Figure S2

A

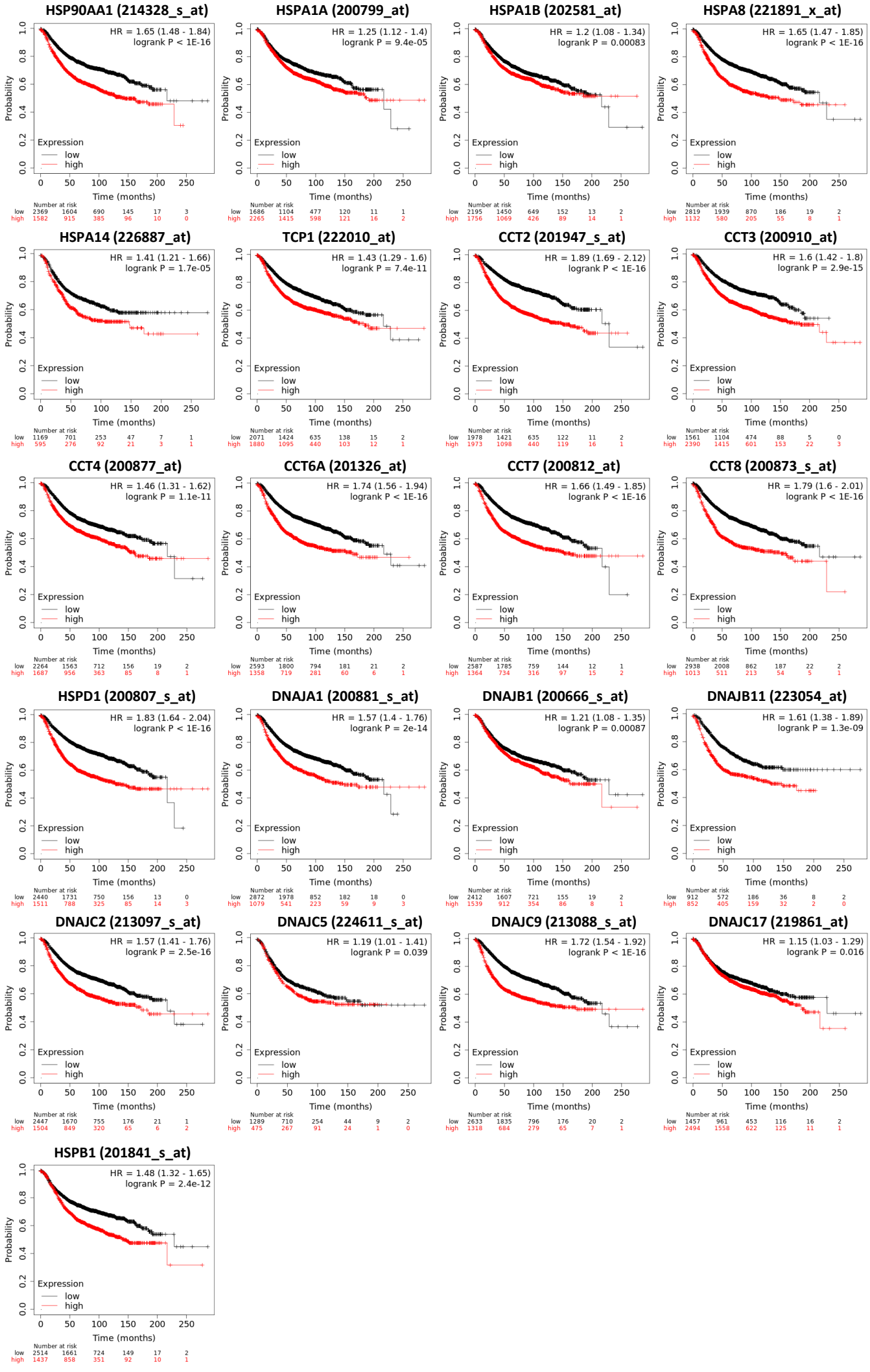

HSP90AA1 (214328_s_at)
HR = 1.65 (1.48 - 1.84)
logrank P < 1E-16
HSPA1A (200799_at)
HR = 1.25 (1.12 - 1.4)
logrank P = 9.4e-05
HSPA1B (202581_at)
HR = 1.2 (1.08 - 1.34)
logrank P = 0.00083
HSPA8 (221891_x_at)
HR = 1.65 (1.47 - 1.85)
logrank P < 1E-16
HSPA14 (226887_at)
HR = 1.41 (1.21 - 1.66)
logrank P = 1.7e-05
TCP1 (222010_at)
HR = 1.43 (1.29 - 1.6)
logrank P = 7.4e-11
CCT2 (201947_s_at)
HR = 1.89 (1.69 - 2.12)
logrank P < 1E-16
CCT3 (200910_at)
HR = 1.6 (1.42 - 1.8)
logrank P = 2.9e-15
CCT4 (200877_at)
HR = 1.46 (1.31 - 1.62)
logrank P = 1.1e-11
CCT6A (201326_at)
HR = 1.74 (1.56 - 1.94)
logrank P < 1E-16
CCT7 (200812_at)
HR = 1.66 (1.49 - 1.85)
logrank P < 1E-16
CCT8 (200873_s_at)
HR = 1.79 (1.6 - 2.01)
logrank P < 1E-16
HSPD1 (200807_s_at)
HR = 1.83 (1.64 - 2.04)
logrank P < 1E-16
DNAJA1 (200881_s_at)
HR = 1.57 (1.4 - 1.76)
logrank P = 2e-14
DNAJB1 (200666_s_at)
HR = 1.21 (1.08 - 1.35)
logrank P = 0.00087
DNAJB11 (223054_at)
HR = 1.61 (1.38 - 1.89)
logrank P = 1.3e-09
DNAJC2 (213097_s_at)
HR = 1.57 (1.41 - 1.76)
logrank P = 2.5e-16
DNAJC5 (224611_s_at)
HR = 1.19 (1.01 - 1.41)
logrank P = 0.039
DNAJC9 (213088_s_at)
HR = 1.72 (1.54 - 1.92)
logrank P < 1E-16
DNAJC17 (219861_at)
HR = 1.15 (1.03 - 1.29)
logrank P = 0.016
HSPB1 (201841_s_at)
HR = 1.48 (1.32 - 1.65)
logrank P = 2.4e-12
Probability
Expression
low
high
Time (months)
Number at risk

Figure S2

B

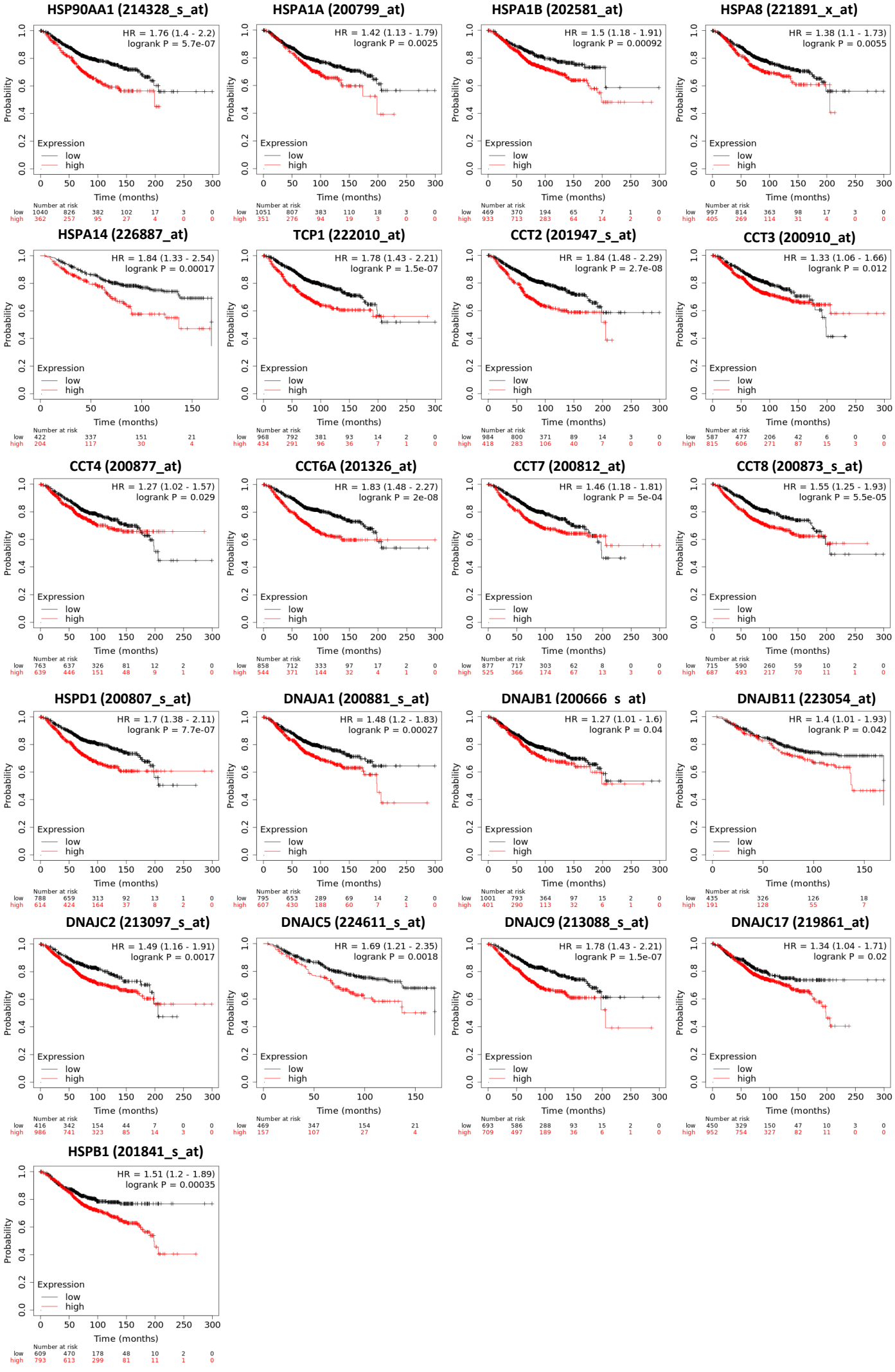

HSP90AA1 (214328_s_at)
HR = 1.76 (1.4 - 2.2)
logrank P = 5.7e-07
HSPA1A (200799_at)
HR = 1.42 (1.13 - 1.79)
logrank P = 0.0025
HSPA1B (202581_at)
HR = 1.5 (1.18 - 1.91)
logrank P = 0.00092
HSPA8 (221891_x_at)
HR = 1.38 (1.1 - 1.73)
logrank P = 0.0055
HSPA14 (226887_at)
HR = 1.84 (1.33 - 2.54)
logrank P = 0.00017
TCP1 (222010_at)
HR = 1.78 (1.43 - 2.21)
logrank P = 1.5e-07
CCT2 (201947_s_at)
HR = 1.84 (1.48 - 2.29)
logrank P = 2.7e-08
CCT3 (200910_at)
HR = 1.33 (1.06 - 1.66)
logrank P = 0.012
CCT4 (200877_at)
HR = 1.27 (1.02 - 1.57)
logrank P = 0.029
CCT6A (201326_at)
HR = 1.83 (1.48 - 2.27)
logrank P = 2e-08
CCT7 (200812_at)
HR = 1.46 (1.18 - 1.81)
logrank P = 5e-04
CCT8 (200873_s_at)
HR = 1.55 (1.25 - 1.93)
logrank P = 5.5e-05
HSPD1 (200807_s_at)
HR = 1.7 (1.38 - 2.11)
logrank P = 7.7e-07
DNAJA1 (200881_s_at)
HR = 1.48 (1.2 - 1.83)
logrank P = 0.00027
DNAJB1 (200666_s_at)
HR = 1.27 (1.01 - 1.6)
logrank P = 0.04
DNAJB11 (223054_at)
HR = 1.4 (1.01 - 1.93)
logrank P = 0.042
DNAJC2 (213097_s_at)
HR = 1.49 (1.16 - 1.91)
logrank P = 0.0017
DNAJC5 (224611_s_at)
HR = 1.69 (1.21 - 2.35)
logrank P = 0.0018
DNAJC9 (213088_s_at)
HR = 1.78 (1.43 - 2.21)
logrank P = 1.5e-07
DNAJC17 (219861_at)
HR = 1.34 (1.04 - 1.71)
logrank P = 0.02
HSPB1 (201841_s_at)
HR = 1.51 (1.2 - 1.89)
logrank P = 0.00035
Probability
Expression
low
high
Time (months)
Number at risk

Figure S2

C

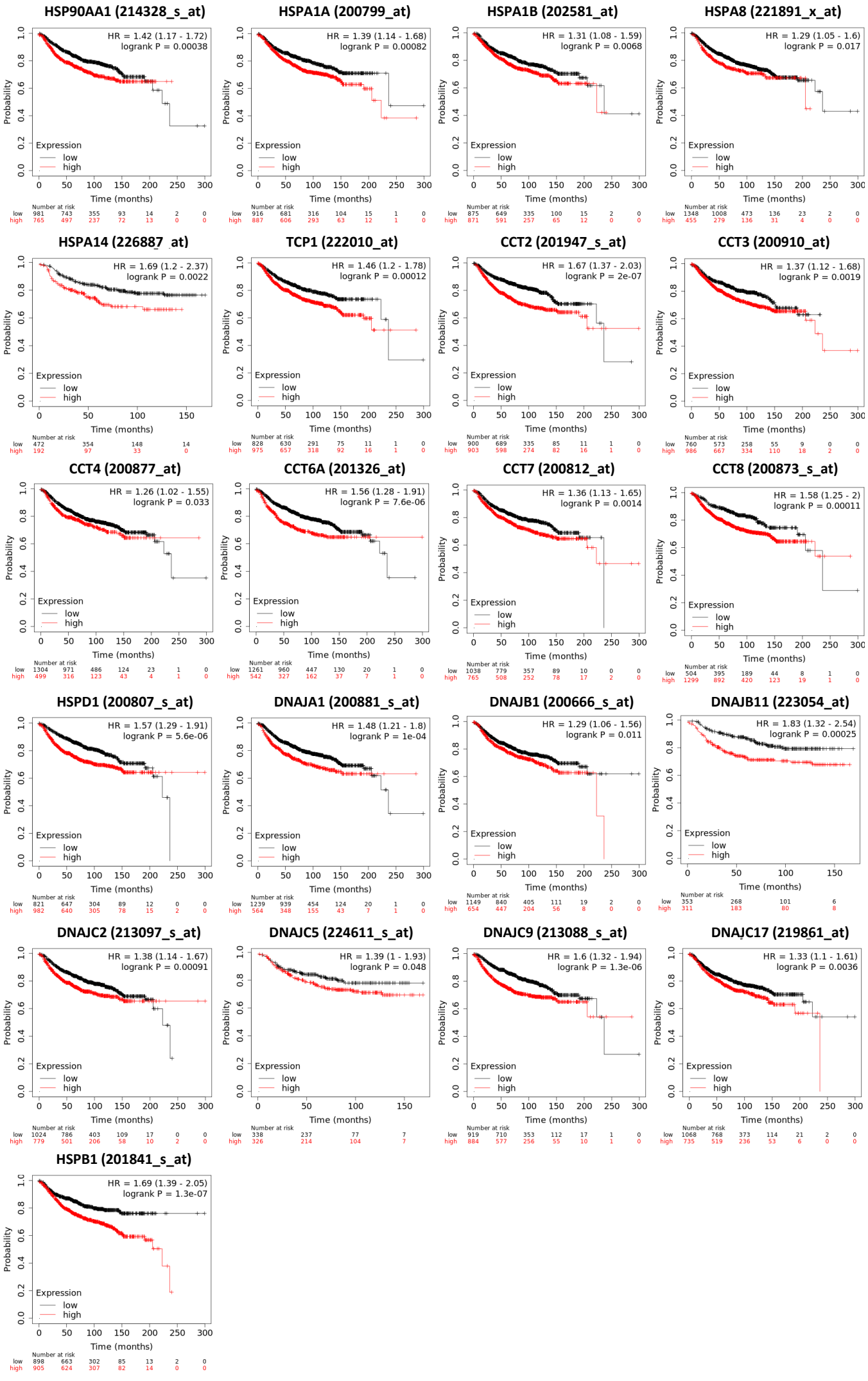

HSP90AA1 (214328_s_at)
HR = 1.42 (1.17 - 1.72)
logrank P = 0.00038
HSPA1A (200799_at)
HR = 1.39 (1.14 - 1.68)
logrank P = 0.00082
HSPA1B (202581_at)
HR = 1.31 (1.08 - 1.59)
logrank P = 0.0068
HSPA8 (221891_x_at)
HR = 1.29 (1.05 - 1.6)
logrank P = 0.017
HSPA14 (226887_at)
HR = 1.69 (1.2 - 2.37)
logrank P = 0.0022
TCP1 (222010_at)
HR = 1.46 (1.2 - 1.78)
logrank P = 0.00012
CCT2 (201947_s_at)
HR = 1.67 (1.37 - 2.03)
logrank P = 2e-07
CCT3 (200910_at)
HR = 1.37 (1.12 - 1.68)
logrank P = 0.0019
CCT4 (200877_at)
HR = 1.26 (1.02 - 1.55)
logrank P = 0.033
CCT6A (201326_at)
HR = 1.56 (1.28 - 1.91)
logrank P = 7.6e-06
CCT7 (200812_at)
HR = 1.36 (1.13 - 1.65)
logrank P = 0.0014
CCT8 (200873_s_at)
HR = 1.58 (1.25 - 2)
logrank P = 0.00011
HSPD1 (200807_s_at)
HR = 1.57 (1.29 - 1.91)
logrank P = 5.6e-06
DNAJA1 (200881_s_at)
HR = 1.48 (1.21 - 1.8)
logrank P = 1e-04
DNAJB1 (200666_s_at)
HR = 1.29 (1.06 - 1.56)
logrank P = 0.011
DNAJB11 (223054_at)
HR = 1.83 (1.32 - 2.54)
logrank P = 0.00025
DNAJC2 (213097_s_at)
HR = 1.38 (1.14 - 1.67)
logrank P = 0.00091
DNAJC5 (224611_s_at)
HR = 1.39 (1 - 1.93)
logrank P = 0.048
DNAJC9 (213088_s_at)
HR = 1.6 (1.32 - 1.94)
logrank P = 1.3e-06
DNAJC17 (219861_at)
HR = 1.33 (1.1 - 1.61)
logrank P = 0.0036
HSPB1 (201841_s_at)
HR = 1.69 (1.39 - 2.05)
logrank P = 1.3e-07
Probability
Expression
low
high
Time (months)
Number at risk

Figure S2

D

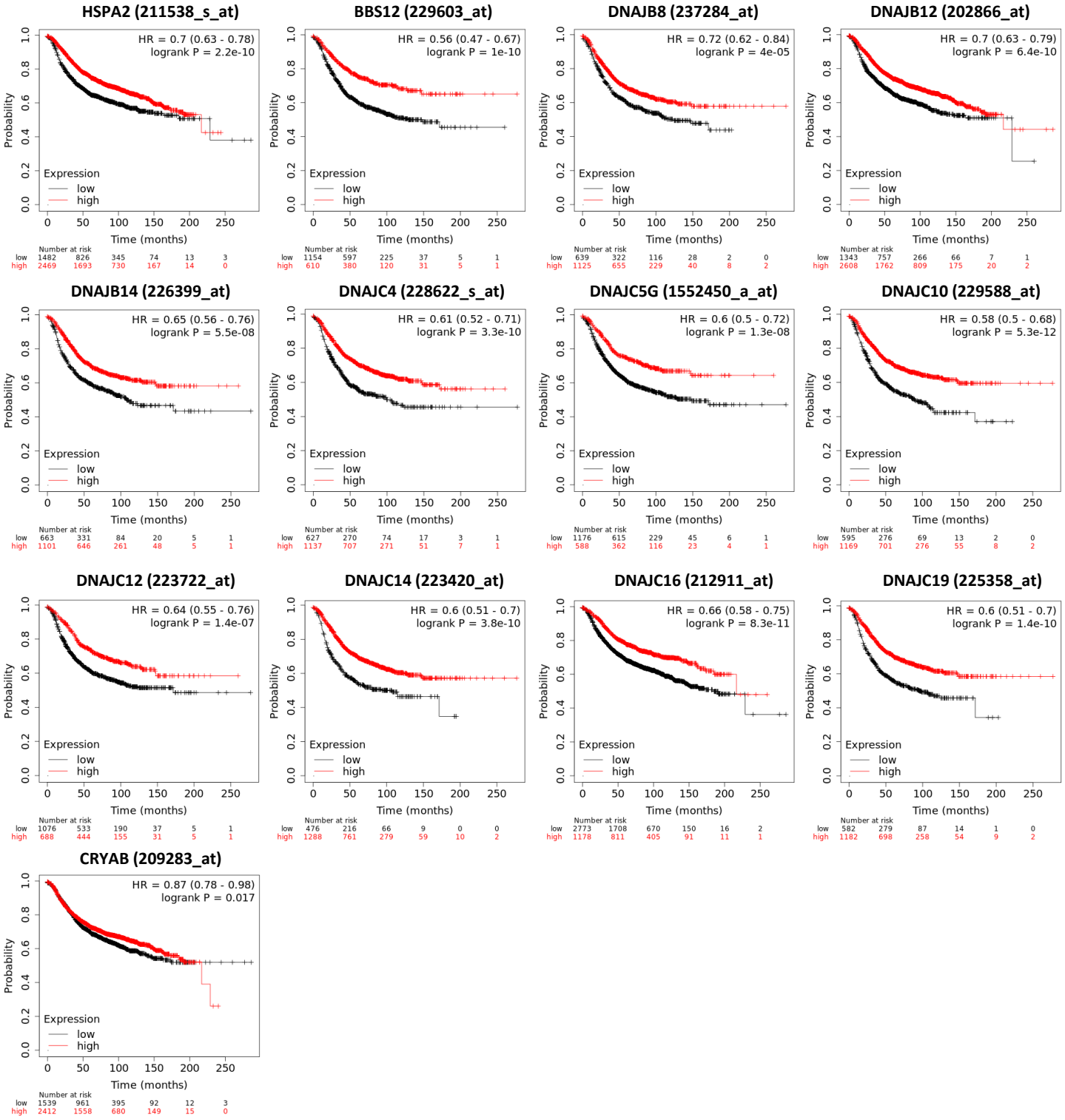

HSPA2 (211538_s_at)
HR = 0.7 (0.63 - 0.78)
logrank P = 2.2e-10
BBS12 (229603_at)
HR = 0.56 (0.47 - 0.67)
logrank P = 1e-10
DNAJB8 (237284_at)
HR = 0.72 (0.62 - 0.84)
logrank P = 4e-05
DNAJB12 (202866_at)
HR = 0.7 (0.63 - 0.79)
logrank P = 6.4e-10
DNAJB14 (226399_at)
HR = 0.65 (0.56 - 0.76)
logrank P = 5.5e-08
DNAJC4 (228622_s_at)
HR = 0.61 (0.52 - 0.71)
logrank P = 3.3e-10
DNAJC5G (1552450_a_at)
HR = 0.6 (0.5 - 0.72)
logrank P = 1.3e-08
DNAJC10 (229588_at)
HR = 0.58 (0.5 - 0.68)
logrank P = 5.3e-12
DNAJC12 (223722_at)
HR = 0.64 (0.55 - 0.76)
logrank P = 1.4e-07
DNAJC14 (223420_at)
HR = 0.6 (0.51 - 0.7)
logrank P = 3.8e-10
DNAJC16 (212911_at)
HR = 0.66 (0.58 - 0.75)
logrank P = 8.3e-11
DNAJC19 (225358_at)
HR = 0.6 (0.51 - 0.7)
logrank P = 1.4e-10
CRYAB (209283_at)
HR = 0.87 (0.78 - 0.98)
logrank P = 0.017
Probability
Time (months)
Expression
low
high
Number at risk

Figure S2

E

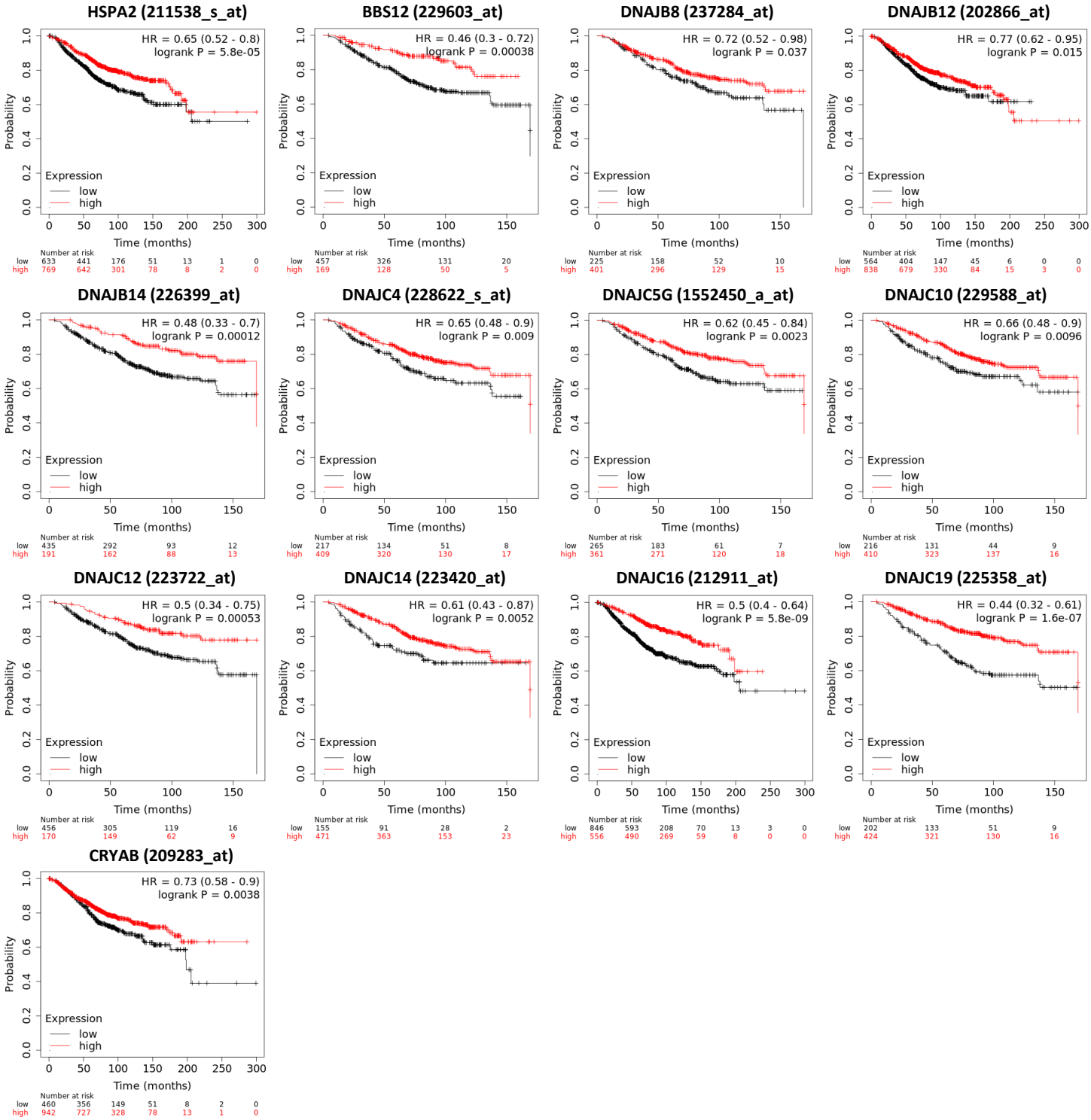
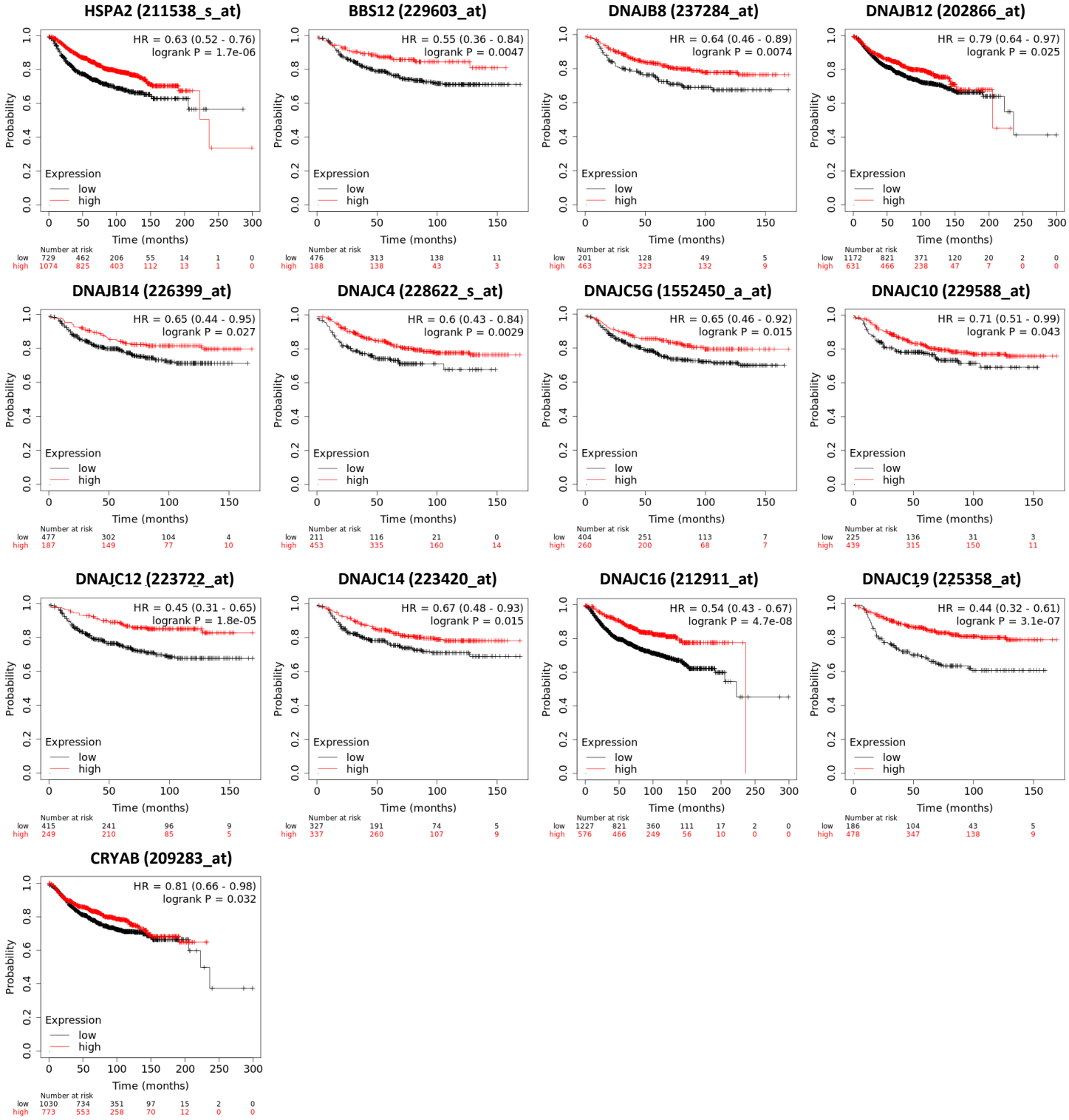

HSPA2 (211538_s_at)
HR = 0.65 (0.52 - 0.8)
logrank P = 5.8e-05
BBS12 (229603_at)
HR = 0.46 (0.3 - 0.72)
logrank P = 0.00038
DNAJB8 (237284_at)
HR = 0.72 (0.52 - 0.98)
logrank P = 0.037
DNAJB12 (202866_at)
HR = 0.77 (0.62 - 0.95)
logrank P = 0.015
DNAJB14 (226399_at)
HR = 0.48 (0.33 - 0.7)
logrank P = 0.00012
DNAJC4 (228622_s_at)
HR = 0.65 (0.48 - 0.9)
logrank P = 0.009
DNAJC5G (1552450_a_at)
HR = 0.62 (0.45 - 0.84)
logrank P = 0.0023
DNAJC10 (229588_at)
HR = 0.66 (0.48 - 0.9)
logrank P = 0.0096
DNAJC12 (223722_at)
HR = 0.5 (0.34 - 0.75)
logrank P = 0.00053
DNAJC14 (223420_at)
HR = 0.61 (0.43 - 0.87)
logrank P = 0.0052
DNAJC16 (212911_at)
HR = 0.5 (0.4 - 0.64)
logrank P = 5.8e-09
DNAJC19 (225358_at)
HR = 0.44 (0.32 - 0.61)
logrank P = 1.6e-07
CRYAB (209283_at)
HR = 0.73 (0.58 - 0.9)
logrank P = 0.0038
Probability
Expression
low
high
Time (months)
Number at risk

Figure S2

F


HSPA2 (211538_s_at)
HR = 0.63 (0.52 - 0.76)
logrank P = 1.7e-06
BBS12 (229603_at)
HR = 0.55 (0.36 - 0.84)
logrank P = 0.0047
DNAJB8 (237284_at)
HR = 0.64 (0.46 - 0.89)
logrank P = 0.0074
DNAJB12 (202866_at)
HR = 0.79 (0.64 - 0.97)
logrank P = 0.025
DNAJB14 (226399_at)
HR = 0.65 (0.44 - 0.95)
logrank P = 0.027
DNAJC4 (228622_s_at)
HR = 0.6 (0.43 - 0.84)
logrank P = 0.0029
DNAJC5G (1552450_a_at)
HR = 0.65 (0.46 - 0.92)
logrank P = 0.015
DNAJC10 (229588_at)
HR = 0.71 (0.51 - 0.99)
logrank P = 0.043
DNAJC12 (223722_at)
HR = 0.45 (0.31 - 0.65)
logrank P = 1.8e-05
DNAJC14 (223420_at)
HR = 0.67 (0.48 - 0.93)
logrank P = 0.015
DNAJC16 (212911_at)
HR = 0.54 (0.43 - 0.67)
logrank P = 4.7e-08
DNAJC19 (225358_at)
HR = 0.44 (0.32 - 0.61)
logrank P = 3.1e-07
CRYAB (209283_at)
HR = 0.81 (0.66 - 0.98)
logrank P = 0.032
Probability
Expression
low
high
Time (months)
Number at risk