Supplemental Table 4. List of candidate off-target peptides shared by TCR_{a3a}, TCR_{A3-05} and TCR_{A3-10}, as predicted from peptide scanning of the MAGE-A3₁₆₈₋₁₇₆ target using TnT-TCR cells as effectors, Related to Figure 6.

2 A16L2 467 505 QVIPVOGRV 62 CLD15 146 154 FFDPLYPGT 122 FR1L6 977 985 QVIPVPAN 182 MARE1 234 242 ENDPVLORI 242 PLD5 387 386 ETDPLTNF 330 A1CF 71 79 ELIPLCEKI 65 DND2 240 248 EIDPMFOKT 123 FSD1L 201 209 EIDPVECLV 163 MEDB 91 99 QVIPLADYYY 243 PP188 107 115 QDIPLSTV 330 ALPIEAN 197 295 FFDPLY 1 64 CNDT1 710 7718 QDIPLSTV 124 GATA1 76 84 QVYPLINGM 184 MEPC 319 327 EGDPVDGTC 244 PP181 107 115 QDIPLSTV 330 ALPIEAN 197 197 318 QDIPLSTV 1 125 QBG 144 52 QHDPLTGV 185 MGST2 58 66 EYPIFIT 248 PR14L 706 713 QTIPLGTK 1 30 ABCC 531 539 QLIPLEAM 65 CNDT 66 CNDT 162 170 EIPLCIR 126 GFD12 217 225 EOIPLVRT 185 MGST2 58 66 EYPIFIT 248 PR14L 706 713 QTIPLGTK 1 30 ABCC 351 539 QFDPLSTV 1 56 CNDT	Department of the control of the con	position position Sequence
2 A16L2 497 508 QVIPVOGRV 62 CLD15 146 156 FFDFLYPGT 122 FR16 977 988 QIVPVAN 182 MARE1 234 242 ENDPVLORI 242 PLD5 387 386 ETDFLTFNF 330 A1CF 77 79 ELIFLCEKI 63 CND2 240 248 EIGDPMCKT 123 FSD1L 201 208 EIDPVECLV 183 MEDB 91 99 QVIPLADMY 244 PPIMB 107 115 QOIPIDSTV 33 A4 AAK1 977 325 EFDPIPVLI 64 CND1 710 778 GIDPLADMT 124 GATA1 76 81 GVYPLDM 183 MEDB 91 99 QVIPLADMY 244 PPIMB 107 115 QOIPIDSTV 33 A4 AAK1 977 325 EFDPIPVLI 64 CND1 710 778 GIDPLADMT 124 GATA1 76 81 GVYPLDM 185 MGSTZ 58 66 EYPIFIIT 248 PR14L 706 773 QTIPICITI 330 ABCC 531 539 QLIPIEAM 65 CND14 377 385 EIDPLSSST 125 GBS 44 52 QHDPLLTGV 185 MGSTZ 58 66 EYPIFIIT 248 PR14L 706 773 QTIPICITI 330 ABCC 337 385 EVDPLSSST 125 GBS 24 A4 52 QHDPLLTGV 185 MGSTZ 58 66 EYPIFIIT 248 PR14L 706 773 QTIPICITI 330 ABCC 337 385 EVDPLSSST 125 QBS 340 EIDPLSSST 125 QBS 340 EIDPLSSST 345 QBS 347 PRD16 862 870 PRDPYSSV 345 ABCC 347	02 THIL 03 TIF1A 04 TIGIT 05 TITIN 06 TKFC 07 TMIL1 08 TMM62 09 TPC11 10 TR49B 11 TR64C 12 TR111 13 TR117 14 TR141	234 242 EVIPVTVTV 924 932 PODPVPLTV 138 146 FOIPLIGAM 24337 24345 ESDPIVAQY 128 138 EGIPVEMV 446 454 EFDPLAPAV 539 547 FNIPLMAYM 993 1001 ENPITTV 22 30 FIDPVTIDC 22 30 FIDPVTIDC
3 AICF 71 79 ELIPLCEKI 63 CND2 240 248 EIDPMCKT 123 FSDIL 201 200 EIDPMCLV 183 MED8 91 99 QVIPLADYY 243 FPIR8 107 115 QQIPLDSTV 330 AAX1 97 925 EDPIPVLI 64 CNOT1 710 718 QIPLAGMT 124 GATA1 76 84 QVVPLINGM 184 MEEV 319 327 EGDIPUSGTC 244 PPMMH 298 306 EIPPMSSEF 33 CARDINARY 185 METS 25 86 66 EPYPHITI 245 PRI4L 705 713 GIPLAGMT 124 GATA1 76 84 QVVPLINGM 184 MEEV 319 327 EGDIPUSGTC 244 PPMMH 298 306 EIPPMSSEF 33 CARDINARY 185 METS 25 86 66 EPYPHITI 245 PRI4L 705 713 GIPLAGMT 137 385 EIPPWSSET 125 GBS 44 52 QMDPLTGV 185 MEST 25 86 66 EPYPHITI 245 PRI4L 705 713 GIPLAGMT 137 385 EIPPWSSET 125 GBS 44 52 QMDPLTGV 185 MEST 25 86 66 EPYPHITI 245 PRIAL 705 713 GIPLAGMT 137 ASC 241 247 PRIDE 247 247 PRIDE 247 247 PRIDE 247 247 PRIDE 247 PRIDE 247 PRIDE 247 PRIDE 247 247 PRIDE 2	03 TIF1A 04 TIGIT 05 TITIN 06 TKFC 07 TM1L1 08 TMM62 09 TPC11 01 TR49B 11 TR84C 12 TR111 13 TR117 14 TR141	138 146 FOIPLLGAM 24337 24345 ESSPIVACY 128 136 EGIPVEMVV 446 454 EFEPLAPAV 539 547 FNIPLMAYM 993 1001 ENIPITTV 22 30 FIDPVTIDC 22 30 FIDPVTIDC
6 ABCAC 531 539 QLIPIEAM 66 CNOT4 377 386 ETIPVSSST 125 GBG5 44 52 QHDPLITGV 185 MGST2 56 66 EPYPIFIIT 245 PR14L 705 713 QTIPIQTKI 330 ABCAC 886 886 QAIPMA, ONT 66 CNOT9 162 170 EIIPLCLR 126 GFTP2 217 225 EQIPILYRT 186 MCNA 208 216 QNDPVTVVV 246 PRC2B 1957 1965 QQIPISLIHT 33 ABCAC 357	05 TITIN 06 TKFC 07 TM1L1 08 TMM62 09 TPC11 10 TR49B 11 TR64C 12 TR111 13 TR117 14 TR141	24337 24345 ESDPIVAQY 128 136 EGIPVEMVV 446 454 EFDPLAPAV 539 547 FINPLMAYM 993 1001 ENIPIITV 22 30 FIDPVTIDC 22 30 FIDPVTIDC
6 ABCAC 858 866 QAIPMLONT 66 CNOTS 162 170 EIIPLCLR 126 GFPTZ 217 228 EQIPLYRT 186 MNDA 208 216 CNDPVTVV 246 PRC28 1957 1968 QQIPISLHT 30 ABCAC 357 ABCGA 35	06 TKFC 07 TM1L1 08 TMM62 09 TPC11 10 TR49B 11 TR64C 12 TR111 13 TR117 14 TR141	128 136 EGIPVEM/V 446 454 EFDFLAPAV 539 547 FNIPLMAYM 993 1001 ENIPIITTV 22 30 FIDPVTIDC 22 30 FIDPVTIDC
Record 367 368 EVOPIESHT 67 CATIN3 524 532 GHOPLIDII 127 GLT16 270 278 EQIPLEQKM 187 MOCS1 549 557 QLIPLOHW 247 PRD16 862 870 PMOPIYSRV 368 368 ABECA 259 257 ESYPLANAF 68 COS646 1982 1900 EIRIVATE 128 GRR18 196 204 FLIPLINII 188 MOCS 65 63 FUPULITIV 248 PRD16 184 192 EIRIPADHT 33 330 ELEPILITY 128 GRR18 128 EIRIPADHT 33 46 ESYPLENAF 170 COG5 268 273 QVIPULGEV 129 GRR13 257 268 OFFYPULIT 189 MRC1 570 578 OFFYPULIT 248 PRD16 144 122 EVOPLYNM 31 11 ACAD 239 EGYPLETIT 71 CP246 390 398 EVYPMCSV 130 GRIA1 531 531 ESTPLETIT 190 MRC18 77 78 QDDPLSEAI 250 FRS10 114 122 EVOPLYNM 31 124 EBF19 196 240 GGIFANAT 72 CP345 216 223 ELPPLETIT 132 GDTC1 51 59 OTFISEHY 128 MRC18 1804 1812 GMPT MRC18 1804 1814	07 TM1L1 08 TMM62 09 TPC11 10 TR49B 11 TR64C 12 TR111 13 TR117 14 TR141	446 454 EFDPLAPAV 539 547 FNIPLMAYM 993 1001 ENIPIITTV 22 30 FIDPVTIDC 22 30 FIDPVTTDC
8 ABEC4 259 287 ESYPLANAF 68 CO8A6 1892 1900 EIIPVVITF 128 GPR18 196 204 FLIPLFIMI 189 MODS 85 93 PVDPLVTTY 248 PRDM8 184 192 EIIPLNDHT 330 PACACA 2174 2182 FLIPLYAGY 70 COS 265 273 QVMPVLGEV 129 GRR4.3 257 256 GPYPVTLRT 189 MORC1 570 578 GMPVDETT 249 PRDM1 42 50 FRYPLDFTF 330 PACACA 2174 229 EGYPLAGY 70 COS 265 273 QVMPVLGEV 130 GRR14 531 539 FLDPLYE! 190 MROBB 70 78 GODPLSEAL 259 PRS10 114 122 EVDPLYYBM 331 11 ACLTA 231 239 EGYPLSTIT 71 CP2A6 390 398 EVYPMLGSV 131 GST2 110 118 FGIPLWOV 191 MTFC2 56 64 ELIPLLINSV 251 PTPR8 1131 1139 GVDPLVGSF 331 12 AEBP1 916 924 GGIPLAGAT 72 CP3A5 215 223 FLOPLYSI 132 GTDC1 51 56 GTRISEHY 192 MORAS 1804 1812 GMPMWSST 252 PLRB 92 100 EMPIEWVC 331 134 AFAD 1275 1225 EAYPIPTOT 73 CPEDI 371 379 FMPVPLVQV 133 HACO3 299 307 GSIPIPNET 192 MORAS 1804 1812 GMPMWSST 252 PLRB 92 100 EMPIEWVC 331 14 AGRE4 300 308 BMPVPGVGI 174 CPUNI 333 371 EFIDLYBLI 134 HACAS 996 1004 ELDPLPEKV 194 MV15B 737 745 EAPLAGGI 254 RBBPB 385 373 GODPLMBV 331 16 AGRC1 14 25 GS EAVPLEACK 77 GRG3 1305 HAMPE 133 HACO3 299 GUIPLNET 196 MV10H 174 134 GRR14 256 GRR45 149 GUIPLNET 197 MV15B 737 745 EAPLAGGI 254 RBBPB 385 373 GODPLMBV 331 16 AGRC1 14 25 GS EAVPLEACK 77 GRG3 1305 HAMPE 135 HACO3 137 HEIDLY 135 HACO3 137 HEIDLY 136 HACO3 137 HEIDLY 137 HACO3 137 HEIDLY 138 HACO3 137 HEIDLY 139 MV15B 737 745 EAPLAGGI 254 RBBPB 385 373 GODPLMBV 331 15 AGRC1 14 25 GS EAVPLEACK 77 GRG3 1305 HAMPE 1434 HACO3 137 HEIDLY 136 HACO3 137 HEIDLY 137 HELD 138 HACO3 137 HEIDLY 138 HACO3 137 HEIDLY 139 MV15B 737 745 EAPLAGGI 254 RBBPB 385 373 GODPLMBV 331 15 AGRC1 14 255 EAVPLEACK 77 GRG3 1305 HAMPE 135 HACO3 137 HELD 135 HA	08 TMM62 09 TPC11 10 TR49B 11 TR64C 12 TR111 13 TR117 14 TR141	539 547 FNIPLMAYM 993 1001 ENIPIITTV 22 30 FIDPVTIDC 22 30 FIDPVTIDC
9 ACACA 2174 2182 FLIPIYHOV 66 COG4 351 356 ELDPILTEV 129 GRNL3 257 286 QFYPVTLRT 189 MORC1 570 578 QFIPVDEIT 240 PRDX1 42 50 FFYPLDFTF 330 ACADM 59 67 EINPVAREY 70 COC5 265 273 QVIPVLEVEV 130 GRNA1 531 538 FLDPLAYEI 190 MROHB 70 78 QODPLEAN 250 PRS10 114 122 EVDPLVYNM 331 114 ACATA 231 239 EGYPLRSTT 71 CP2A6 390 398 EVYPMLGSV 131 GST2 110 118 FGIPLWOV 191 MTR2 56 64 ELIPLILISV 251 PTRB 1131 1139 QVDLVQSF 331 12 AEBP1 916 924 QGIPLANT 72 CP3A5 215 223 FLDPLRSI 132 GTDC1 51 59 QTIPLSETH 192 MORA5 1804 1812 QMPMSST 252 PUR6 92 100 EMPIEWOV 331 131 AFAD 1217 1225 EAVIPITOT 73 CPED1 371 379 FMPWLVQV 133 HADG3 299 307 QSIPIPNET 193 MORA5 2544 2552 FHDPISEKI 253 RASHB 289 297 QLIPLEET 331 154 AGR66 716 724 QVDPLASVI 76 CPLN1 363 371 EFIPLHPUI 134 HADG4 996 1004 ELDPLPEVV 194 MY15B 737 745 EAIPLAPGI 254 RBBP6 365 373 QODPLMPV 331 156 AGR66 716 724 QVDPLASVI 75 CPLN1 2416 2424 QLIPLENLI 135 HD 2801 ELDPLEVV 194 MY15B 737 745 EAIPLAPGI 254 RBBP6 365 373 QODPLADV 331 156 AGR66 716 724 QVDPLASVI 75 CPLN1 2416 2424 QLIPLENLI 135 HD 2801 ELDPLEVV 194 MY15B 737 745 EAIPLAPGI 254 RBBP6 365 753 751 FAYPLOWSI 331 156 AGR66 716 724 QVDPLASVI 75 CPLN1 2416 2424 QLIPLENLI 135 HD 2801 ELDPLEVV 195 MYBB 388 386 386 ELLIPLSTST 255 RBM25 753 751 FAYPLOWSI 331 156 AGR66 756 CRBG3 1635 1645 EVIPMMPEV 136 FELI 160 ELDPLEVV 196 MY01H 174 182 QGIPVGGH 256 RBP1 190 198 FGIPLADAV 331 17 AGR13 37 45 ESYPIELRC 77 CRBG3 170 ETIPLY 176 EVIPMENT 177 ELIPLMEVT 196 MY01H 174 182 QGIPVLTC 256 RBP1 190 198 FGIPLADAV 331 18 ALPK3 1702 1710 EILPLYLIV 75 CRBM 79 87 QVIPVLPOV 138 HERC1 4483 4291 QCIPVLQV 199 MVOZ2 242 250 EMPIVIT 257 RELCH 541 549 ELIPLICT 331 19 AMPE 711 719 ELIPLMEVT 71 719 ELIPLMEVT 71 719 ELIPLMEVT 256 FGELN 1228 EDIPLICT 331 19 AMPE 711 719 ELIPLMEVT 71 719 ELIPLMEV	09 TPC11 10 TR49B 11 TR64C 12 TR111 13 TR117 14 TR141	993 1001 ENIPIITTV 22 30 FIDPVTIDC 22 30 FIDPVTIDC
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16 AGRL1 42 50 EGYPIELRC 76 CRBG3 1638 1648 EVIPMMPEV 136 HELQ 1009 1017 ELIPLMEVT 198 MYO1H 174 182 QGIPVGGHI 256 R8P1 199 198 FGIPLADAV 31 17 AGRL3 37 45 ESYPIELRC 77 CRBG3 1706 1714 EVIPVLAM 137 HELZ2 612 620 QTDPVLQV 197 MYO2Z 242 250 EMPIVITT 257 RELCH 541 549 ELIPLICT 31 18 ALPK3 1702 1710 EIIPLYLIY 78 CRBM 79 87 QVIPVLPQV 138 HERC1 4283 4291 QGIPVLGV 198 NCHL1 145 153 EGDPVLPC 258 RELN 1228 1234 QIPVLDV 31 199 MELT 17 719 ELIPMINETY 79 CSDE1 279 287 (NOPPLPGN 139 HERC1 4483 4291 QGIPVLGVV 199 NCHL1 15 153 FG ENDIVLTE 259 REVSL 2217 2225 ENDICALF 31	15 TRI51	27 35 FTDPVSIGC
17 AGRL3 37 45 ESYPIELRC 77 CR8G3 1706 1714 EVIPVTLAM 137 HELZ2 612 620 QTDPVTLQY 197 MYOZZ 242 250 ENIPIVITT 257 RELCH 541 549 ELIPLILCT 31 18 ALPK3 1702 1710 EVIPVTLY 78 CR8N 79 87 QVIPVLPQV 138 HERC1 4283 4291 QQIPVLAGV 198 NCHL1 145 153 ECOPVLPC 258 RELN 1226 1224 QVIPVNPT 31 199 AMPE 711 719 ELYPMIEEY 79 CSDE1 279 287 QNDPLPGRI 139 HERC1 4647 4655 EMIPLDSFV 199 NELFD 571 579 EMDPVTEFI 259 REV3L 2317 2325 EFDPICALF 31		22 30 FLDPVTIDC
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19 AMPE 711 719 ELYPMIEEY 79 CSDE1 279 287 ONDPLPGRI 139 HERC1 4647 4655 EMIPLDSRV 199 NELFD 571 579 EMDPVTEFI 259 REV3L 2317 2325 EFDPICALF 31	17 TRIM4 18 TRIO	19 27 FQDPVSIEC 2687 2695 FVIPLSEVT
	19 TSH1	466 474 QSIPLPPTT
20 AN13A 377 385 QVIPIIDLM 80 CSKP 767 775 FAYPIPHTT 140 HEXD 82 90 EVIPLYQTF 200 NEMF 388 396 QGDPVASAI 260 RFIP3 118 126 ELDPLFSWT 32	20 TSHB	95 103 FSYPVALSC
	21 TT21A	441 449 QGIPLGSEY
22 ANR16 147 155 EGDPLILQY 82 CTR4 540 548 FQIPM/PU 142 HNRC1 250 258 EGDPLDDDV 202 NEP1 178 186 FSIPV/V5DV 262 RHAG 49 57 ELYPLFODV 32	22 TT30A	438 446 ENYPMVEKV
	23 TT30B	438 446 ENYPMVEKI
	24 TTC14	535 543 ECYPVPANT
	25 TTLL2	176 184 QFIPLTFVM
	26 TWF2 27 TX13C	246 254 EGDPLESVV 154 162 QVYPMPQDF
	27 TX13C 28 UBE2W	76 84 ENIPVHPHV
	29 UBP38	242 250 QHIPLQMIT
	30 UBP8	658 666 FLDPITGTF
31 AT134 544 552 QGDPLDLKM 91 DIP28 426 434 EVIPVPIEV 151 QCN 341 349 QTYPVVSVT 211 NUMB 509 517 QSYPVANGM 271 RUSD2 494 502 ETDPLCAEC 33	31 URAS1	24 32 FLIPVALRV
32 ATP7A 660 668 FCIPVMGLM 92 DLG1 822 830 QLYPLISFI 152 IRF6 393 401 QVIPVVARM 212 OS1A7 27 35 FSIPICLMY 272 S12A3 487 495 QLYPLIGFF 33	32 UTP20	2326 2334 FFIPLCLMT
	33 VAFNB	87 95 EMYPVPPIT
	34 VATA	417 425 FSDPVTSAT
	35 VLDLR	126 134 QCIPVSWRC
	36 VW5B1 37 WASH1	49 57 FVYPLDECT 188 196 FLDPLAGAV
	37 WASH1 38 WDFY4	313 321 EGYPLLKV
	39 WDR27	491 499 EAYPVECAV
	40 WFS1	343 351 FFIPLVIFY
41 BUD13 541 549 EGDPMANFI 101 DYH7 836 844 QHIPLIQVI 161 KDMZA 303 311 FNIPMDLKI 221 OR6K6 55 63 FFIPLLLIY 281 SCNNA 110 118 FSYPVSLNI 34	41 XPO2	40 48 QNYPLLLLT
	42 XPO2	913 921 EHDPVGQMV
	43 XPOT	741 749 EFIPLINQI
	44 ZC3HE	438 446 QIDPVMAET
	45 ZDHC5 46 ZDHC8	196 204 FFIPVAGLT 196 204 FFIPVIGLT
	46 ZDHC8 47 ZMYM2	902 910 QNIPVPTTV
	48 ZMYM4	1005 1013 FGIPVPMPV
	49 ZN395	416 424 FQIPVSPHI
	50 ZN541	916 924 QSIPVVPVT
	51 ZN608	1382 1390 FHYPVYGKM
	52 ZN791	393 401 FNYPLDLKI
	53 ZNF8	240 248 QAIPITELT
	54 ZSWM8	488 496 EAYPLPGVT
55 CEAMI 406 414 QSDPIMLNV 115 FBX30 717 725 EAIPLPCMC 175 LTBP4 1124 1132 EFDPMTGRC 235 PGCA 53 61 FIDPMHPVT 295 TDRD9 414 422 QVYPLHSSV 56 CEAMS 225 233 FSDPVTLNV 116 FCERA 206 214 FFIPLLVVI 176 LYAM2 357 365 QQIPVCEAF 236 PHB2 75 83 FQYPIIVDI 296 TEAD1 178 186 QAYPIQPAV	+ -	
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59 CFA46 1999 1977 QADPVHPTC 119 RIP1 200 208 EVIPVTSTT 179 MAGA3 168 176 EVDPIGHLY 239 PI4KA 369 377 FSDPLYLTM 299 TF3C4 449 457 QLIPIFTDV		
60 CFA47 2661 2669 QIPLVNCT 120 PNBP1 240 248 QVIPLIGKC 180 MAGA6 168 176 EVDPIGHVY 240 PIGB 362 370 FYPVLPFC 300 TF65 480 488 QGIPVAPHT		