

Supplementary Materials: The Role of PTEN Loss in Immune Escape, Melanoma Prognosis and Therapy Response

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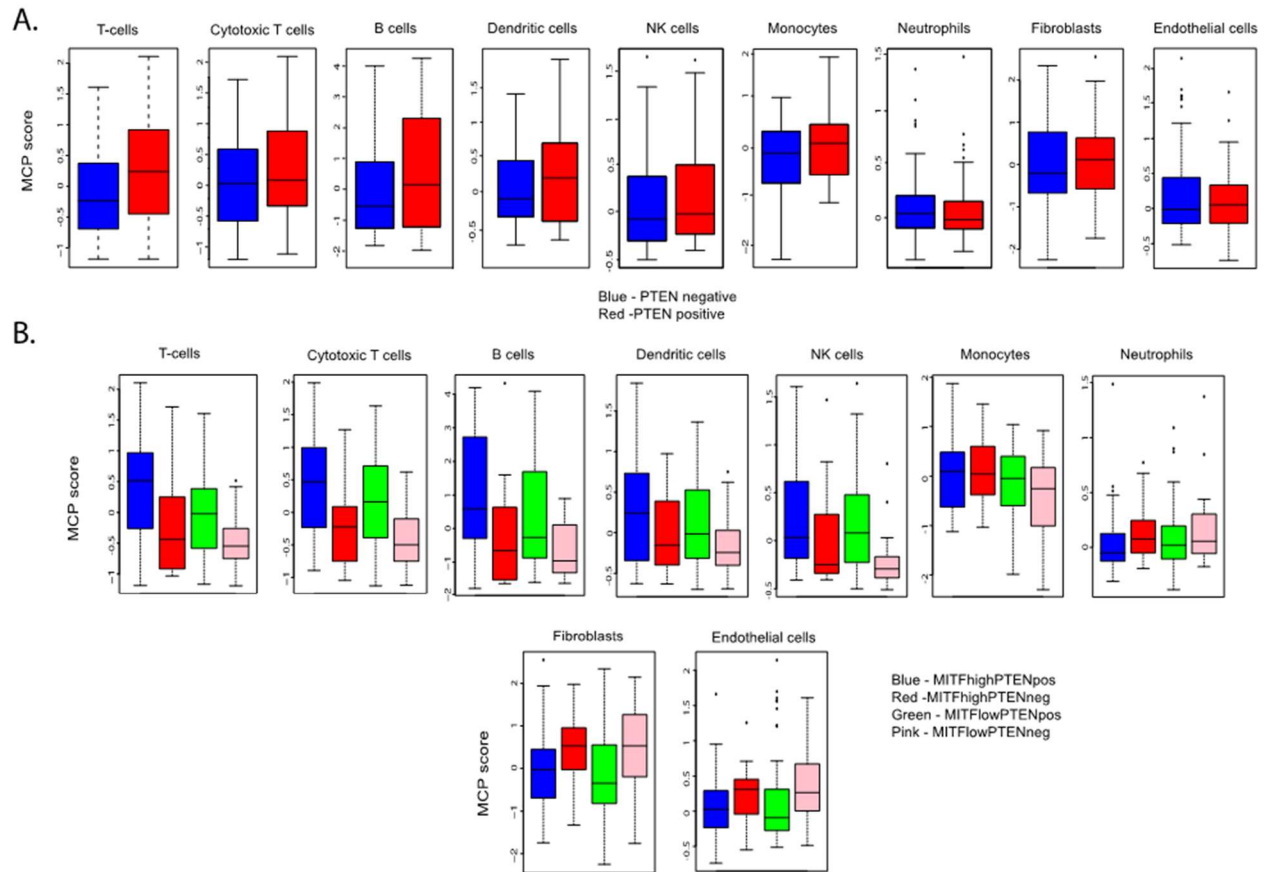


Figure S1. MCP immune scores in melanoma tumors. (A) MCP scores in PTEN negative (blue) and PTEN positive (red) tumors. (B) MCP scores in groups based on PTEN and MITF protein.

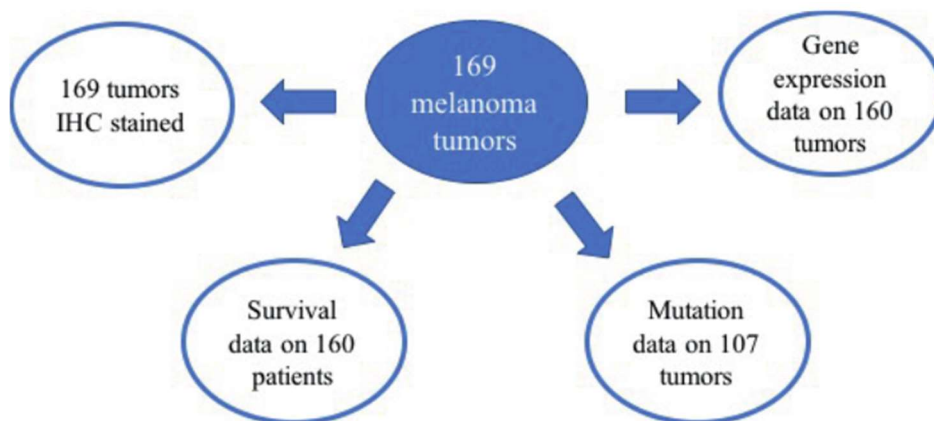


Figure S2. Flowchart of different data levels included in the study.

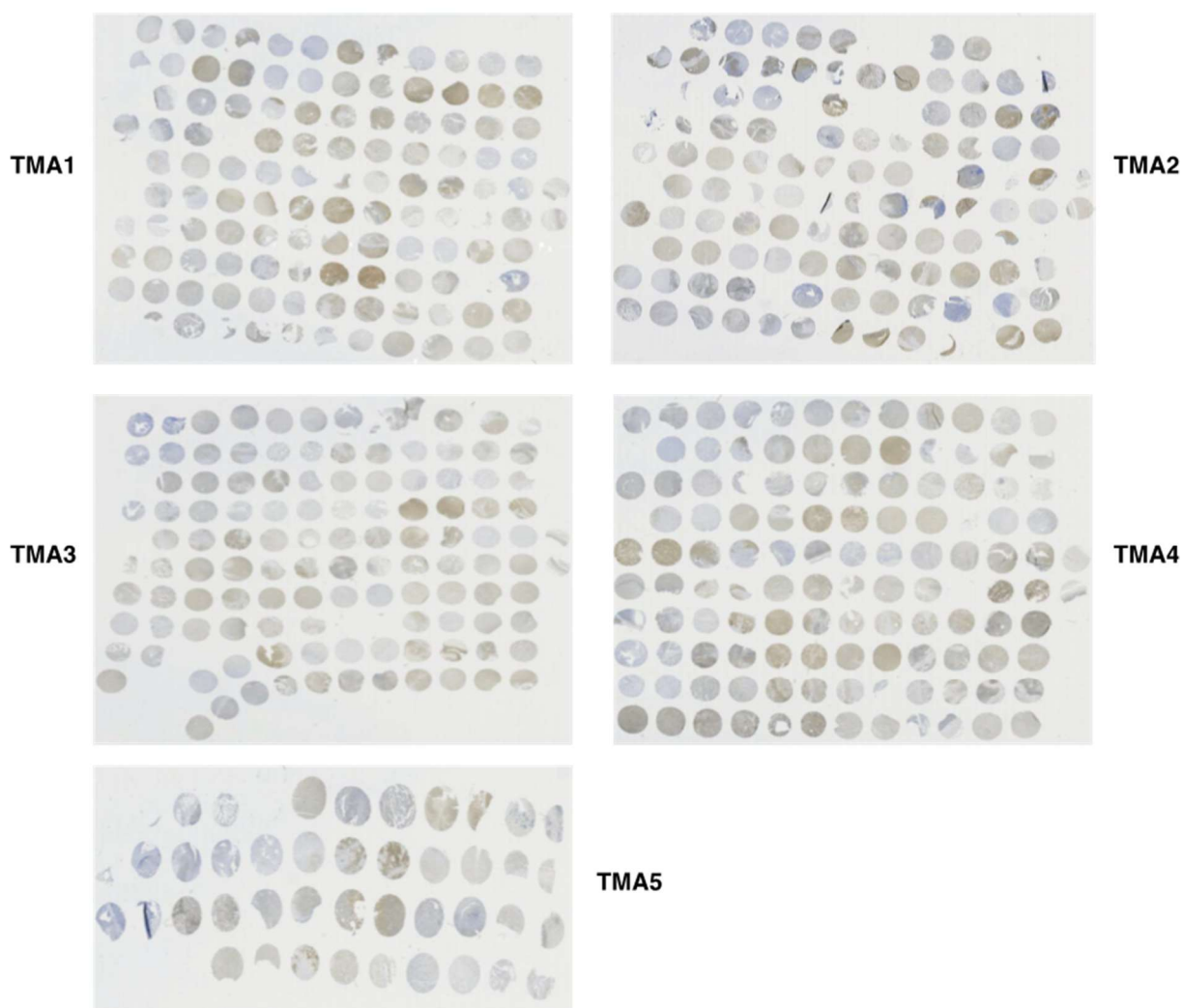


Figure S3. PTEN staining in the five tissue microarray (TMA) blocks.

Table S1. Multivariate cox regression model analysis of MITF/PTEN groupings adjusted for stage of disease, with hazard ratios and confidence intervals.

| Group | HR (CI) | <i>p</i> -Value |
|--|--------------------|-----------------|
| MITF/PTEN | | |
| MITF ^{high} PTEN ^{pos} | 1 | - |
| MITF ^{high} PTEN ^{neg} | 0.96 (0.56, 1.66) | 0.89 |
| MITF ^{low} PTEN ^{pos} | 0.90 (0.34, 2.41) | 0.84 |
| MITF ^{low} PTEN ^{neg} | 1.76 (0.89, 3.51) | 0.11 |
| Stage | | |
| II | 1 | - |
| III | 1.89 (0.26, 13.95) | 0.53 |
| IV | 7.29 (0.99, 53.73) | 0.05 |

Table S2. Genes and log2 expression value included in the centroid.

| Genes down in MITF ^{low} /PTEN ^{neg} | Value (log2 Expression) | Genes up in MITF ^{low} /PTEN ^{neg} | Value (log2 Expression) |
|---|-------------------------|--|-------------------------|
| ITGB7 | -0.6637874 | CADM1 | 1.646150435 |
| CD79A | -1.1278003 | DACT3 | 1.284409212 |
| TRIB1 | -0.627541 | SLIT2 | 1.069307448 |
| CXCL10 | -0.9425556 | PCDH18 | 1.272287668 |
| HLA-C | -0.6630066 | DIAPH3 | 1.049453117 |
| PLEK | -0.8497991 | C5orf13 | 1.463647947 |
| SDCBP | -0.6168393 | EFNB2 | 1.037998007 |
| VNN2 | -0.7245285 | LOC730994 | 0.816222521 |
| CSTB | -0.6004923 | HOXB5 | 0.890360545 |
| CD8A | -0.9936763 | MOXD1 | 1.464179261 |
| ANKRD22 | -0.6451211 | PCDH17 | 0.945373709 |
| CD52 | -1.1162775 | PHLDB2 | 0.937604318 |
| NCF1 | -0.6778936 | CDH2 | 1.690163862 |
| BHLHB3 | -0.748998 | DDAH1 | 0.904631728 |
| HSD17B14 | -0.6270172 | POSTN | 1.497294478 |
| PARP10 | -0.590338 | ID3 | 1.086589022 |
| BACE2 | -0.7382248 | FAM46A | 0.87567871 |
| IRF8 | -0.9998088 | SBK1 | 1.009141014 |
| GMPR | -0.7436598 | UBE2E3 | 0.702786405 |
| UPP1 | -0.6648916 | ERRFI1 | 1.163913307 |
| BIRC3 | -0.9908605 | NTM | 1.054703811 |
| PTPLA | -0.7938529 | PMEPA1 | 1.072653617 |
| SOX13 | -0.5899106 | ANGPT1 | 1.237634849 |
| ROPN1B | -0.8454191 | PCOLCE | 1.201493612 |
| TBC1D7 | -0.7636963 | FZD1 | 0.747943092 |
| CCL5 | -0.9803421 | PFTK1 | 0.727311372 |
| SNCA | -0.8087163 | SPRY2 | 0.991593274 |
| BCL2A1 | -0.6826054 | CDK6 | 0.99478154 |
| SLC11A2 | -0.6300387 | VCAN | 1.210859619 |
| NAPSB | -1.0304074 | DPYSL3 | 0.992375697 |
| MAZ | -0.5829541 | H19 | 1.385217542 |
| ISG20 | -1.0010139 | TUSC3 | 1.081069528 |
| FBXW5 | -0.6088084 | EDNRA | 0.748145714 |
| TAPBP | -0.6346449 | FZD2 | 0.640606599 |
| RAB32 | -0.6490322 | LMNB2 | 0.699635474 |
| INPP4B | -0.723364 | FLJ21986 | 0.98835468 |
| SWAP70 | -0.6226537 | ANTXR1 | 0.773896172 |
| MICAL1 | -0.6903372 | LOC100132901 | 0.628405303 |
| PHF15 | -0.6475682 | PXDN | 1.09475001 |
| BIRC7 | -0.9480172 | MLLT11 | 1.107994167 |
| UBE2L6 | -0.6231672 | S1PR3 | 0.701131508 |
| RARRES3 | -0.9509742 | ATP2B1 | 0.690079064 |
| LYN | -0.7620077 | CLIP3 | 0.842692443 |
| HCG4 | -0.7113144 | RBBP4 | 0.641617936 |
| GSDMD | -0.6139003 | SPON1 | 1.228882684 |
| RAC2 | -0.9906747 | CBX2 | 0.809775067 |
| SLC9A3R1 | -0.6598906 | HOXA5 | 0.87906882 |
| RUNX3 | -0.9927214 | LPHN2 | 0.959883606 |
| FAM69A | -0.6461069 | DKK3 | 0.96728898 |
| FCRLA | -1.2602482 | LOC100134134 | 1.132476535 |
| ST3GAL6 | -0.726269 | IGFBP2 | 1.175978716 |
| IRF1 | -0.855961 | LOX | 0.884465254 |

| | | | |
|-----------|------------|--------------|-------------|
| ECHDC3 | -0.6181085 | FOXC1 | 0.816302065 |
| CD38 | -0.7704543 | PAM | 0.799031548 |
| SIGIRR | -0.5547153 | LIMCH1 | 1.012433761 |
| ROPN1 | -0.9194562 | CPXM1 | 0.789543917 |
| FYB | -0.9039163 | ZNF286C | 0.591466564 |
| SNX10 | -0.6588212 | LOC730525 | 0.964789048 |
| HCST | -0.9173861 | LAMB1 | 0.836327667 |
| HERC5 | -0.8169465 | KLF12 | 0.635473272 |
| OAS2 | -0.9269088 | LOC730101 | 0.552153478 |
| MCOLN2 | -0.7311837 | BCAT1 | 0.86543584 |
| APOE | -0.8097618 | UBA52 | 0.71648786 |
| IFIH1 | -0.6855511 | ZNF84 | 0.57808192 |
| SAT1 | -0.5598643 | LOC100131139 | 0.746655149 |
| RAB38 | -1.0610584 | TSPAN5 | 0.759215173 |
| CD79B | -1.0782803 | TMTC4 | 0.660299277 |
| SOD2 | -0.7568718 | TCF12 | 0.670699942 |
| GNPTAB | -0.5581993 | AUTS2 | 0.785922266 |
| PSMB9 | -0.7884907 | ID1 | 0.782572387 |
| TMEM140 | -0.7171343 | TRIL | 1.002751536 |
| GZMB | -1.1113204 | KDM5B | 0.592819139 |
| HPS1 | -0.5510096 | OLFML2A | 0.657865462 |
| RPS6KA5 | -0.5670544 | LOC653344 | 0.557543286 |
| MFSD6 | -0.654256 | LRP5 | 0.677173513 |
| PRKCB1 | -1.0735466 | EFEMP2 | 0.795275881 |
| QPCT | -1.3166014 | ZNF512 | 0.539659589 |
| IFI27 | -1.2217255 | EXTL2 | 0.626539861 |
| RGS10 | -0.7862767 | KDEL1 | 0.58089505 |
| GSTO1 | -0.6408478 | GPC6 | 0.539021279 |
| CXCL9 | -1.2573445 | TRO | 0.739533058 |
| NKG7 | -1.0087833 | TMEM47 | 0.877205827 |
| PLXNC1 | -0.7586767 | LOC100133760 | 0.610353211 |
| RAB27A | -0.7035185 | RBP1 | 0.904180363 |
| KAT2B | -0.7054284 | RCAN2 | 0.815825294 |
| SLC45A2 | -1.3923654 | ZNF395 | 0.466799304 |
| ADAMDEC1 | -1.2114369 | COL8A1 | 1.104412183 |
| HIGD1A | -0.5175034 | LOC645166 | 0.708179748 |
| IL18BP | -0.9325305 | TLE4 | 0.855509022 |
| GPNMB | -0.8294207 | RGS4 | 0.808223983 |
| PHACTR1 | -0.89301 | ENC1 | 0.70166635 |
| CD3D | -1.3203313 | SOX4 | 0.746289566 |
| FAM65B | -0.9658844 | PEG10 | 0.912269898 |
| GBP5 | -0.977453 | LAMC1 | 0.640956415 |
| PARP9 | -0.6920852 | PKIA | 0.800216308 |
| HLA-B | -1.035889 | CNPY4 | 0.526343004 |
| LOC390557 | -0.711938 | CENPV | 0.637370896 |
| CD48 | -1.3089793 | LOC728732 | 0.588492945 |
| HCP5 | -1.1201781 | SSFA2 | 0.591275715 |
| HLA-F | -0.9897719 | FSCN1 | 0.845168008 |
| TRIM63 | -1.0193057 | MEIS2 | 0.698440229 |
| PLA1A | -1.3654242 | TCF3 | 0.511660304 |
| TYSND1 | -0.5696128 | SOCS2 | 0.711013533 |
| CXCL13 | -1.3775115 | ZNF532 | 0.661004424 |
| MGC13057 | -0.8628089 | NR4A2 | 0.792124417 |
| CABLES1 | -0.9646213 | SKP2 | 0.550286352 |
| MCOLN3 | -1.1803748 | STK32B | 0.771656854 |
| HRK | -0.9979237 | COL16A1 | 0.57506291 |

| | | | |
|------------|------------|-----------|-------------|
| OSTM1 | -0.7563323 | TMSB15A | 0.974167792 |
| ASB9 | -0.7821446 | SALL2 | 0.523090784 |
| FAIM3 | -1.2771405 | SYT11 | 0.685722901 |
| VAT1 | -0.8184453 | SMA4 | 0.490970275 |
| ARL6IP5 | -0.5472999 | COL5A1 | 0.915868645 |
| S100A1 | -1.2352086 | ITGB5 | 0.759491207 |
| GYG2 | -1.1467153 | ESM1 | 0.802307819 |
| LY96 | -0.8734827 | ENAH | 0.536317296 |
| CD44 | -0.8390348 | COL12A1 | 0.736125394 |
| HLA-E | -0.8544589 | LOC645385 | 0.519091808 |
| FBXO6 | -0.7278542 | PRDM8 | 0.58278025 |
| HLA-G | -0.9261851 | EMILIN1 | 0.681825454 |
| MITF | -1.0316963 | PRICKLE1 | 0.673123973 |
| RTP4 | -0.7450338 | COL5A2 | 0.880003949 |
| SERPINB1 | -0.765538 | CPE | 0.766764884 |
| CBR3 | -0.7069554 | SDC2 | 0.812521025 |
| ST6GALNAC2 | -1.0839479 | HNRPA1L-2 | 0.564260655 |
| MYO5A | -0.8894011 | PRICKLE2 | 0.62877922 |
| PSMB8 | -0.8434825 | SPRY1 | 0.757695396 |
| PRKCD | -0.7213112 | | |
| CEACAM1 | -1.3066298 | | |
| IRF4 | -1.0175048 | | |
| IFI35 | -0.8226885 | | |
| STX7 | -0.7898291 | | |
| TRPV2 | -0.9233402 | | |
| CHCHD6 | -1.3239017 | | |
| CASP1 | -1.1742113 | | |
| STAT1 | -1.0606008 | | |
| P2RX7 | -0.9106965 | | |
| LYST | -1.1111641 | | |
| CAPN3 | -1.5022855 | | |
| MBP | -1.1579674 | | |
| PIR | -1.1636714 | | |
| TAP1 | -1.0964007 | | |
| LGALS3 | -1.1585214 | | |
| TNFRSF14 | -1.0275618 | | |
| WARS | -1.182983 | | |
| GPR56 | -1.3043394 | | |
| UBD | -1.6572527 | | |
| RENBP | -1.1237332 | | |
| D4S234E | -2.0144985 | | |
| TYR | -2.3769119 | | |
| SILV | -2.8824723 | | |
| MLANA | -2.3966163 | | |



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