726027      2 high
## 284 3.06301370      1 low
## 285 2.16712329      1 high
## 286 0.59726027      1 low
## 287 3.11232877      2 low
## 288 1.80273973      1 high
## 289 2.16712329      1 low
## 290 1.59452055      2 low
## 291 5.06027397      1 high
## 292 0.48219178      2 low
## 293 1.32602740      1 low
## 294 3.47397260      2 low
## 295 6.15890411      1 low
## 296 2.85753425      2 low
## 297 2.41643836      1 low
## 298 0.93150685      2 low
## 299 3.27123288      2 low
## 300 0.88493151      1 low
## 301 2.60000000      2 low
## 302 2.47945205      2 low
## 303 1.64657534      1 high
## 304 1.00000000      1 low
## 306 1.42465753      1 low
## 307 2.36164384      1 low
## 308 4.10684932      2 low
## 309 0.93972603      2 low
## 310 1.72602740      1 low
## 311 6.02465753      1 low
## 312 1.30410959      1 low
## 313 0.77260274      2 low
## 314 1.11780822      1 low
## 315 0.38082192      2 low
## 316 1.13698630      1 low
## 317 2.25753425      1 low
## 318 NA                1 low
## 319 2.45479452      2 low
## 320 2.19178082      2 high
## 321 0.71232877      1 low
## 322 2.49863014      1 low
## 323 3.50684932      1 low
## 324 1.70958904      1 high
## 325 3.44657534      2 low
## 326 0.72328767      1 low
## 327 NA                1 low
## 328 1.31780822      1 low

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## 329 1.62739726      2 low
## 330 0.16986301      2 low
## 331 3.48493151      1 low
## 332 0.27123288      2 low
## 333 2.45479452      2 low
## 334 0.07671233      1 low
## 335 4.18630137      2 low
## 336 0.36438356      1 low
## 337 2.93972603      2 low
## 338 2.00273973      2 low
## 339 3.17534247      1 high
## 340 2.52602740      2 low
## 341 1.68219178      1 low
## 342 1.98356164      1 low
## 343 4.17260274      1 low
## 344 1.02739726      2 low
## 345 4.27123288      1 low
## 346 1.41095890      1 low
## 347 0.82191781      2 low
## 348 1.72054795      2 low
## 349 1.52602740      2 low
## 350 1.65205479      1 low
## 351 1.57260274      2 low
## 352 3.74520548      1 low
## 353 2.16712329      1 low
## 354 0.66575342      2 low
## 355 4.47123288      2 low
## 356 2.90410959      1 low
## 357 0.87945205      2 high
## 358 5.55342466      2 low
## 359 1.34794521      1 low
## 360 1.65753425      1 low
## 361 1.20547945      2 low
## 362 2.21095890      2 low
## 363 1.12602740      1 low
## 364 4.90410959      2 low
## 365 1.93150685      1 low
## 366 0.96986301      1 low
## 367 5.66301370      1 low
## 368 10.29863014     1 low
## 369 2.20821918      1 low
## 370 1.30410959      1 low
## 371 2.43561644      1 low
## 372 1.12054795      2 low
## 373 1.85479452      1 low
## 374 2.30684932      1 low
## 375 9.20821918      2 low
## 376 0.24931507      2 low
## 377 1.18904110      2 low
## 378 2.02465753      1 low
## 379 1.26575342      1 low
## 380 0.03835616      1 low
## 381 2.11780822      1 low
## 382 3.70136986      1 high

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## 383 1.92328767          2 low
## 384 0.55342466          1 low
## 385 1.53698630          2 low
## 386 2.22739726          1 high
## 387 1.49589041          1 low
## 388 1.70958904          2 low
## 389 1.21643836          2 low
## 390 1.89315068          1 low
## 391 1.50958904          1 low
## 392 2.41643836          1 high
## 393 4.61095890          1 low
## 394 5.01369863          2 low
## 395 1.78904110          2 low
## 396 3.92328767          1 low
## 397 8.47671233          1 low
## 398 1.78630137          1 low
## 399 2.08493151          2 low
## 400 1.66849315          1 low
## 401 3.89315068          2 low
## 402 1.65205479          1 low
## 403 1.54794521          1 low
## 404 1.51506849          1 low
## 405 0.78082192          1 low
## 406 3.06575342          1 low
## 407 3.71780822          2 low
## 408 1.84657534          1 low
## 409 0.47397260          2 low
## 410 0.75068493          2 low
## 411 2.12328767          1 low
## 412 0.84931507          1 low
## 413 2.36712329          1 low
## 414 0.92328767          1 low
## 415 0.46849315          2 low
## 416 0.75342466          2 low
## 417 0.61369863          1 low
## 418 0.12054795          1 low
## 419 1.43013699          1 low
## 420 3.05479452          2 low
## 421 1.55342466          1 low
## 423 1.05479452          1 low
## 424 6.19452055          1 low
## 425 0.23013699          1 low
## 426 0.20273973          2 low
## 427 0.44931507          2 low
## 428 0.92054795          2 low
## 429 1.71232877          2 high
## 430 3.53150685          1 high
## 431 13.67671233         1 high
## 432 1.83287671          1 low
## 433 1.16164384          1 low
## 434 3.65205479          1 low
## 435 1.66575342          1 high
## 436 1.73698630          1 low
## 437 2.49315068          1 low

```

```

## 438 1.05479452      1 low
## 439 1.21095890      1 high
## 440 0.03561644      1 low
## 441 2.04657534      1 high
## 442 3.08493151      1 low
## 443 6.89041096      1 low
## 444 1.60547945      2 high
## 445 0.64931507      2 low
## 446 0.12054795      1 low
## 447 5.95616438      2 low
## 448 1.33150685      1 low
## 449 1.50684932      2 low
## 450 0.35342466      1 low
## 451 1.03287671      1 low
## 452 1.83561644      1 low
## 453 4.73424658      1 low
## 454 NA                2 low
## 455 1.15890411      1 high
## 456 3.32876712      2 low
## 457 1.88767123      1 low
## 458 1.69041096      1 low
## 459 1.58356164      1 high
## 460 1.46301370      1 low
## 461 13.59178082     2 low
## 462 1.94794521      2 low
## 463 1.05479452      2 low
## 464 0.52876712      2 low
## 465 10.06575342     1 high
## 466 3.25753425      1 low
## 467 0.16438356      1 low
## 468 1.81917808      1 low
## 469 1.19178082      1 low
## 470 4.52876712      2 low
## 471 5.12328767      1 low
## 472 4.05205479      1 low
## 473 7.17808219      2 low
## 474 3.21917808      1 low
## 475 5.38356164      1 low
## 476 3.92054795      1 low
## 477 0.09589041      1 low
## 478 1.17260274      2 low
## 479 13.05479452     1 low
## 480 0.71232877      2 low
## 481 19.85753425     1 low
## 482 2.08493151      1 low
## 483 1.93150685      1 high
## 484 2.67671233      1 low
## 485 2.27123288      1 low
## 486 3.52876712      2 low
## 487 2.12876712      2 low
## 488 1.28219178      2 low
## 490 1.38356164      1 high
## 491 2.83835616      1 low
## 492 3.62739726      1 low

```

```

## 493 3.52054795      1 low
## 494 1.92054795      2 low
## 495 1.14520548      1 low
## 496 2.77534247      1 high
## 497          NA      2 low
## 498 6.09315068      1 low
## 499 1.46849315      1 low
## 500 1.36712329      1 low
## 501 6.35068493      2 low
## 502 2.38904110      1 low
## 503 1.72328767      1 low
## 504 1.21643836      2 low
## 505 0.47671233      1 low
## 506 0.49863014      1 low
## 507 4.72602740      2 low
## 508 1.27123288      2 low
## 509 0.00000000      1 low
## 510 2.27397260      1 low
## 511 3.14246575      2 low
## 512 0.32602740      2 low
## 513 1.67123288      1 low
## 514 2.73698630      2 low
## 515 1.71506849      1 low
## 516 2.86575342      2 low
## 517 1.54520548      1 low
## 518 1.14246575      1 low
## 519 0.00000000      1 low
## 520          NA      1 high
## 521 3.56438356      1 low
## 522 1.48219178      1 low

```

```

#Determine the level of miRNA expression
fit <- survfit(Surv(os, vital_status)~GENE, data=df1.cat) #Modeling based on expression
ggsurvplot(fit,data = df1.cat,title    = "hsa-mir-184", size = 1 ,xlim = c(0, 5) ,ylim=c(1,100), risk.t
font.subtitle = c(20, "bold.italic", "black"),
font.caption = c(25, "italic", "black"),
font.x = c(25, "italic", "black"),
font.y = c(25, "italic", "black"),
font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by =1)

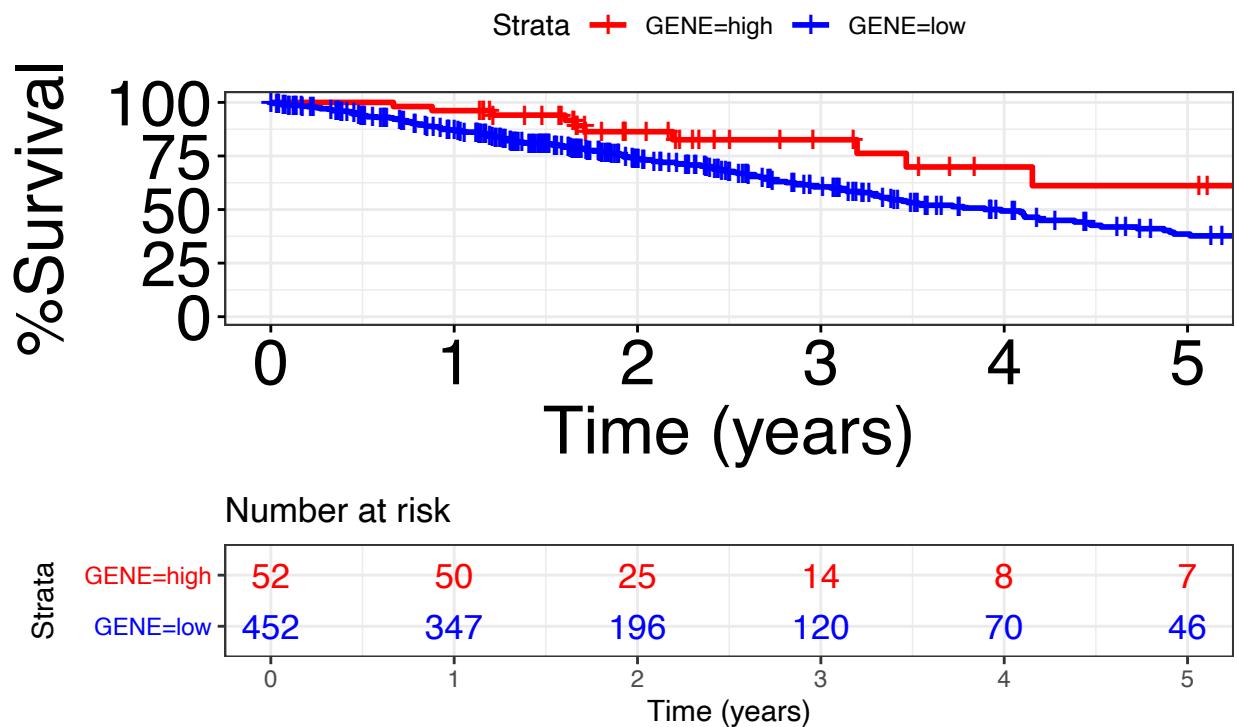
```

```

## Warning: Vectorized input to 'element_text()' is not officially supported.
## Results may be unexpected or may change in future versions of ggplot2.

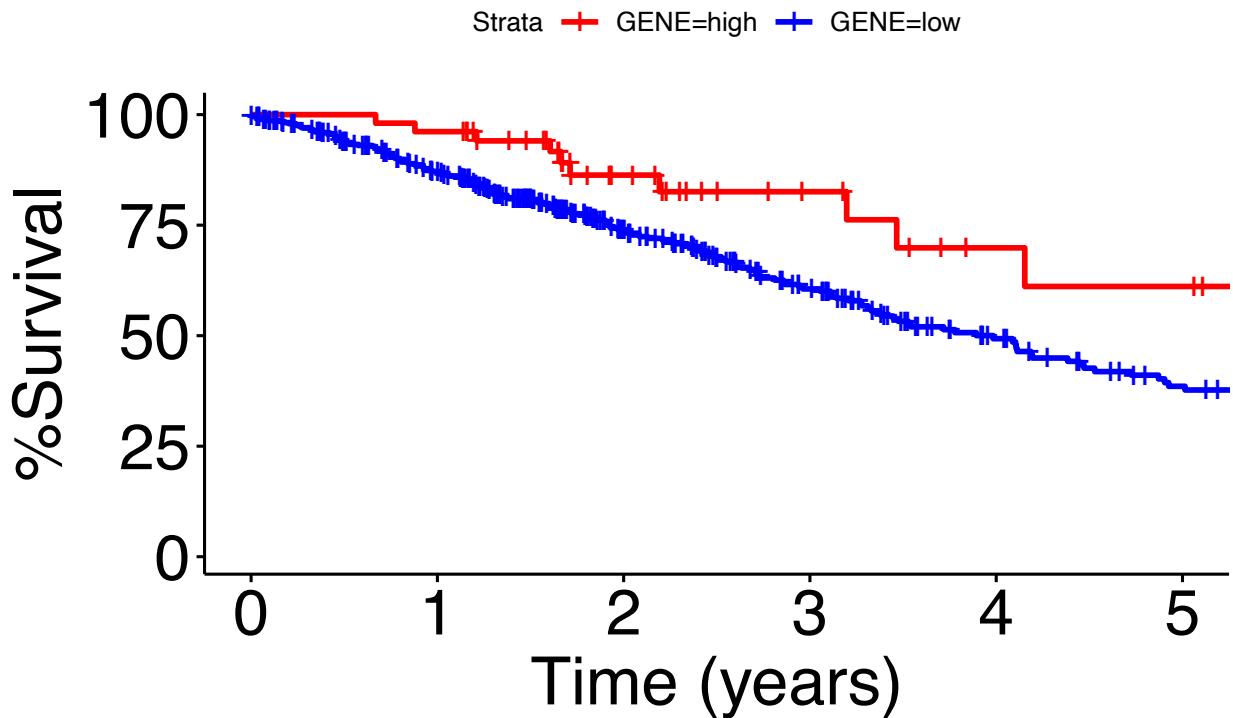
```

hsa-mir-184



```
ggsurvplot(fit,data = df1.cat,title      = "hsa-mir-184", size = 1,xlim = c(0, 5),ylim=c(1,100),xlab = "T  
font.subtitle = c(20, "bold.italic", "black"),  
font.caption = c(25, "italic", "black"),  
font.x = c(25, "italic", "black"),  
font.y = c(25, "italic", "black"),  
font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by =1)
```

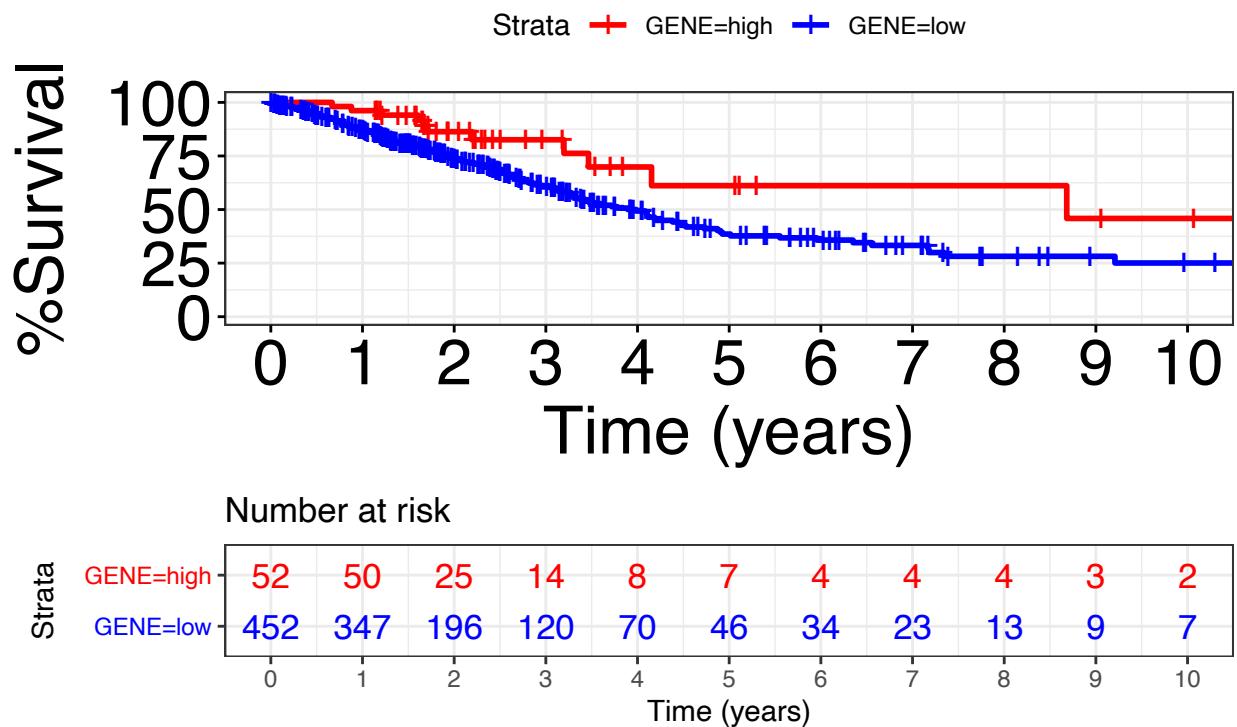
hsa-mir-184



```
ggsurvplot(fit, data = df1.cat, title      = "hsa-mir-184", size = 1, xlim = c(0, 10)  , ylim=c(1,100), end.1 = 10, end.2 = 10, font.subtitle = c(20, "bold", "black"), font.caption = c(25, "plain", "black"), font.x = c(25, "plain", "black"), font.y = c(25, "plain", "black"), font.tickslab = c(24, "plain", "black"), fun = "pct", break.x.by = 1)

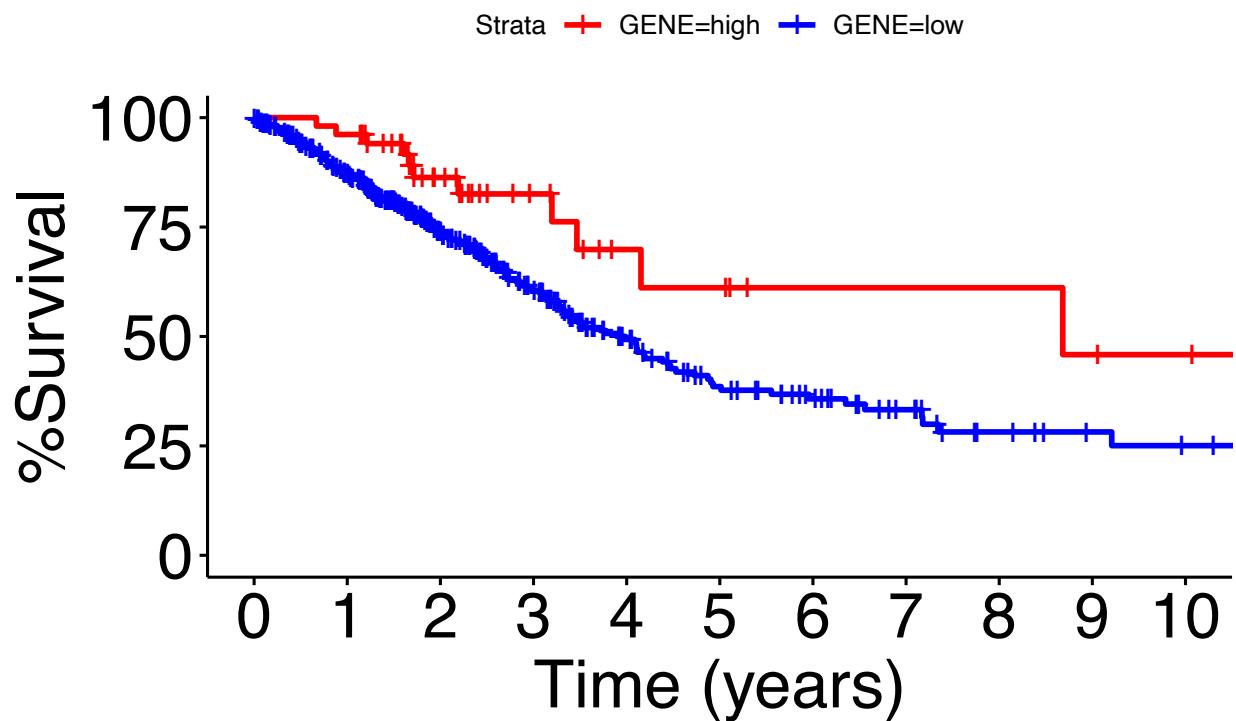
## Warning: Vectorized input to 'element_text()' is not officially supported.
## Results may be unexpected or may change in future versions of ggplot2.
```

hsa-mir-184

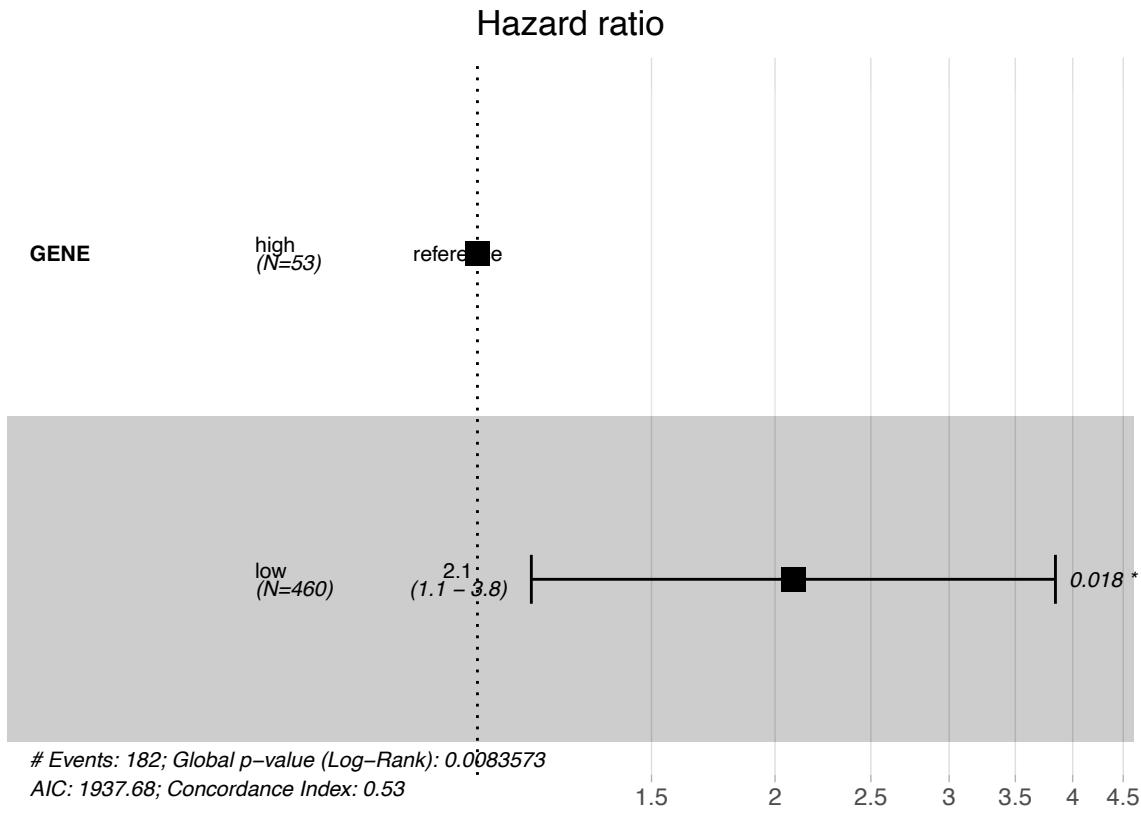


```
ggsurvplot(fit,data = df1.cat,title      = "hsa-mir-184", size = 1,xlim = c(0, 10), ylim = c(0, 100),xlab  
font.subtitle = c(20, "bold.italic", "black"),  
font.caption = c(25, "italic", "black"),  
font.x = c(25, "italic", "black"),  
font.y = c(25, "italic", "black"),  
font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by =1)
```

hsa-mir-184



```
#Cox_Regression  
fit.coxph1<- coxph(Surv(os, vital_status) ~ GENE, data=df1.cat)  
ggforest(fit.coxph1,data = df1.cat)
```



```
print(fit)
```

```
## Call: survfit(formula = Surv(os, vital_status) ~ GENE, data = df1.cat)
##
##      9 observations deleted due to missingness
##          n events median 0.95LCL 0.95UCL
## GENE=high  52      11    8.68    4.15      NA
## GENE=low   452     171    3.98    3.33    4.53
```

```
print(fit.coxph1)
```

```
## Call:
## coxph(formula = Surv(os, vital_status) ~ GENE, data = df1.cat)
##
##      coef exp(coef)  se(coef)      z      p
## GENElow 0.7359    2.0873   0.3114  2.363 0.0181
##
## Likelihood ratio test=6.96 on 1 df, p=0.008357
## n= 504, number of events= 182
## (9 observations deleted due to missingness)
```

```
coxph(Surv(os, vital_status) ~ GENE, data=df1.cat) %>%
gtsummary::tbl_regression(exp = TRUE)
```

```

## Table printed with 'knitr::kable()', not {gt}. Learn why at
## http://www.danielsjoberg.com/gtsummary/articles/rmarkdown.html
## To suppress this message, include 'message = FALSE' in code chunk header.

```

Characteristic	HR	95% CI	p-value
GENE			
high			
low	2.09	1.13, 3.84	0.018

```
summary(fit)$table
```

```

##      records n.max n.start events    *rmean *se(rmean)   median  0.95LCL
## GENE=high      52     52      52      11 11.483411   2.204069 8.682192 4.153425
## GENE=low       452    452     452     171  7.023464   0.694736 3.983562 3.328767
##      0.95UCL
## GENE=high      NA
## GENE=low     4.528767

#hsa-mir-22

#Get mirna data
#Screen the tumor sample barcode from the results: TP (primary solid tumor)

samplesTP <- TCGAquery_SampleTypes(barcode = colnames(miR_matrix), typesample = c("TP"))
mir22_exp <- miR_matrix[c("hsa-mir-22"), samplesTP]
names(mir22_exp) <- sapply(strsplit(names(mir22_exp), '-'), function(x) paste(x[1:3], collapse="-"))
clinical$GENE <- mir22_exp[clinical$submitter_id]

#Integrate vital status, deaths , last follow up visit.
df2<-subset(clinical,select=c(submitter_id,vital_status,days_to_death,days_to_last_follow_up,GENE))
df2$years_death<- df2$days_to_death/365#convert days to years
df2$years_to_last_follow_up <- df2$days_to_last_follow_up/365
df2$os<-ifelse(df2$vital_status=='Alive',df2$years_to_last_follow_up,df2$years_death)#calculate it as d
df2 <- df2[!is.na(df2$GENE),]#Remove samples with 0 expression
df2[df2$vital_status=='Dead']$vital_status <- 2
df2[df2$vital_status=='Alive']$vital_status <- 1
df2$vital_status <- as.numeric(df2$vital_status)

#Determine the optimal cutpoint of variables
df2.cut <- surv_cutpoint(df2,
  time = "os",
  event = "vital_status",
  variables = c("GENE"))
summary(df2.cut)

##      cutpoint statistic
## GENE 81157.96  1.112411

```

```

#Categorize variables
df2.cat<-surv_categorize(df2.cut,variables = NULL, labels = c("low", "high"))
head(df2.cat)

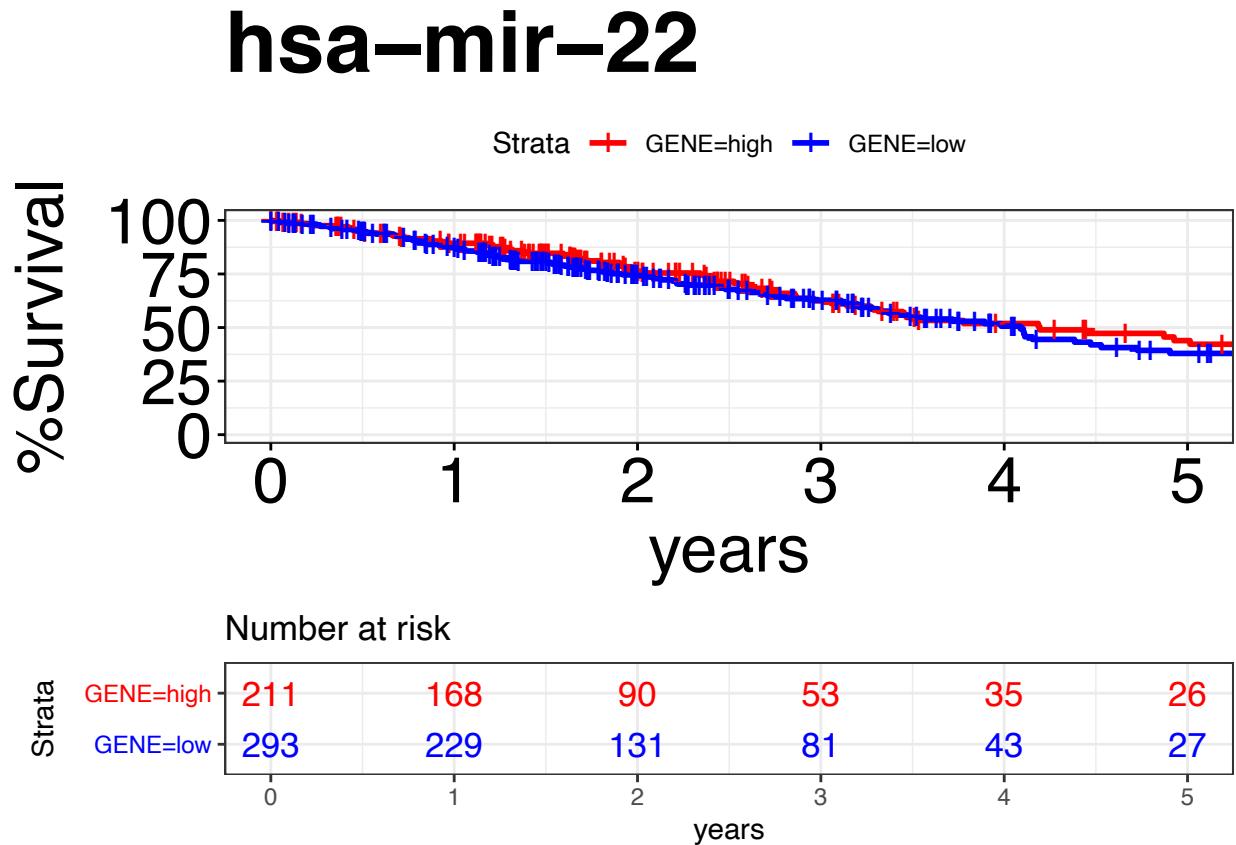
##          os vital_status GENE
## 1 2.5945205           1  low
## 2 2.9616438           2 high
## 3 0.7342466           2  low
## 4 0.8410959           2  low
## 5 0.0000000           2  low
## 6 1.4931507           1 high

#get/plot the level of miRNA expression
fit2 <- survfit(Surv(os, vital_status)~GENE, data=df2.cat) #Modeling based on expression.

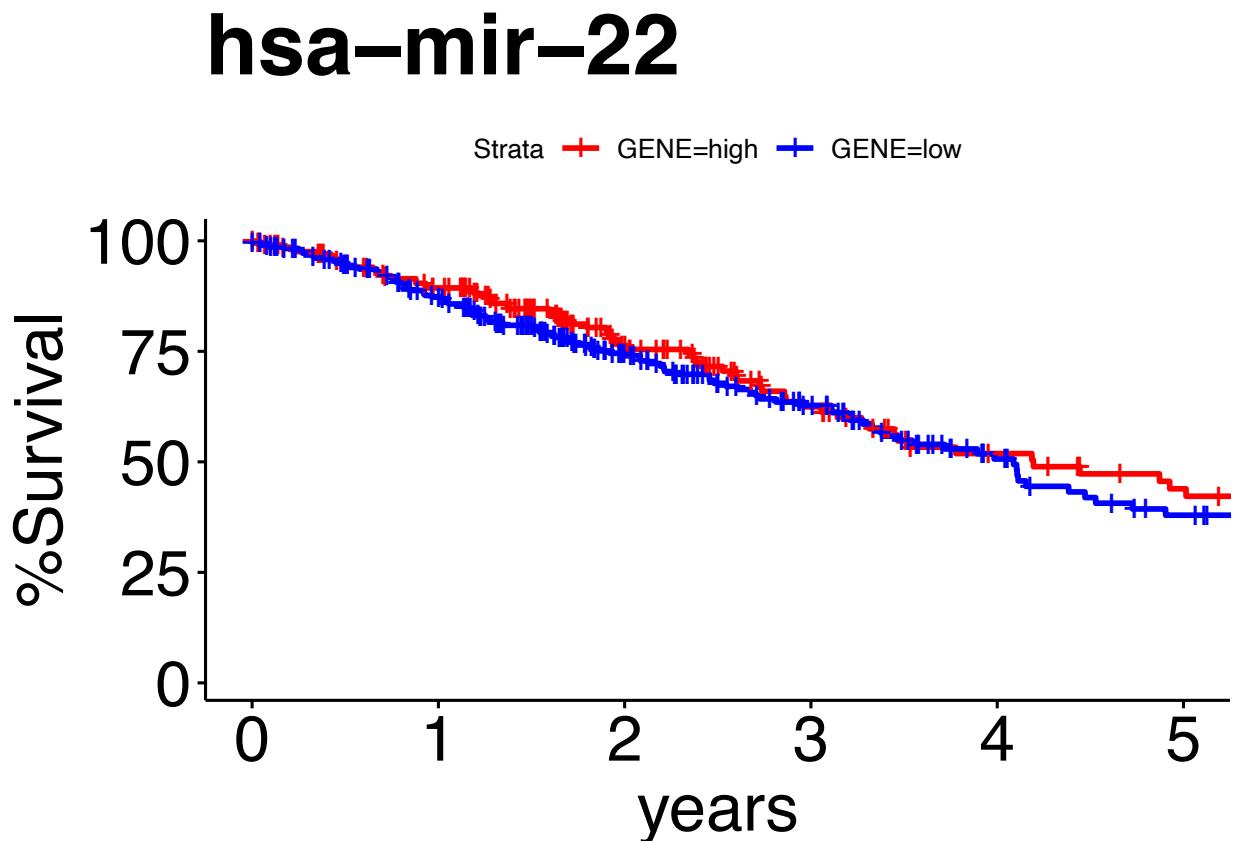
#plotKM
ggsurvplot(fit2,data = df2.cat,title = "hsa-mir-22", size = 1, xlim = c(0, 5),ylim=c(1,100), risk.table
  font.subtitle = c(20, "bold.italic", "black"),
  font.caption = c(25, "italic", "black"),
  font.x = c(25, "italic", "black"),
  font.y = c(25, "italic", "black"),
  font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by =1)

## Warning: Vectorized input to 'element_text()' is not officially supported.
## Results may be unexpected or may change in future versions of ggplot2.

```



```
ggsurvplot(fit2,data = df2.cat,title = "hsa-mir-22", size = 1, xlim = c(0, 5),ylim=c(1,100),xlab = "years"
font.subtitle = c(20, "bold.italic", "black"),
font.caption = c(25, "plain", "black"),
font.x = c(25, "plain", "black"),
font.y = c(25, "plain", "black"),
font.tickslab = c(24, "plain", "black"), fun = "pct", break.x.by =1)
```



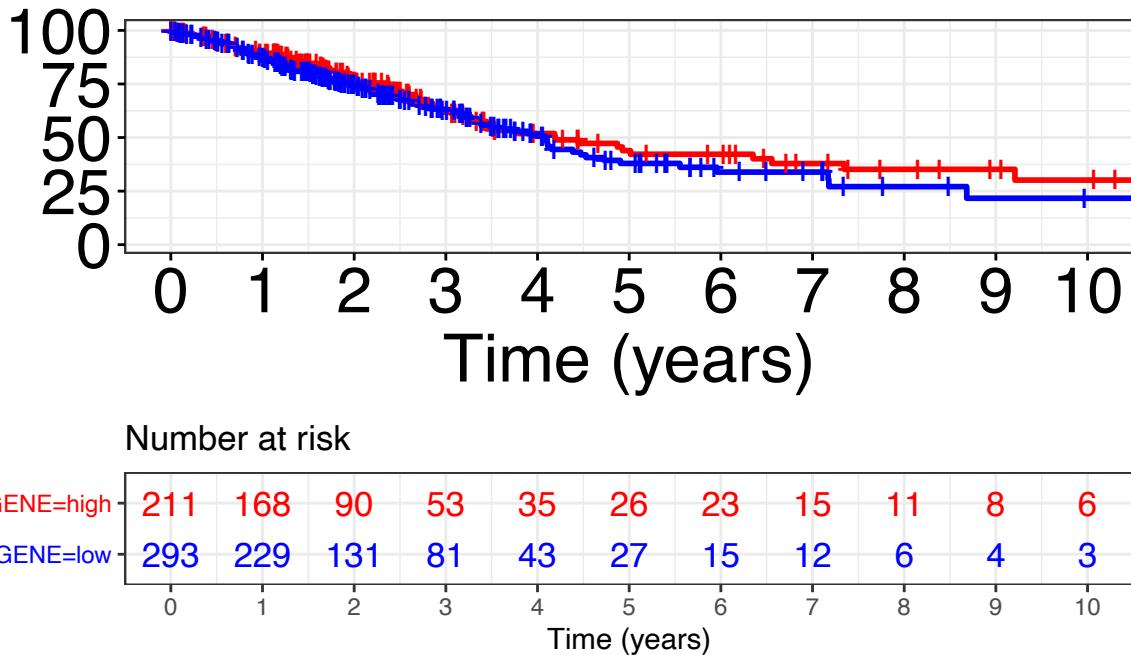
```
ggsurvplot(fit2,data = df2.cat,title = "hsa-mir-22", size = 1, xlim = c(0, 10),ylim=c(1,100), risk.table
```

```
## Warning: Vectorized input to 'element_text()' is not officially supported.
## Results may be unexpected or may change in future versions of ggplot2.
```

Survival probability (%)

hsa-mir-22

Strata + GENE=high + GENE=low

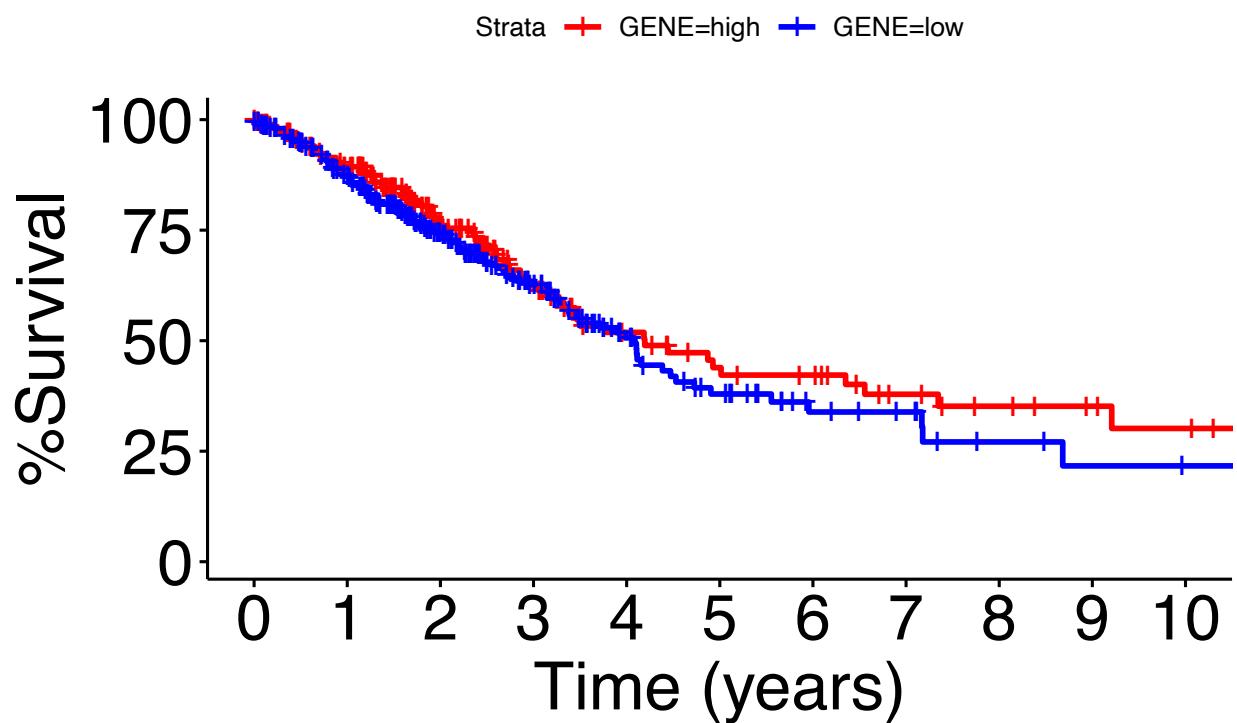


```
ggsurvplot(fit2, data = df2.cat, title = "hsa-mir-22", size = 1, xlim = c(0, 10), ylim=c(1,100), pval=TRUE,
font.subtitle = c(20, "bold.italic", "black"),
font.caption = c(25, "italic", "black"),
font.x = c(25, "italic", "black"),
font.y = c(25, "italic", "black"),
font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by =1)
```

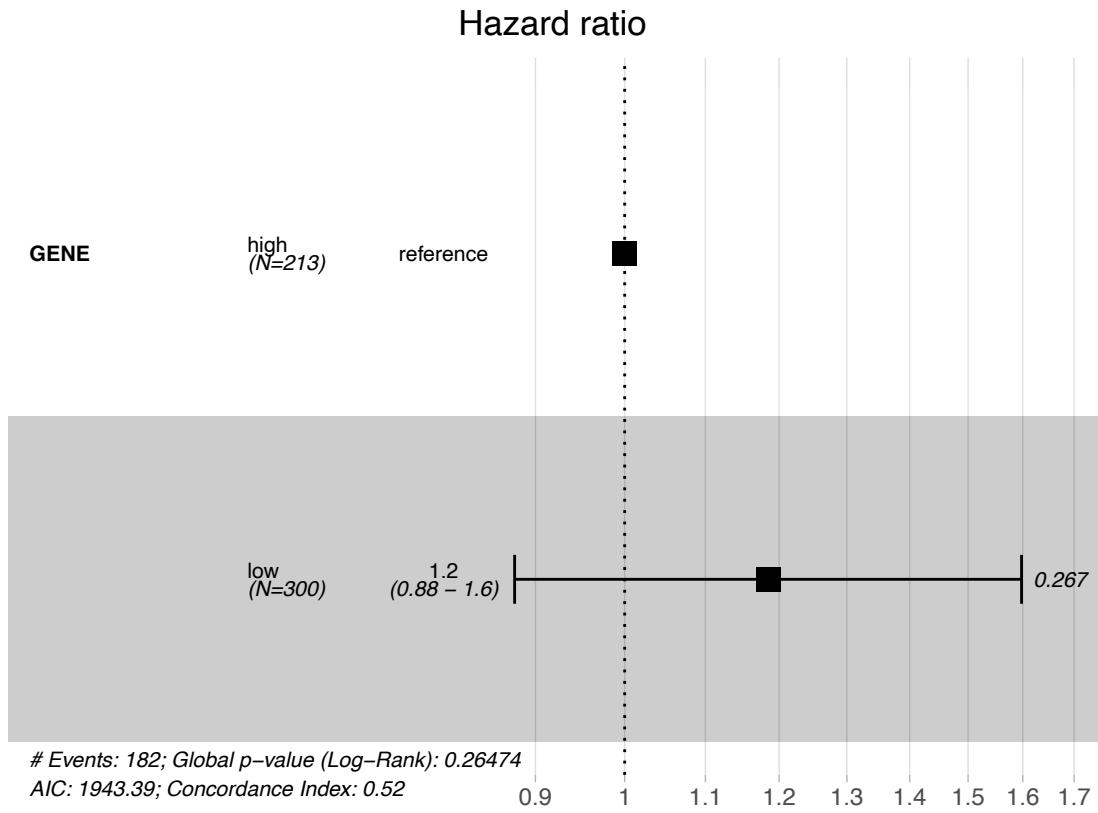
```
## Warning: Removed 1 rows containing missing values (geom_text).
```

```
## Warning: Removed 1 rows containing missing values (geom_text).
```

hsa-mir-22



```
#Cox_Regression  
fit.coxph2<- coxph(Surv(os, vital_status)~ GENE, data=df2.cat)  
ggforest(fit.coxph2,data = df2.cat)
```



```
print(fit2)
```

```
## Call: survfit(formula = Surv(os, vital_status) ~ GENE, data = df2.cat)
##
##      9 observations deleted due to missingness
##          n events median 0.95LCL 0.95UCL
## GENE=high 211     71    4.19    3.31    7.35
## GENE=low   293    111    4.09    3.37    4.73
```

```
print(fit.coxph2)
```

```
## Call:
## coxph(formula = Surv(os, vital_status) ~ GENE, data = df2.cat)
##
##      coef exp(coef)  se(coef)      z      p
## GENElow 0.1693    1.1845   0.1527  1.109 0.267
##
## Likelihood ratio test=1.24  on 1 df, p=0.2647
## n= 504, number of events= 182
## (9 observations deleted due to missingness)
```

```
coxph(Surv(os, vital_status)~GENE, data=df2.cat) %>%
gtsummary::tbl_regression(exp = TRUE)
```

```
## Table printed with 'knitr::kable()', not {gt}. Learn why at
## http://www.danielsjoberg.com/gtsummary/articles/rmarkdown.html
## To suppress this message, include 'message = FALSE' in code chunk header.
```

Characteristic	HR	95% CI	p-value
GENE			
high			
low	1.18	0.88, 1.60	0.3

```
summary(fit2)$table
```

```
##          records n.max n.start events    *rmean *se(rmean)   median 0.95LCL
## GENE=high      211     211      211      71 8.357805 0.9818366 4.194521 3.312329
## GENE=low       293     293      293     111 6.332918 0.8630399 4.087671 3.367123
##          0.95UCL
## GENE=high 7.345205
## GENE=low  4.726027
```

```
#hsa-mir-22+184
```

```
merged6 <- merge(df1.cat,df2.cat)
merged6
```

```
##          os vital_status GENE
## 1 0.00000000 1 low
## 2 0.00000000 1 low
## 3 0.00000000 1 low
## 4 0.00000000 1 low
## 5 0.00000000 1 low
## 6 0.00000000 1 low
## 7 0.00000000 2 low
## 8 0.04109589 1 low
## 9 0.04931507 2 low
## 10 0.06027397 2 low
## 11 0.07671233 1 low
## 12 0.09041096 2 low
## 13 0.09863014 1 low
## 14 0.12054795 1 low
## 15 0.12054795 1 low
## 16 0.13150685 1 low
## 17 0.16986301 1 low
## 18 0.16986301 2 low
## 19 0.21643836 1 low
## 20 0.23013699 1 low
## 21 0.23013699 1 low
## 22 0.23013699 1 low
## 23 0.23013699 1 low
## 24 0.24931507 2 low
## 25 0.26575342 2 low
## 26 0.27123288 2 low
```

## 27	0.31780822	2	low
## 28	0.32602740	1	low
## 29	0.32602740	2	low
## 30	0.33150685	2	low
## 31	0.38082192	2	low
## 32	0.38630137	1	low
## 33	0.41369863	1	low
## 34	0.46849315	2	low
## 35	0.47397260	2	low
## 36	0.47671233	1	low
## 37	0.49041096	1	low
## 38	0.49041096	2	low
## 39	0.49863014	1	low
## 40	0.50410959	1	low
## 41	0.50958904	1	low
## 42	0.50958904	1	low
## 43	0.51232877	2	low
## 44	0.52876712	2	low
## 45	0.55342466	1	low
## 46	0.57534247	2	low
## 47	0.61643836	1	low
## 48	0.63013699	1	low
## 49	0.66849315	2	low
## 50	0.66849315	2	low
## 51	0.70410959	2	low
## 52	0.70684932	2	low
## 53	0.72328767	1	low
## 54	0.73424658	2	low
## 55	0.75342466	2	low
## 56	0.76986301	2	low
## 57	0.77260274	2	low
## 58	0.78082192	1	low
## 59	0.78630137	1	low
## 60	0.79726027	2	low
## 61	0.82191781	2	low
## 62	0.83013699	2	low
## 63	0.84109589	1	low
## 64	0.84109589	2	low
## 65	0.84383562	2	low
## 66	0.84931507	1	low
## 67	0.87945205	2	high
## 68	0.87945205	2	high
## 69	0.88493151	1	low
## 70	0.92054795	2	low
## 71	0.92054795	2	low
## 72	0.92054795	2	low
## 73	0.92054795	2	low
## 74	0.92876712	2	low
## 75	0.96164384	1	low
## 76	0.96986301	2	low
## 77	1.00000000	1	low
## 78	1.01369863	2	low
## 79	1.01917808	1	low
## 80	1.02739726	2	low

## 81	1.03013699	2 low
## 82	1.05479452	1 low
## 83	1.05479452	1 low
## 84	1.05479452	2 low
## 85	1.12054795	2 low
## 86	1.13424658	1 low
## 87	1.13424658	2 low
## 88	1.13698630	1 high
## 89	1.13698630	1 high
## 90	1.14246575	1 low
## 91	1.16164384	1 low
## 92	1.16986301	1 low
## 93	1.17260274	2 low
## 94	1.18904110	2 low
## 95	1.18904110	2 low
## 96	1.19178082	1 high
## 97	1.19178082	1 high
## 98	1.19178082	1 low
## 99	1.19178082	1 low
## 100	1.20547945	2 low
## 101	1.21643836	2 low
## 102	1.21643836	2 low
## 103	1.22191781	1 low
## 104	1.22739726	1 low
## 105	1.22739726	1 low
## 106	1.22739726	1 low
## 107	1.22739726	1 low
## 108	1.24657534	1 low
## 109	1.25205479	2 low
## 110	1.26027397	2 low
## 111	1.28219178	2 low
## 112	1.28219178	2 low
## 113	1.30410959	1 low
## 114	1.30410959	1 low
## 115	1.30410959	1 low
## 116	1.30410959	1 low
## 117	1.30410959	1 low
## 118	1.30410959	1 low
## 119	1.30410959	1 low
## 120	1.30410959	1 low
## 121	1.30410959	1 low
## 122	1.30958904	1 low
## 123	1.30958904	2 low
## 124	1.31780822	1 low
## 125	1.32602740	1 low
## 126	1.33150685	1 low
## 127	1.33424658	1 low
## 128	1.33698630	2 low
## 129	1.34794521	1 low
## 130	1.38356164	1 high
## 131	1.42465753	1 low
## 132	1.44109589	1 low
## 133	1.45479452	1 low
## 134	1.47123288	1 low

```

## 135 1.47671233      1 high
## 136 1.48219178      1 low
## 137 1.49863014      1 low
## 138 1.50684932      2 low
## 139 1.51506849      1 low
## 140 1.52602740      2 low
## 141 1.53698630      2 low
## 142 1.54520548      1 low
## 143 1.54794521      1 low
## 144 1.55342466      1 low
## 145 1.55616438      1 low
## 146 1.55616438      1 low
## 147 1.55616438      1 low
## 148 1.55616438      1 low
## 149 1.55616438      1 low
## 150 1.55616438      1 low
## 151 1.55616438      1 low
## 152 1.55616438      1 low
## 153 1.55616438      1 low
## 154 1.57260274      2 low
## 155 1.58356164      1 high
## 156 1.58356164      1 high
## 157 1.58356164      1 high
## 158 1.58356164      1 high
## 159 1.58356164      1 low
## 160 1.61917808      1 low
## 161 1.62191781      1 low
## 162 1.62465753      2 low
## 163 1.65205479      1 high
## 164 1.65205479      1 high
## 165 1.65205479      1 high
## 166 1.65753425      1 low
## 167 1.65753425      1 low
## 168 1.66301370      2 high
## 169 1.66849315      1 low
## 170 1.67123288      1 high
## 171 1.67123288      1 low
## 172 1.67123288      1 low
## 173 1.67123288      1 low
## 174 1.67123288      1 low
## 175 1.69041096      1 low
## 176 1.69041096      1 low
## 177 1.70958904      1 high
## 178 1.71506849      1 high
## 179 1.71506849      1 low
## 180 1.71506849      2 low
## 181 1.72054795      2 low
## 182 1.72602740      1 low
## 183 1.73698630      1 low
## 184 1.78356164      1 low
## 185 1.78356164      1 low
## 186 1.78356164      1 low
## 187 1.78356164      1 low
## 188 1.78630137      1 low

```

```

## 189 1.78630137      1 low
## 190 1.78630137      1 low
## 191 1.78630137      1 low
## 192 1.78904110      1 low
## 193 1.78904110      2 low
## 194 1.80273973      1 high
## 195 1.80273973      1 high
## 196 1.81917808      1 low
## 197 1.82465753      2 low
## 198 1.83287671      1 low
## 199 1.83561644      1 low
## 200 1.83561644      1 low
## 201 1.83561644      1 low
## 202 1.83561644      1 low
## 203 1.85479452      1 low
## 204 1.85479452      2 low
## 205 1.88767123      1 low
## 206 1.89041096      1 low
## 207 1.89315068      1 low
## 208 1.92054795      1 high
## 209 1.92054795      2 low
## 210 1.92876712      1 high
## 211 1.92876712      1 high
## 212 1.92876712      1 high
## 213 1.92876712      1 high
## 214 1.93150685      1 high
## 215 1.93150685      1 low
## 216 1.96712329      1 low
## 217 1.96986301      1 low
## 218 1.98082192      1 low
## 219 1.99452055      1 low
## 220 10.06575342     1 high
## 221 10.79452055    1 low
## 222 13.59178082    2 low
## 223 13.67671233    1 high
## 224 19.85753425    1 low
## 225 2.00273973     2 low
## 226 2.03013699     1 low
## 227 2.08219178     2 low
## 228 2.08493151     1 low
## 229 2.08493151     1 low
## 230 2.08493151     2 low
## 231 2.11780822     1 low
## 232 2.12328767     1 low
## 233 2.12876712     2 low
## 234 2.16712329     1 high
## 235 2.16712329     1 low
## 236 2.16712329     1 low
## 237 2.16712329     1 low
## 238 2.16712329     1 low
## 239 2.16712329     1 low
## 240 2.16712329     1 low
## 241 2.16712329     1 low
## 242 2.16712329     1 low

```

```

## 243 2.16712329      1 low
## 244 2.20547945      1 high
## 245 2.21095890      2 low
## 246 2.21369863      2 low
## 247 2.22739726      1 high
## 248 2.25753425      1 low
## 249 2.25753425      1 low
## 250 2.25753425      1 low
## 251 2.25753425      1 low
## 252 2.26301370      2 low
## 253 2.26575342      1 low
## 254 2.27123288      1 low
## 255 2.27397260      1 low
## 256 2.29863014      1 high
## 257 2.30684932      1 low
## 258 2.31506849      1 low
## 259 2.36438356      1 low
## 260 2.36712329      1 low
## 261 2.38904110      1 low
## 262 2.41643836      1 high
## 263 2.41643836      1 low
## 264 2.45479452      2 low
## 265 2.45479452      2 low
## 266 2.45479452      2 low
## 267 2.45479452      2 low
## 268 2.47945205      2 low
## 269 2.49315068      1 low
## 270 2.49863014      1 low
## 271 2.50136986      1 high
## 272 2.54520548      2 low
## 273 2.54794521      1 low
## 274 2.59452055      1 low
## 275 2.60821918      2 low
## 276 2.67397260      2 low
## 277 2.70410959      2 low
## 278 2.70684932      1 low
## 279 2.72602740      2 low
## 280 2.72602740      2 low
## 281 2.81095890      2 low
## 282 2.83835616      1 low
## 283 2.84931507      1 low
## 284 2.90410959      1 low
## 285 2.93424658      1 low
## 286 2.93698630      1 low
## 287 2.93972603      2 low
## 288 3.00547945      1 low
## 289 3.08219178      1 low
## 290 3.08493151      1 low
## 291 3.08493151      1 low
## 292 3.08493151      1 low
## 293 3.08493151      1 low
## 294 3.10958904      2 low
## 295 3.11232877      2 low
## 296 3.14520548      1 low

```

```

## 297 3.16986301      1 low
## 298 3.20821918      2 low
## 299 3.21917808      1 low
## 300 3.22739726      1 low
## 301 3.25753425      1 low
## 302 3.25753425      1 low
## 303 3.25753425      1 low
## 304 3.25753425      1 low
## 305 3.27945205      2 low
## 306 3.32876712      2 low
## 307 3.36712329      2 low
## 308 3.37808219      1 low
## 309 3.38356164      2 low
## 310 3.46575342      2 high
## 311 3.47397260      2 low
## 312 3.48493151      1 low
## 313 3.52054795      1 low
## 314 3.53150685      1 high
## 315 3.54246575      2 low
## 316 3.56438356      1 low
## 317 3.57534247      1 low
## 318 3.62739726      1 low
## 319 3.65205479      1 low
## 320 3.71780822      2 low
## 321 3.74520548      1 low
## 322 3.75068493      1 low
## 323 3.89315068      2 low
## 324 3.91506849      1 low
## 325 3.92054795      1 low
## 326 3.92328767      1 low
## 327 3.98356164      2 low
## 328 4.03835616      1 low
## 329 4.05205479      1 low
## 330 4.08767123      2 low
## 331 4.10410959      2 low
## 332 4.10684932      2 low
## 333 4.11232877      2 low
## 334 4.17260274      1 low
## 335 4.38356164      2 low
## 336 4.47123288      2 low
## 337 4.52876712      2 low
## 338 4.61095890      1 low
## 339 4.72602740      2 low
## 340 4.73424658      1 low
## 341 4.79452055      1 low
## 342 4.90410959      2 low
## 343 5.12328767      1 low
## 344 5.38356164      1 low
## 345 5.40821918      1 low
## 346 5.55342466      2 low
## 347 5.65753425      1 low
## 348 5.66301370      1 low
## 349 5.77808219      1 low
## 350 5.92054795      1 low

```

```

## 351 5.95616438      2 low
## 352 6.19452055      1 low
## 353 6.48767123      1 low
## 354 6.89041096      1 low
## 355 7.09589041      1 low
## 356 7.10958904      1 low
## 357 7.16986301      2 low
## 358 7.17808219      2 low
## 359 7.33150685      1 low
## 360 7.75890411      1 low
## 361 8.47671233      1 low
## 362 9.05479452      1 high
## 363 9.95890411      1 low
## 364 NA               1 high
## 365 NA               1 high
## 366 NA               1 low
## 367 NA               1 low
## 368 NA               1 low
## 369 NA               1 low
## 370 NA               1 low
## 371 NA               1 low
## 372 NA               1 low
## 373 NA               1 low
## 374 NA               1 low
## 375 NA               1 low
## 376 NA               1 low
## 377 NA               1 low
## 378 NA               2 low
## 379 NA               2 low
## 380 NA               2 low
## 381 NA               2 low
## 382 NA               2 low
## 383 NA               2 low
## 384 NA               2 low
## 385 NA               2 low
## 386 NA               2 low
## 387 NA               2 low
## 388 NA               2 low
## 389 NA               2 low
## 390 NA               2 low
## 391 NA               2 low
## 392 NA               2 low
## 393 NA               2 low

```

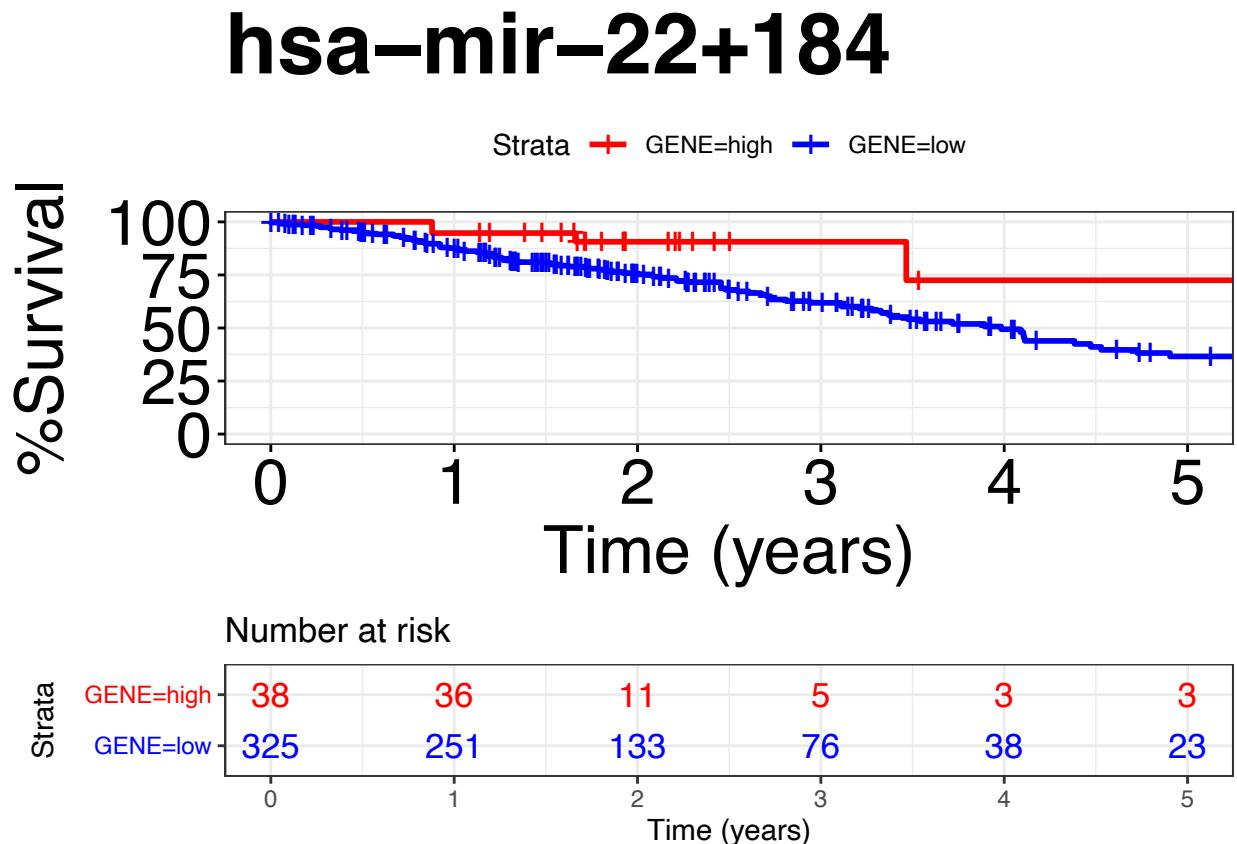
```

fit8<- survfit(Surv(os, vital_status)~ GENE, data=merged6)

#Plot_KM
ggsurvplot(fit8,data = merged6,title     = "hsa-mir-22+184", size = 1 ,xlim = c(0, 5), risk.table = TRUE
            font.subtitle = c(20, "bold.italic", "black"),
            font.caption = c(25, "italic", "black"),
            font.x = c(25, "italic", "black"),
            font.y = c(25, "italic", "black"),
            font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by =1, ylim=c(0,100))

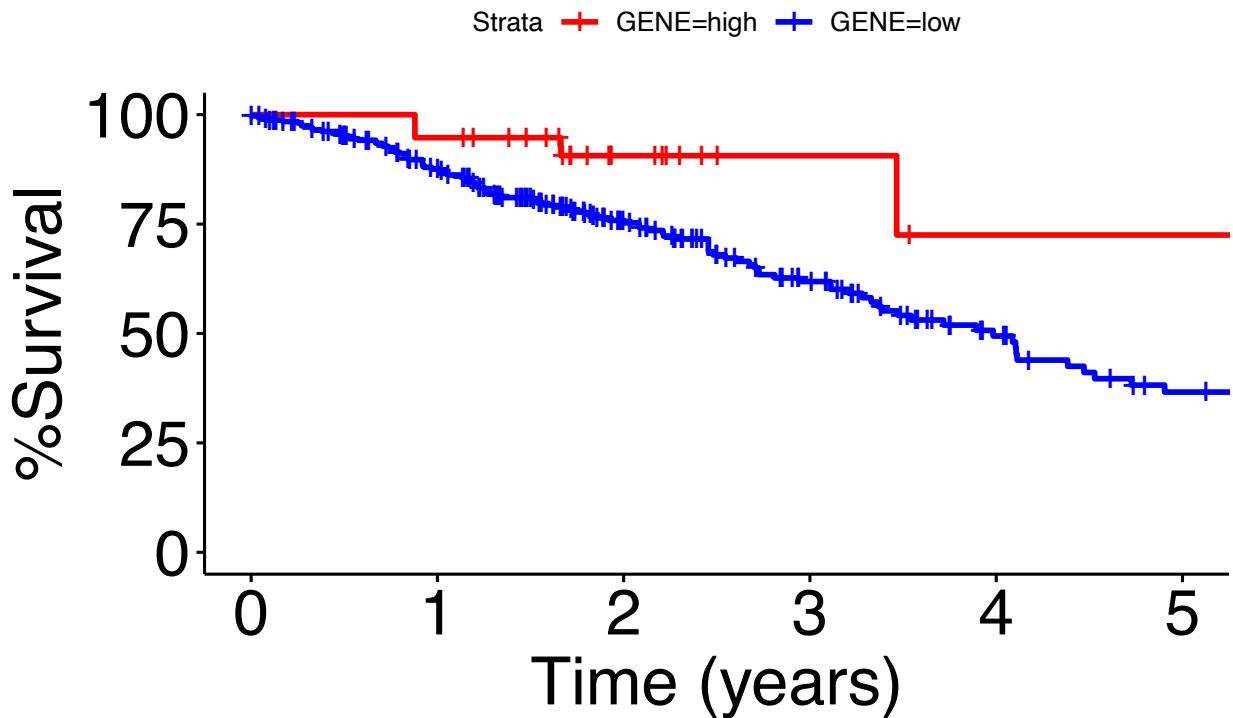
```

```
## Warning: Vectorized input to 'element_text()' is not officially supported.
## Results may be unexpected or may change in future versions of ggplot2.
```



```
ggsurvplot(fit8, data = merged6, title      = "hsa-mir-22+184", size = 1, xlim = c(0, 5), xlab = "Time (years)",
            font.subtitle = c(20, "bold.italic", "black"),
            font.caption = c(25, "italic", "black"),
            font.x = c(25, "italic", "black"),
            font.y = c(25, "italic", "black"),
            font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by = 1, ylim=c(0,100))
```

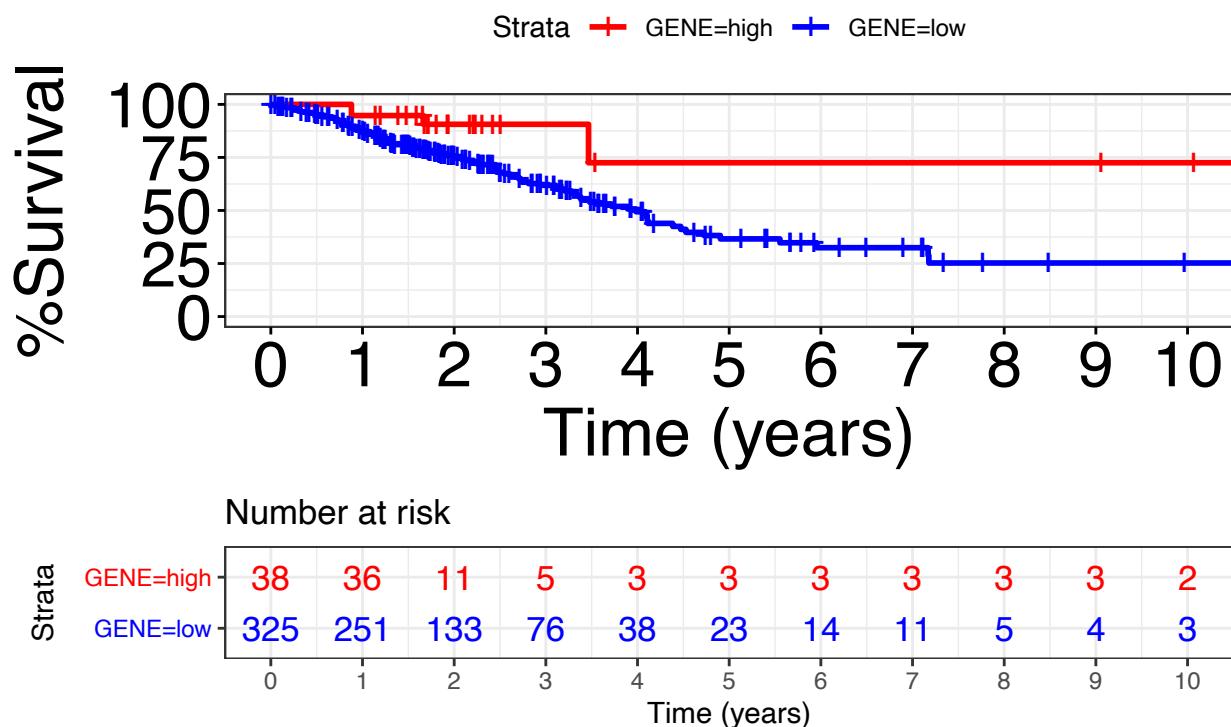
hsa-mir-22+184



```
ggsurvplot(fit8,data = merged6,title      = "hsa-mir-22+184", size = 1 ,xlim = c(0, 10), risk.table = TRUE  
font.subtitle = c(20, "bold.italic", "black"),  
font.caption = c(25, "italic", "black"),  
font.x = c(25, "italic", "black"),  
font.y = c(25, "italic", "black"),  
font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by =1, ylim=c(0,100))
```

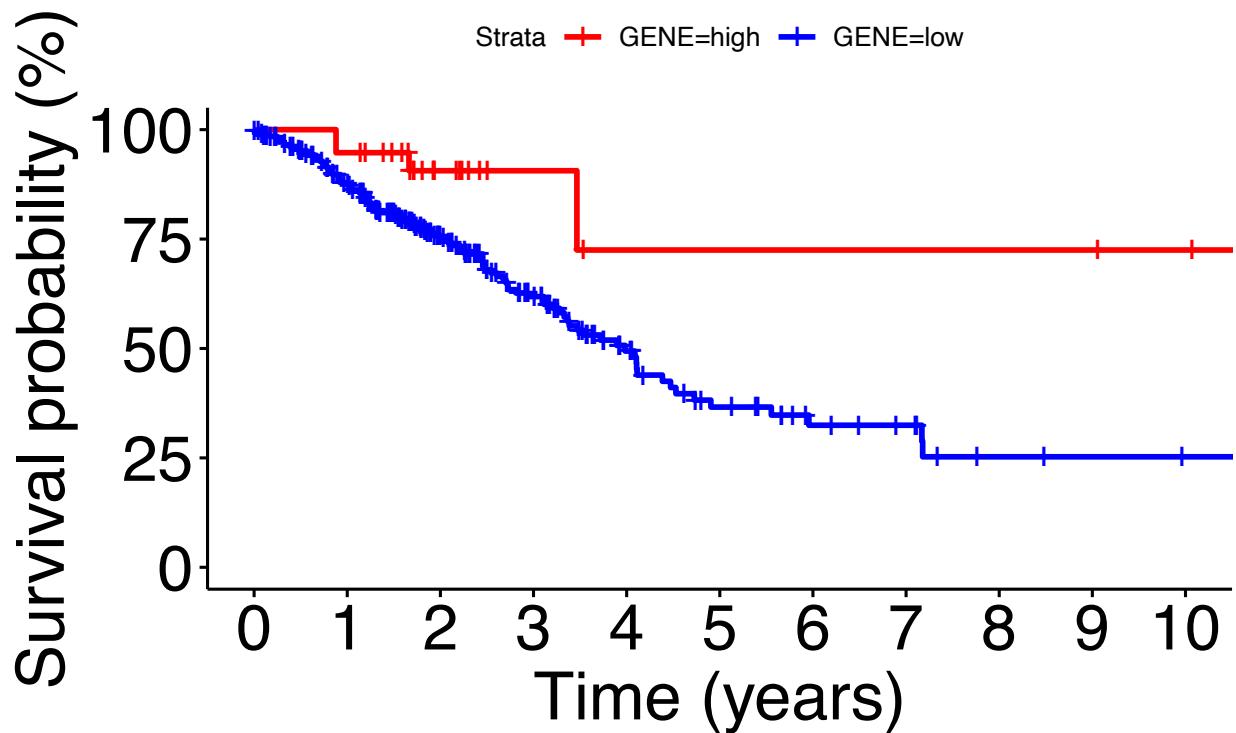
```
## Warning: Vectorized input to 'element_text()' is not officially supported.  
## Results may be unexpected or may change in future versions of ggplot2.
```

hsa-mir-22+184

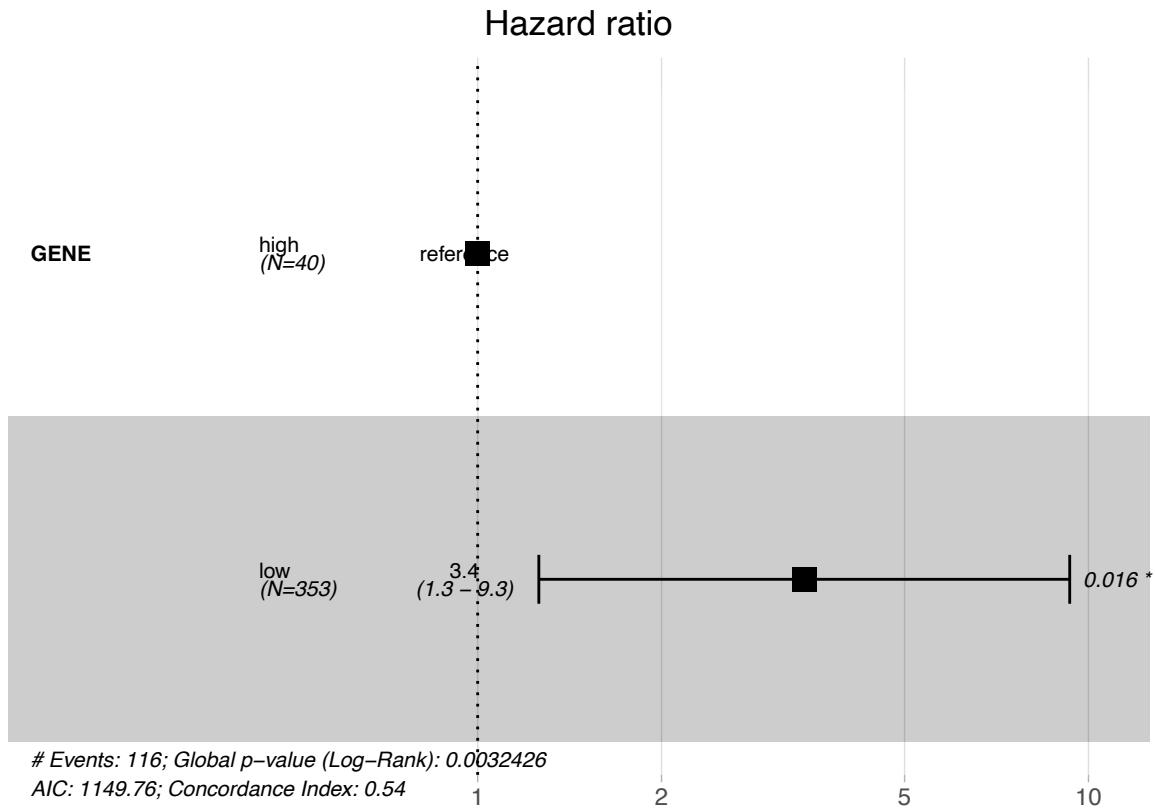


```
ggsurvplot(fit8,data = merged6,title      = "hsa-mir-22+184", size = 1,xlim = c(0, 10), xlab = "Time (year",  
font.subtitle = c(20, "bold.italic", "black"),  
font.caption = c(25, "italic", "black"),  
font.x = c(25, "italic", "black"),  
font.y = c(25, "italic", "black"),  
font.tickslab = c(24, "italic", "black"), fun = "pct", break.x.by =1, ylim=c(0,100))
```

hsa-mir-22+184



```
#Cox_Regression  
fit.coxph1<- coxph(Surv(os, vital_status)~ GENE, data=merged6)  
ggforest(fit.coxph1,data = merged6)
```



```
print(fit8)
```

```
## Call: survfit(formula = Surv(os, vital_status) ~ GENE, data = merged6)
##
##      30 observations deleted due to missingness
##          n events median 0.95LCL 0.95UCL
## GENE=high  38      4     NA    3.47      NA
## GENE=low   325    112    3.98    3.37    4.73
```

```
print(fit.coxph1)
```

```
## Call:
## coxph(formula = Surv(os, vital_status) ~ GENE, data = merged6)
##
##      coef exp(coef)  se(coef)      z      p
## GENElow 1.2313    3.4255   0.5105  2.412 0.0159
##
## Likelihood ratio test=8.67  on 1 df, p=0.003243
## n= 363, number of events= 116
## (30 observations deleted due to missingness)
```

```
coxph(Surv(os, vital_status) ~ GENE, data=merged6) %>%
gtsummary::tbl_regression(exp = TRUE)
```

```
## Table printed with 'knitr::kable()', not {gt}. Learn why at
## http://www.danielsjoberg.com/gtsummary/articles/rmarkdown.html
## To suppress this message, include 'message = FALSE' in code chunk header.
```

Characteristic	HR	95% CI	p-value
GENE			
high			
low	3.43	1.26, 9.32	0.016

```
summary(fit8)$table
```

```
##          records n.max n.start events    *rmean *se(rmean)   median 0.95LCL
## GENE=high      38     38      38      4 15.138481  2.7782600      NA 3.465753
## GENE=low       325    325     325     112  6.559784  0.9027472 3.983562 3.367123
##          0.95UCL
## GENE=high      NA
## GENE=low     4.726027
```

Combining Plasma Extracellular Vesicle Let-7b-5p, miR-184 and Circulating miR-22-3p Levels for NSCLC Diagnosis and Drug Resistance Prediction

G. P. Vadla¹, B. Daghpat¹, N. Patterson¹, V. Ahmad¹, G. Perez^{1*}, A. Garcia^{1*}, Y. Manjunath², J.T. Kaifi^{2,3}, G. Li², and C.Y. Chabu^{1,2,3†}

Supplementary information 2. R session information for covariate analyses

```
## – Session info _____
## setting value
## version R version 4.1.0 (2021-05-18)
## os   Ubuntu 20.04.2 LTS
## system x86_64, linux-gnu
## ui    X11
## language (EN)
## collate en_US.UTF-8
## ctype en_US.UTF-8
## tz   Etc/UTC
## date 2021-08-05

## – Packages _____
package      * version date   lib source
## abind       1.4-5  2016-07-21 [1] RSPM (R 4.0.3)
## assertthat   0.2.1  2019-03-21 [1] RSPM (R 4.1.0)
## backports    1.2.1  2020-12-09 [1] RSPM (R 4.1.0)
## Biobase     * 2.52.0 2021-05-19 [1] Bioconductor
## BiocGenerics * 0.38.0 2021-05-19 [1] Bioconductor
## broom        0.7.9  2021-07-27 [1] RSPM (R 4.0.5)
## bslib        0.2.5.1 2021-05-18 [1] RSPM (R 4.1.0)
## cachem       1.0.5  2021-05-15 [1] RSPM (R 4.1.0)
## callr        3.7.0  2021-04-20 [1] RSPM (R 4.1.0)
## car          3.0-11 2021-06-27 [1] RSPM (R 4.0.5)
## carData      3.0-4   2020-05-22 [1] RSPM (R 4.0.3)
## cellranger   1.1.0  2016-07-27 [1] RSPM (R 4.1.0)
## cli          3.0.1  2021-07-17 [1] RSPM (R 4.0.5)
## colorspace   2.0-2   2021-06-24 [1] RSPM (R 4.1.0)
## crayon       1.4.1  2021-02-08 [1] RSPM (R 4.1.0)
## crosstalk   1.1.1  2021-01-12 [1] RSPM (R 4.0.3)
## curl         4.3.2  2021-06-23 [1] RSPM (R 4.1.0)
## data.table   1.14.0 2021-02-21 [1] RSPM (R 4.1.0)
## DBI          1.1.1  2021-01-15 [1] RSPM (R 4.1.0)
## desc         1.3.0  2021-03-05 [1] RSPM (R 4.1.0)
## devtools     2.4.2  2021-06-07 [1] RSPM (R 4.1.0)
## digest       0.6.27 2020-10-24 [1] RSPM (R 4.1.0)
## dplyr       * 1.0.7  2021-06-18 [1] RSPM (R 4.1.0)
## DT          * 0.18   2021-04-14 [1] RSPM (R 4.0.4)
```

```
## edgeR      * 3.34.0 2021-05-19 [1] Bioconductor
## ellipsis    0.3.2  2021-04-29 [1] RSPM (R 4.1.0)
## evaluate    0.14   2019-05-28 [1] RSPM (R 4.1.0)
## fansi       0.5.0   2021-05-25 [1] RSPM (R 4.1.0)
## farver      2.1.0   2021-02-28 [1] RSPM (R 4.1.0)
## fastmap     1.1.0   2021-01-25 [1] RSPM (R 4.1.0)
## forcats     0.5.1   2021-01-27 [1] RSPM (R 4.1.0)
## foreign     0.8-81  2020-12-22 [2] CRAN (R 4.1.0)
## fs          1.5.0   2020-07-31 [1] RSPM (R 4.1.0)
## generics    0.1.0   2020-10-31 [1] RSPM (R 4.1.0)
## GGally      * 2.1.2   2021-06-21 [1] RSPM (R 4.0.5)
## ggplot2     * 3.3.5   2021-06-25 [1] RSPM (R 4.1.0)
## ggpubr      * 0.4.0   2020-06-27 [1] RSPM (R 4.0.3)
## ggrepel     * 0.9.1   2021-01-15 [1] RSPM (R 4.0.5)
## ggsignif    0.6.2   2021-06-14 [1] RSPM (R 4.0.5)
## glue         1.4.2   2020-08-27 [1] RSPM (R 4.1.0)
## gtable      0.3.0   2019-03-25 [1] RSPM (R 4.1.0)
## haven        2.4.1   2021-04-23 [1] RSPM (R 4.1.0)
## highr        0.9    2021-04-16 [1] RSPM (R 4.1.0)
## hms          1.1.0   2021-05-17 [1] RSPM (R 4.1.0)
## htmltools    0.5.1.1 2021-01-22 [1] RSPM (R 4.1.0)
## htmlwidgets  1.5.3   2020-12-10 [1] RSPM (R 4.1.0)
## httr         1.4.2   2020-07-20 [1] RSPM (R 4.1.0)
## jquerylib    0.1.4   2021-04-26 [1] RSPM (R 4.1.0)
## jsonlite     1.7.2   2020-12-09 [1] RSPM (R 4.1.0)
## kableExtra   * 1.3.4   2021-02-20 [1] RSPM (R 4.0.3)
## knitr        1.33    2021-04-24 [1] RSPM (R 4.1.0)
## labeling     0.4.2   2020-10-20 [1] RSPM (R 4.1.0)
## lattice      0.20-44  2021-05-02 [2] CRAN (R 4.1.0)
## lazyeval     0.2.2   2019-03-15 [1] RSPM (R 4.0.3)
## lifecycle    1.0.0   2021-02-15 [1] RSPM (R 4.1.0)
## limma        * 3.48.1  2021-06-24 [1] Bioconductor
## locfit       1.5-9.4 2020-03-25 [1] RSPM (R 4.0.3)
## magrittr     * 2.0.1   2020-11-17 [1] RSPM (R 4.1.0)
## Matrix        1.3-3   2021-05-04 [2] CRAN (R 4.1.0)
## memoise      2.0.0   2021-01-26 [1] RSPM (R 4.1.0)
## mgcv         1.8-35  2021-04-18 [2] CRAN (R 4.1.0)
## mime          0.11   2021-06-23 [1] RSPM (R 4.1.0)
## munsell      0.5.0   2018-06-12 [1] RSPM (R 4.1.0)
## nlme         3.1-152  2021-02-04 [2] CRAN (R 4.1.0)
## openxlsx     4.2.4   2021-06-16 [1] RSPM (R 4.0.5)
## pcaMethods   * 1.84.0  2021-05-19 [1] Bioconductor
## pheatmap     * 1.0.12  2019-01-04 [1] RSPM (R 4.0.3)
```

```
## pillar      1.6.1 2021-05-16 [1] RSPM (R 4.1.0)
## pkgbuild    1.2.0 2020-12-15 [1] RSPM (R 4.1.0)
## pkgconfig   2.0.3 2019-09-22 [1] RSPM (R 4.1.0)
## pkgload     1.2.1 2021-04-06 [1] RSPM (R 4.1.0)
## plotly      * 4.9.4.1 2021-06-18 [1] RSPM (R 4.0.5)
## plyr       * 1.8.6 2020-03-03 [1] RSPM (R 4.0.5)
## prettyunits 1.1.1 2020-01-24 [1] RSPM (R 4.1.0)
## processx    3.5.2 2021-04-30 [1] RSPM (R 4.1.0)
## ps          1.6.0 2021-02-28 [1] RSPM (R 4.1.0)
## purrr      0.3.4 2020-04-17 [1] RSPM (R 4.1.0)
## R6          2.5.0 2020-10-28 [1] RSPM (R 4.1.0)
## RColorBrewer * 1.1-2 2014-12-07 [1] RSPM (R 4.1.0)
## Rcpp        1.0.7 2021-07-07 [1] RSPM (R 4.1.0)
## readr       * 2.0.0 2021-07-20 [1] RSPM (R 4.0.5)
## readxl      * 1.3.1 2019-03-13 [1] RSPM (R 4.1.0)
## remotes     2.4.0 2021-06-02 [1] RSPM (R 4.1.0)
## reshape     0.8.8 2018-10-23 [1] RSPM (R 4.0.3)
## rio         0.5.27 2021-06-21 [1] RSPM (R 4.0.5)
## rlang       0.4.11 2021-04-30 [1] RSPM (R 4.1.0)
## rmarkdown   2.9   2021-06-15 [1] RSPM (R 4.1.0)
## ROCR       * 1.0-11 2020-05-02 [1] RSPM (R 4.0.0)
## rprojroot   2.0.2 2020-11-15 [1] RSPM (R 4.1.0)
## rstatix    0.7.0 2021-02-13 [1] RSPM (R 4.0.3)
## rstudioapi  0.13 2020-11-12 [1] RSPM (R 4.1.0)
## Rtsne       * 0.15 2018-11-10 [1] RSPM (R 4.0.5)
## rvest       1.0.1 2021-07-26 [1] RSPM (R 4.0.5)
## sass        0.4.0 2021-05-12 [1] RSPM (R 4.1.0)
## scales      1.1.1 2020-05-11 [1] RSPM (R 4.1.0)
## sessioninfo 1.1.1 2018-11-05 [1] RSPM (R 4.1.0)
## stringi     * 1.7.3 2021-07-16 [1] RSPM (R 4.0.5)
## stringr     * 1.4.0 2019-02-10 [1] RSPM (R 4.1.0)
## svglite     2.0.0 2021-02-20 [1] RSPM (R 4.0.5)
## systemfonts 1.0.2 2021-05-11 [1] RSPM (R 4.0.5)
## testthat    3.0.4 2021-07-01 [1] RSPM (R 4.1.0)
## tibble      * 3.1.3 2021-07-23 [1] RSPM (R 4.0.5)
## tidyverse    * 1.1.3 2021-03-03 [1] RSPM (R 4.1.0)
## tidyselect   1.1.1 2021-04-30 [1] RSPM (R 4.1.0)
## tzdb        0.1.2 2021-07-20 [1] RSPM (R 4.0.5)
## usethis     2.0.1 2021-02-10 [1] RSPM (R 4.1.0)
## utf8        1.2.2 2021-07-24 [1] RSPM (R 4.0.5)
## vctrs        0.3.8 2021-04-29 [1] RSPM (R 4.1.0)
## viridisLite 0.4.0 2021-04-13 [1] RSPM (R 4.1.0)
## webshot     0.5.2 2019-11-22 [1] RSPM (R 4.1.0)
```

```
## withr      2.4.2 2021-04-18 [1] RSPM (R 4.1.0)
## WriteXLS   * 6.3.0 2021-04-01 [1] RSPM (R 4.0.4)
## xfun       0.24  2021-06-15 [1] RSPM (R 4.1.0)
## xml2       1.3.2  2020-04-23 [1] RSPM (R 4.1.0)
## yaml       2.2.1  2020-02-01 [1] RSPM (R 4.1.0)
## zip        2.2.0  2021-05-31 [1] RSPM (R 4.1.0)
##
## [1] /usr/local/lib/R/site-library
## [2] /usr/local/lib/R/library
```

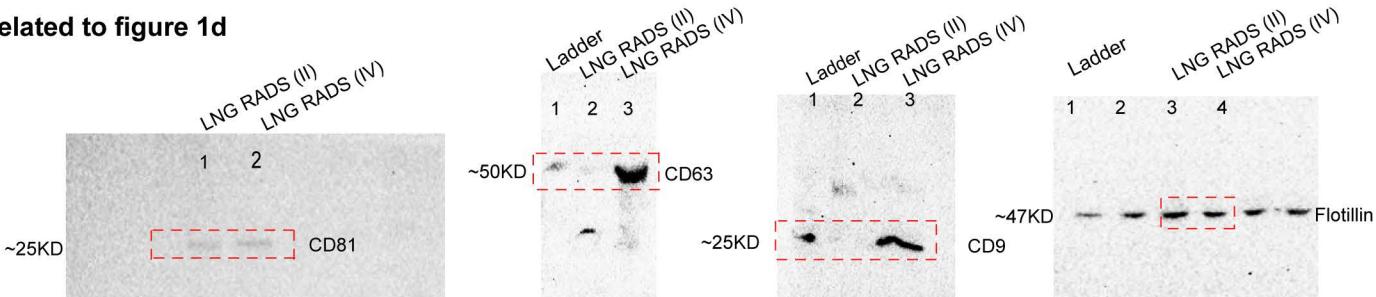
Combining Plasma Extracellular Vesicle Let-7b-5p, miR-184 and Circulating miR-22-3p Levels for NSCLC Diagnosis and Drug Resistance Prediction

G. P. Vadla¹, B. Daghpat¹, N. Patterson¹, V. Ahmad¹, G. Perez^{1*}, A. Garcia^{1*}, Y. Manjunath², J.T. Kaifi^{2,3}, G. Li^{2,3}, and C.Y. Chabu^{1,2,3‡}

Supplementary information 3. Full images of Western blots

Red dashlines denote the area presented in the main figures.

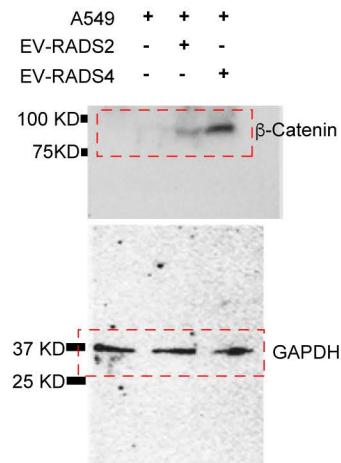
Related to figure 1d



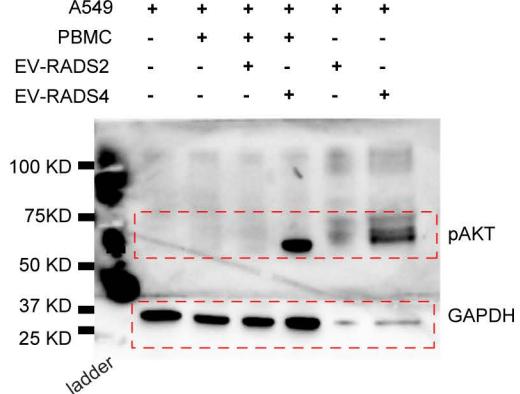
Related to figure 1e



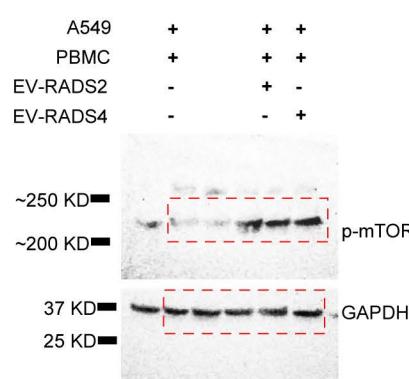
Related to figure 3e



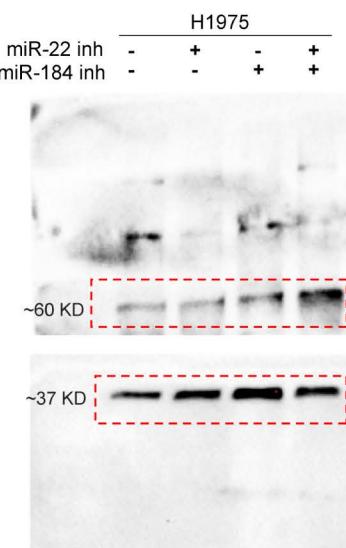
Related to figure 3f



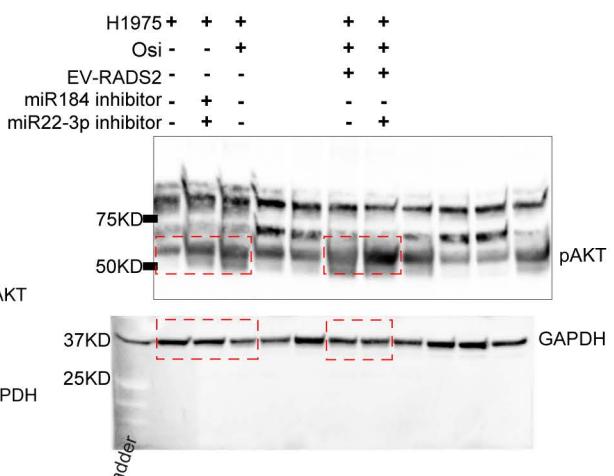
Related to figure 3g



Related to figure 3k



Related to figure 3l



Related to supplementary figure 3

