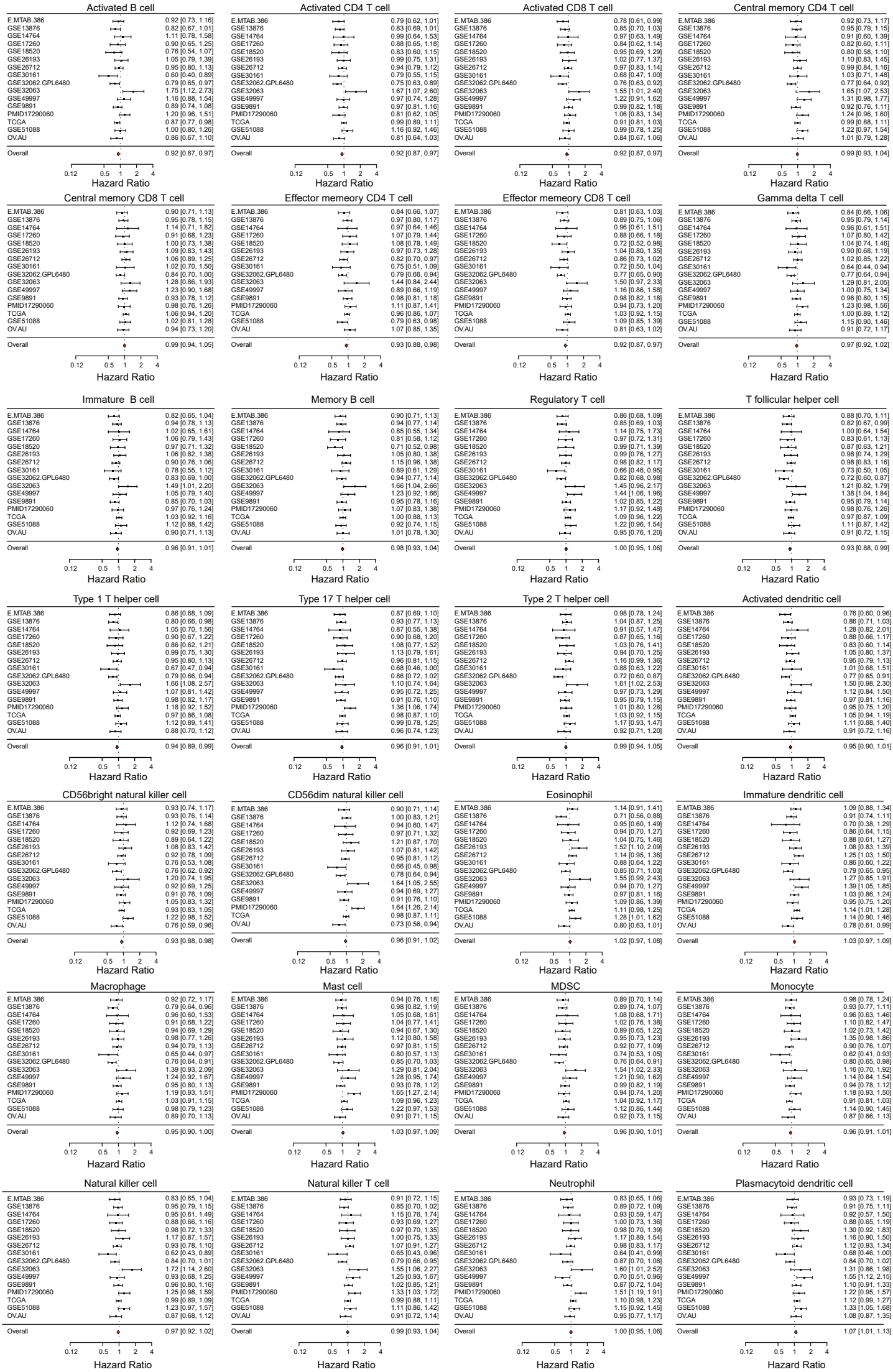
Activated B cell
E.MTAB.386 0.92 [0.73, 1.16]
GSE13876 0.82 [0.67, 1.01]
GSE14764 1.11 [0.78, 1.58]
GSE17260 0.90 [0.65, 1.25]
GSE18520 0.76 [0.54, 1.07]
GSE26193 1.05 [0.79, 1.39]
GSE26712 0.95 [0.80, 1.13]
GSE30161 0.60 [0.40, 0.89]
GSE32062.GPL6480 0.79 [0.65, 0.97]
GSE32063 1.75 [1.12, 2.73]
GSE49997 1.16 [0.88, 1.54]
GSE9891 0.89 [0.74, 1.08]
PMID17290060 1.20 [0.96, 1.51]
TCGA 0.87 [0.77, 0.98]
GSE51088 1.00 [0.80, 1.26]
OV.AU 0.86 [0.67, 1.10]
Overall 0.92 [0.87, 0.97]
0.12 0.5 1 2 4
Hazard Ratio
Activated CD4 T cell
E.MTAB.386 0.79 [0.62, 1.01]
GSE13876 0.83 [0.69, 1.01]
GSE14764 0.99 [0.64, 1.53]
GSE17260 0.88 [0.65, 1.18]
GSE18520 0.83 [0.60, 1.15]
GSE26193 0.99 [0.75, 1.31]
GSE26712 0.94 [0.79, 1.12]
GSE30161 0.79 [0.55, 1.15]
GSE32062.GPL6480 0.75 [0.63, 0.89]
GSE32063 1.67 [1.07, 2.60]
GSE49997 0.97 [0.74, 1.28]
GSE9891 0.97 [0.81, 1.16]
PMID17290060 0.81 [0.62, 1.05]
TCGA 0.99 [0.89, 1.11]
GSE51088 1.16 [0.92, 1.46]
OV.AU 0.81 [0.64, 1.03]
Overall 0.92 [0.87, 0.97]
0.12 0.5 1 2 4
Hazard Ratio
Activated CD8 T cell
E.MTAB.386 0.78 [0.61, 0.99]
GSE13876 0.85 [0.70, 1.03]
GSE14764 0.97 [0.63, 1.49]
GSE17260 0.84 [0.62, 1.14]
GSE18520 0.95 [0.69, 1.29]
GSE26193 1.02 [0.77, 1.37]
GSE26712 0.97 [0.83, 1.14]
GSE30161 0.68 [0.47, 1.00]
GSE32062.GPL6480 0.76 [0.63, 0.92]
GSE32063 1.55 [1.01, 2.40]
GSE49997 1.22 [0.91, 1.62]
GSE9891 0.99 [0.82, 1.18]
PMID17290060 1.06 [0.83, 1.34]
TCGA 0.91 [0.81, 1.03]
GSE51088 0.99 [0.78, 1.25]
OV.AU 0.84 [0.67, 1.06]
Overall 0.92 [0.87, 0.97]
0.12 0.5 1 2 4
Hazard Ratio
Central memory CD4 T cell
E.MTAB.386 0.92 [0.73, 1.17]
GSE13876 0.95 [0.79, 1.15]
GSE14764 0.91 [0.60, 1.39]
GSE17260 0.82 [0.60, 1.11]
GSE18520 0.80 [0.58, 1.10]
GSE26193 1.10 [0.83, 1.45]
GSE26712 0.99 [0.84, 1.16]
GSE30161 1.03 [0.71, 1.48]
GSE32062.GPL6480 0.77 [0.64, 0.92]
GSE32063 1.65 [1.07, 2.53]
GSE49997 1.31 [0.98, 1.77]
GSE9891 0.92 [0.76, 1.11]
PMID17290060 1.24 [0.96, 1.60]
TCGA 0.99 [0.88, 1.11]
GSE51088 1.22 [0.97, 1.54]
OV.AU 1.01 [0.79, 1.28]
Overall 0.99 [0.93, 1.04]
0.12 0.5 1 2 4
Hazard Ratio
Central memory CD8 T cell
E.MTAB.386 0.90 [0.71, 1.13]
GSE13876 0.95 [0.78, 1.15]
GSE14764 1.14 [0.71, 1.82]
GSE17260 0.91 [0.68, 1.23]
GSE18520 1.00 [0.73, 1.38]
GSE26193 1.09 [0.83, 1.43]
GSE26712 1.06 [0.89, 1.25]
GSE30161 1.02 [0.70, 1.50]
GSE32062.GPL6480 0.84 [0.70, 1.00]
GSE32063 1.28 [0.86, 1.93]
GSE49997 1.23 [0.90, 1.68]
GSE9891 0.93 [0.78, 1.12]
PMID17290060 0.98 [0.76, 1.26]
TCGA 1.06 [0.94, 1.20]
GSE51088 1.02 [0.81, 1.28]
OV.AU 0.94 [0.73, 1.20]
Overall 0.99 [0.94, 1.05]
0.12 0.5 1 2 4
Hazard Ratio
Effector memeory CD4 T cell
E.MTAB.386 0.84 [0.66, 1.07]
GSE13876 0.97 [0.80, 1.17]
GSE14764 0.97 [0.64, 1.46]
GSE17260 1.07 [0.79, 1.44]
GSE18520 1.08 [0.78, 1.49]
GSE26193 0.97 [0.73, 1.28]
GSE26712 0.82 [0.70, 0.97]
GSE30161 0.75 [0.51, 1.09]
GSE32062.GPL6480 0.79 [0.66, 0.94]
GSE32063 1.44 [0.84, 2.44]
GSE49997 0.89 [0.66, 1.19]
GSE9891 0.98 [0.81, 1.18]
PMID17290060 1.11 [0.87, 1.41]
TCGA 0.96 [0.86, 1.07]
GSE51088 0.79 [0.63, 0.98]
OV.AU 1.07 [0.85, 1.35]
Overall 0.93 [0.88, 0.98]
0.12 0.5 1 2 4
Hazard Ratio
Effector memeory CD8 T cell
E.MTAB.386 0.81 [0.63, 1.03]
GSE13876 0.89 [0.75, 1.06]
GSE14764 0.96 [0.61, 1.51]
GSE17260 0.88 [0.66, 1.18]
GSE18520 0.72 [0.52, 0.98]
GSE26193 1.04 [0.80, 1.35]
GSE26712 0.86 [0.73, 1.02]
GSE30161 0.72 [0.50, 1.04]
GSE32062.GPL6480 0.77 [0.65, 0.90]
GSE32063 1.50 [0.97, 2.33]
GSE49997 1.16 [0.86, 1.58]
GSE9891 0.98 [0.82, 1.18]
PMID17290060 0.94 [0.73, 1.20]
TCGA 1.03 [0.92, 1.15]
GSE51088 1.09 [0.85, 1.39]
OV.AU 0.81 [0.63, 1.02]
Overall 0.92 [0.87, 0.97]
0.12 0.5 1 2 4
Hazard Ratio
Gamma delta T cell
E.MTAB.386 0.84 [0.66, 1.06]
GSE13876 0.95 [0.79, 1.14]
GSE14764 0.96 [0.61, 1.51]
GSE17260 1.07 [0.80, 1.42]
GSE18520 1.04 [0.74, 1.46]
GSE26193 0.90 [0.68, 1.19]
GSE26712 1.02 [0.85, 1.22]
GSE30161 0.64 [0.44, 0.94]
GSE32062.GPL6480 0.77 [0.64, 0.94]
GSE32063 1.29 [0.81, 2.05]
GSE49997 1.00 [0.75, 1.34]
GSE9891 0.96 [0.80, 1.15]
PMID17290060 1.23 [0.98, 1.56]
TCGA 1.00 [0.89, 1.12]
GSE51088 1.15 [0.90, 1.46]
OV.AU 0.91 [0.72, 1.17]
Overall 0.97 [0.92, 1.02]
0.12 0.5 1 2 4
Hazard Ratio
Immature B cell
E.MTAB.386 0.82 [0.65, 1.04]
GSE13876 0.94 [0.78, 1.13]
GSE14764 1.02 [0.65, 1.61]
GSE17260 1.06 [0.79, 1.43]
GSE18520 0.97 [0.71, 1.32]
GSE26193 1.06 [0.82, 1.38]
GSE26712 0.90 [0.76, 1.06]
GSE30161 0.78 [0.55, 1.12]
GSE32062.GPL6480 0.83 [0.69, 1.00]
GSE32063 1.49 [1.01, 2.20]
GSE49997 1.05 [0.79, 1.40]
GSE9891 0.85 [0.70, 1.03]
PMID17290060 0.97 [0.76, 1.24]
TCGA 1.03 [0.92, 1.16]
GSE51088 1.12 [0.88, 1.42]
OV.AU 0.90 [0.71, 1.13]
Overall 0.96 [0.91, 1.01]
0.12 0.5 1 2 4
Hazard Ratio
Memory B cell
E.MTAB.386 0.90 [0.71, 1.13]
GSE13876 0.94 [0.77, 1.14]
GSE14764 0.85 [0.55, 1.34]
GSE17260 0.81 [0.58, 1.12]
GSE18520 0.71 [0.52, 0.98]
GSE26193 1.05 [0.80, 1.38]
GSE26712 1.15 [0.96, 1.38]
GSE30161 0.89 [0.61, 1.29]
GSE32062.GPL6480 0.94 [0.77, 1.14]
GSE32063 1.66 [1.04, 2.66]
GSE49997 1.23 [0.92, 1.66]
GSE9891 0.95 [0.78, 1.16]
PMID17290060 1.07 [0.83, 1.38]
TCGA 1.00 [0.88, 1.13]
GSE51088 0.92 [0.74, 1.15]
OV.AU 1.01 [0.78, 1.30]
Overall 0.98 [0.93, 1.04]
0.12 0.5 1 2 4
Hazard Ratio
Regulatory T cell
E.MTAB.386 0.86 [0.68, 1.09]
GSE13876 0.85 [0.69, 1.03]
GSE14764 1.14 [0.75, 1.73]
GSE17260 0.97 [0.72, 1.31]
GSE18520 0.99 [0.71, 1.39]
GSE26193 0.99 [0.76, 1.27]
GSE26712 0.98 [0.82, 1.17]
GSE30161 0.66 [0.46, 0.95]
GSE32062.GPL6480 0.82 [0.68, 0.98]
GSE32063 1.45 [0.96, 2.17]
GSE49997 1.44 [1.06, 1.96]
GSE9891 1.02 [0.85, 1.22]
PMID17290060 1.17 [0.92, 1.48]
TCGA 1.09 [0.96, 1.22]
GSE51088 1.22 [0.96, 1.54]
OV.AU 0.95 [0.76, 1.20]
Overall 1.00 [0.95, 1.06]
0.12 0.5 1 2 4
Hazard Ratio
T follicular helper cell
E.MTAB.386 0.88 [0.70, 1.11]
GSE13876 0.82 [0.67, 0.99]
GSE14764 1.00 [0.64, 1.54]
GSE17260 0.83 [0.61, 1.13]
GSE18520 0.87 [0.63, 1.21]
GSE26193 0.98 [0.74, 1.29]
GSE26712 0.98 [0.83, 1.16]
GSE30161 0.73 [0.50, 1.05]
GSE32062.GPL6480 0.72 [0.60, 0.87]
GSE32063 1.21 [0.82, 1.79]
GSE49997 1.38 [1.04, 1.84]
GSE9891 0.95 [0.79, 1.14]
PMID17290060 0.98 [0.76, 1.26]
TCGA 0.97 [0.87, 1.09]
GSE51088 1.11 [0.87, 1.42]
OV.AU 0.91 [0.72, 1.15]
Overall 0.93 [0.88, 0.99]
0.12 0.5 1 2 4
Hazard Ratio
Type 1 T helper cell
E.MTAB.386 0.86 [0.68, 1.09]
GSE13876 0.80 [0.66, 0.98]
GSE14764 1.05 [0.70, 1.56]
GSE17260 0.90 [0.67, 1.22]
GSE18520 0.86 [0.62, 1.21]
GSE26193 0.99 [0.75, 1.30]
GSE26712 0.95 [0.80, 1.13]
GSE30161 0.67 [0.47, 0.94]
GSE32062.GPL6480 0.79 [0.66, 0.94]
GSE32063 1.66 [1.08, 2.57]
GSE49997 1.07 [0.81, 1.42]
GSE9891 0.98 [0.82, 1.17]
PMID17290060 1.18 [0.92, 1.52]
TCGA 0.97 [0.86, 1.08]
GSE51088 1.12 [0.89, 1.41]
OV.AU 0.88 [0.70, 1.12]
Overall 0.94 [0.89, 0.99]
0.12 0.5 1 2 4
Hazard Ratio
Type 17 T helper cell
E.MTAB.386 0.87 [0.69, 1.10]
GSE13876 0.93 [0.77, 1.13]
GSE14764 0.87 [0.55, 1.38]
GSE17260 0.90 [0.68, 1.20]
GSE18520 1.08 [0.77, 1.52]
GSE26193 1.13 [0.79, 1.61]
GSE26712 0.96 [0.81, 1.15]
GSE30161 0.68 [0.46, 1.00]
GSE32062.GPL6480 0.86 [0.72, 1.02]
GSE32063 1.10 [0.74, 1.64]
GSE49997 0.95 [0.72, 1.25]
GSE9891 0.91 [0.76, 1.10]
PMID17290060 1.36 [1.06, 1.74]
TCGA 0.98 [0.87, 1.10]
GSE51088 0.99 [0.78, 1.25]
OV.AU 0.96 [0.74, 1.23]
Overall 0.96 [0.91, 1.01]
0.12 0.5 1 2 4
Hazard Ratio
Type 2 T helper cell
E.MTAB.386 0.98 [0.78, 1.24]
GSE13876 1.04 [0.87, 1.25]
GSE14764 0.91 [0.57, 1.47]
GSE17260 0.87 [0.65, 1.16]
GSE18520 1.03 [0.76, 1.41]
GSE26193 0.94 [0.70, 1.25]
GSE26712 1.16 [0.99, 1.36]
GSE30161 0.88 [0.63, 1.22]
GSE32062.GPL6480 0.72 [0.60, 0.87]
GSE32063 1.61 [1.02, 2.53]
GSE49997 0.97 [0.73, 1.29]
GSE9891 0.95 [0.79, 1.15]
PMID17290060 1.01 [0.80, 1.28]
TCGA 1.03 [0.92, 1.15]
GSE51088 1.17 [0.93, 1.47]
OV.AU 0.92 [0.71, 1.20]
Overall 0.99 [0.94, 1.05]
0.12 0.5 1 2 4
Hazard Ratio
Activated dendritic cell
E.MTAB.386 0.76 [0.60, 0.96]
GSE13876 0.86 [0.71, 1.03]
GSE14764 1.28 [0.82, 2.01]
GSE17260 0.88 [0.66, 1.17]
GSE18520 0.83 [0.60, 1.14]
GSE26193 1.05 [0.80, 1.37]
GSE26712 0.95 [0.79, 1.13]
GSE30161 1.01 [0.68, 1.51]
GSE32062.GPL6480 0.77 [0.65, 0.91]
GSE32063 1.50 [0.98, 2.30]
GSE49997 1.12 [0.84, 1.50]
GSE9891 0.97 [0.81, 1.16]
PMID17290060 0.95 [0.75, 1.20]
TCGA 1.05 [0.94, 1.19]
GSE51088 1.11 [0.88, 1.40]
OV.AU 0.91 [0.72, 1.16]
Overall 0.95 [0.90, 1.01]
0.12 0.5 1 2 4
Hazard Ratio
CD56bright natural killer cell
E.MTAB.386 0.93 [0.74, 1.17]
GSE13876 0.93 [0.76, 1.14]
GSE14764 1.12 [0.74, 1.68]
GSE17260 0.92 [0.69, 1.23]
GSE18520 0.89 [0.64, 1.22]
GSE26193 1.08 [0.83, 1.42]
GSE26712 0.92 [0.78, 1.09]
GSE30161 0.76 [0.53, 1.08]
GSE32062.GPL6480 0.76 [0.62, 0.92]
GSE32063 1.20 [0.74, 1.95]
GSE49997 0.92 [0.69, 1.25]
GSE9891 0.91 [0.76, 1.09]
PMID17290060 1.05 [0.83, 1.32]
TCGA 0.93 [0.83, 1.05]
GSE51088 1.22 [0.98, 1.52]
OV.AU 0.76 [0.59, 0.96]
Overall 0.93 [0.88, 0.98]
0.12 0.5 1 2 4
Hazard Ratio
CD56dim natural killer cell
E.MTAB.386 0.90 [0.71, 1.14]
GSE13876 1.00 [0.83, 1.21]
GSE14764 0.94 [0.60, 1.47]
GSE17260 0.97 [0.71, 1.32]
GSE18520 1.21 [0.87, 1.70]
GSE26193 1.07 [0.81, 1.42]
GSE26712 0.95 [0.81, 1.12]
GSE30161 0.66 [0.45, 0.98]
GSE32062.GPL6480 0.78 [0.64, 0.94]
GSE32063 1.64 [1.05, 2.55]
GSE49997 0.94 [0.69, 1.27]
GSE9891 0.91 [0.76, 1.10]
PMID17290060 1.64 [1.26, 2.14]
TCGA 0.98 [0.87, 1.11]
OV.AU 0.73 [0.56, 0.94]
Overall 0.96 [0.91, 1.02]
0.12 0.5 1 2 4
Hazard Ratio
Eosinophil
E.MTAB.386 1.14 [0.91, 1.41]
GSE13876 0.71 [0.56, 0.88]
GSE14764 0.95 [0.60, 1.49]
GSE17260 0.94 [0.70, 1.27]
GSE18520 1.04 [0.75, 1.46]
GSE26193 1.52 [1.10, 2.09]
GSE26712 1.14 [0.95, 1.36]
GSE30161 0.88 [0.64, 1.22]
GSE32062.GPL6480 0.85 [0.71, 1.03]
GSE32063 1.55 [0.99, 2.43]
GSE49997 0.94 [0.70, 1.27]
GSE9891 0.97 [0.81, 1.16]
PMID17290060 1.09 [0.86, 1.39]
TCGA 1.11 [0.98, 1.25]
GSE51088 1.28 [1.01, 1.62]
OV.AU 0.80 [0.63, 1.01]
Overall 1.02 [0.97, 1.08]
0.12 0.5 1 2 4
Hazard Ratio
Immature dendritic cell
E.MTAB.386 1.09 [0.88, 1.34]
GSE13876 0.91 [0.74, 1.11]
GSE14764 0.70 [0.38, 1.29]
GSE17260 0.86 [0.64, 1.15]
GSE18520 0.88 [0.61, 1.27]
GSE26193 1.08 [0.83, 1.39]
GSE26712 1.25 [1.03, 1.50]
GSE30161 0.86 [0.60, 1.22]
GSE32062.GPL6480 0.79 [0.65, 0.95]
GSE32063 1.27 [0.85, 1.91]
GSE49997 1.39 [1.05, 1.85]
GSE9891 1.03 [0.86, 1.24]
PMID17290060 0.95 [0.75, 1.20]
TCGA 1.14 [1.01, 1.28]
GSE51088 1.14 [0.90, 1.46]
OV.AU 0.78 [0.61, 0.99]
Overall 1.03 [0.97, 1.09]
0.12 0.5 1 2 4
Hazard Ratio
Macrophage
E.MTAB.386 0.92 [0.72, 1.17]
GSE13876 0.79 [0.64, 0.96]
GSE14764 0.96 [0.60, 1.53]
GSE17260 0.91 [0.68, 1.22]
GSE18520 0.94 [0.69, 1.29]
GSE26193 0.98 [0.77, 1.26]
GSE26712 0.94 [0.79, 1.13]
GSE30161 0.65 [0.44, 0.97]
GSE32062.GPL6480 0.76 [0.64, 0.91]
GSE32063 1.39 [0.93, 2.09]
GSE49997 1.24 [0.92, 1.67]
GSE9891 0.95 [0.80, 1.13]
PMID17290060 1.19 [0.93, 1.51]
TCGA 1.03 [0.91, 1.15]
GSE51088 0.98 [0.79, 1.23]
OV.AU 0.89 [0.70, 1.13]
Overall 0.95 [0.90, 1.00]
0.12 0.5 1 2 4
Hazard Ratio
Mast cell
E.MTAB.386 0.94 [0.76, 1.18]
GSE13876 0.98 [0.82, 1.19]
GSE14764 1.05 [0.68, 1.61]
GSE17260 1.04 [0.77, 1.41]
GSE18520 0.94 [0.67, 1.30]
GSE26193 1.12 [0.80, 1.58]
GSE26712 0.97 [0.81, 1.15]
GSE30161 0.80 [0.57, 1.13]
GSE32062.GPL6480 0.85 [0.70, 1.03]
GSE32063 1.29 [0.81, 2.04]
GSE49997 1.28 [0.95, 1.74]
GSE9891 0.93 [0.78, 1.12]
PMID17290060 1.65 [1.27, 2.14]
TCGA 1.09 [0.96, 1.23]
GSE51088 1.22 [0.97, 1.53]
OV.AU 0.91 [0.71, 1.15]
Overall 1.03 [0.97, 1.09]
0.12 0.5 1 2 4
Hazard Ratio
MDSC
E.MTAB.386 0.89 [0.70, 1.14]
GSE13876 0.89 [0.74, 1.07]
GSE14764 1.08 [0.68, 1.71]
GSE17260 1.02 [0.76, 1.38]
GSE18520 0.89 [0.65, 1.22]
GSE26193 0.95 [0.73, 1.23]
GSE26712 0.92 [0.77, 1.09]
GSE30161 0.74 [0.53, 1.05]
GSE32062.GPL6480 0.76 [0.64, 0.91]
GSE32063 1.54 [1.02, 2.33]
GSE49997 1.21 [0.90, 1.62]
GSE9891 0.99 [0.82, 1.19]
PMID17290060 0.94 [0.74, 1.20]
TCGA 1.04 [0.92, 1.17]
GSE51088 1.12 [0.86, 1.44]
OV.AU 0.92 [0.73, 1.15]
Overall 0.96 [0.90, 1.01]
0.12 0.5 1 2 4
Hazard Ratio
Monocyte
E.MTAB.386 0.98 [0.78, 1.24]
GSE13876 0.93 [0.77, 1.11]
GSE14764 0.96 [0.63, 1.46]
GSE17260 1.10 [0.82, 1.47]
GSE18520 1.02 [0.73, 1.42]
GSE26193 1.35 [0.98, 1.86]
GSE26712 0.90 [0.76, 1.07]
GSE30161 0.62 [0.41, 0.93]
GSE32062.GPL6480 0.80 [0.65, 0.98]
GSE32063 1.16 [0.70, 1.92]
GSE49997 1.14 [0.84, 1.54]
GSE9891 0.94 [0.78, 1.12]
PMID17290060 1.18 [0.93, 1.50]
TCGA 0.91 [0.81, 1.03]
GSE51088 1.14 [0.90, 1.45]
OV.AU 0.87 [0.66, 1.13]
Overall 0.96 [0.91, 1.01]
0.12 0.5 1 2 4
Hazard Ratio
Natural killer cell
E.MTAB.386 0.83 [0.65, 1.04]
GSE13876 0.95 [0.79, 1.15]
GSE14764 0.95 [0.61, 1.49]
GSE17260 0.88 [0.66, 1.16]
GSE18520 0.98 [0.72, 1.33]
GSE26193 1.17 [0.87, 1.57]
GSE26712 0.93 [0.78, 1.10]
GSE30161 0.62 [0.43, 0.89]
GSE32062.GPL6480 0.84 [0.70, 1.01]
GSE32063 1.72 [1.14, 2.60]
GSE49997 0.93 [0.68, 1.25]
GSE9891 0.96 [0.80, 1.16]
PMID17290060 1.25 [0.98, 1.59]
TCGA 0.99 [0.89, 1.09]
GSE51088 1.23 [0.97, 1.57]
OV.AU 0.87 [0.68, 1.12]
Overall 0.97 [0.92, 1.02]
0.12 0.5 1 2 4
Hazard Ratio
Natural killer T cell
E.MTAB.386 0.91 [0.72, 1.15]
GSE13876 0.85 [0.70, 1.02]
GSE14764 1.15 [0.76, 1.74]
GSE17260 0.93 [0.69, 1.27]
GSE18520 0.97 [0.70, 1.35]
GSE26193 1.00 [0.75, 1.33]
GSE26712 1.07 [0.91, 1.27]
GSE30161 0.65 [0.43, 0.96]
GSE32062.GPL6480 0.79 [0.66, 0.95]
GSE32063 1.55 [1.06, 2.27]
GSE49997 1.25 [0.93, 1.67]
GSE9891 1.02 [0.85, 1.21]
PMID17290060 1.33 [1.03, 1.72]
TCGA 0.99 [0.88, 1.11]
GSE51088 1.11 [0.86, 1.42]
OV.AU 0.91 [0.72, 1.14]
Overall 0.99 [0.93, 1.04]
0.12 0.5 1 2 4
Hazard Ratio
Neutrophil
E.MTAB.386 0.83 [0.65, 1.06]
GSE13876 0.89 [0.72, 1.09]
GSE14764 0.93 [0.59, 1.47]
GSE17260 1.00 [0.73, 1.36]
GSE18520 0.98 [0.70, 1.39]
GSE26193 1.17 [0.89, 1.54]
GSE26712 0.98 [0.83, 1.17]
GSE30161 0.64 [0.41, 0.99]
GSE32062.GPL6480 0.87 [0.70, 1.08]
GSE32063 1.60 [1.01, 2.52]
GSE49997 0.70 [0.51, 0.96]
GSE9891 0.87 [0.72, 1.04]
PMID17290060 1.51 [1.19, 1.91]
TCGA 1.10 [0.98, 1.23]
GSE51088 1.15 [0.92, 1.45]
OV.AU 0.95 [0.77, 1.17]
Overall 1.00 [0.95, 1.06]
0.12 0.5 1 2 4
Hazard Ratio
Plasmacytoid dendritic cell
E.MTAB.386 0.93 [0.73, 1.19]
GSE13876 0.91 [0.75, 1.11]
GSE14764 0.92 [0.57, 1.50]
GSE17260 0.88 [0.65, 1.19]
GSE18520 1.30 [0.92, 1.83]
GSE26193 1.16 [0.90, 1.50]
GSE26712 1.12 [0.93, 1.34]
GSE30161 0.68 [0.46, 1.00]
GSE32062.GPL6480 0.84 [0.70, 1.02]
GSE32063 1.31 [0.86, 1.98]
GSE49997 1.55 [1.12, 2.15]
GSE9891 1.10 [0.91, 1.33]
PMID17290060 1.22 [0.95, 1.57]
TCGA 1.12 [0.99, 1.27]
GSE51088 1.33 [1.05, 1.68]
OV.AU 1.08 [0.87, 1.35]
Overall 1.07 [1.01, 1.13]
0.12 0.5 1 2 4
Hazard Ratio